TC-Python Documentation

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Thermo-Calc Software AB

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CHAPTER

ONE

INSTALLATION GUIDE

This guide helps you to get a working TC-Python API installation.

There is a PDF guide included with your installation. In the Thermo-Calc menu, select $Help \rightarrow Manuals$ Folder. Then double-click to open the Software Development Kits (SDKs) folder.

Note: A license is required to run TC-Python.

1.1 What type of installation should I choose?

There are two possibilities to install TC-Python:

- 1. Using the Python-interpreter bundled to Thermo-Calc: This interpreter has TC-Python preinstalled together with some popular Python-packages. This is the recommended option for new users to TC-Python, but it is limited to the preinstalled packages.
- 2. Installing TC-Python into the Python-interpreter of your choice: This is the recommended option for any more advanced usage and provides full flexibility.

1.2 Using the Python-interpreter bundled to Thermo-Calc

Note: A Python-interpreter is bundled to Thermo-Calc beginning with version 2021a.

1.2.1 Limitations

The bundled Python 3.7.2 interpreter is containing the following major packages:

Package	Version		
matplotlib	3.3.2		
numpy	1.19.2		
scikit-learn	0.23.2		
scipy	1.5.2		
TC-Python	2022b		

Please contact the Thermo-Calc support if you think that further packages might be useful in future releases.

Note: The following TC-Python examples are requiring additional packages that are not available in the bundled Python-interpreter, they can therefore not be run:

- pyex_M_01_Input_from_file.py
- pyex_M_02_Output_to_file.py

Warning: The Python-interpreter bundled to Thermo-Calc is also used for running the property models in Thermo-Calc. **Any changes to the interpreter packages can therefore break Thermo-Calc and should be avoided**. If the installation gets broken, it can be fixed by reinstalling Thermo-Calc after having removed it.

1.2.2 Step 1: Install an IDE (Integrated Development Environment)

Any editor can be used to write the Python code, but an IDE is recommended, e.g. PyCharm. These instructions are based on the use of PyCharm.

Use of an IDE will give you access to code completion, which is of great help when you use the API as it will give you the available methods on the objects you are working with.

- 1. Navigate to the PyCharm website: https://www.jetbrains.com/pycharm/download.
- 2. Click to choose your OS and then click **Download**. You can use the **Community** version of Py-Charm.
- 3. Follow the instructions. It is recommended you keep all the defaults.

Note: For Mac installations, you also need to set some environment variables as described below in *Mac OS: Setting Environment Variables.*

1.2.3 Step 2: Configure PyCharm to use the bundled Python-interpreter

Open PyCharm and configure the interpreter:

- 1. Go the menu **File** \rightarrow **Settings**.
- 2. Navigate in the tree to Project: YourProjectName and choose Project Interpreter.
- 3. Click on the settings symbol close to the Project Interpreter dropdown menu and choose Add.
- 4. Now choose **System Interpreter** and add the bundled Thermo-Calc Python 3 interpreter. It is located in different places depending on the operating system:

Operating system						
Windows C:\Program Files\Thermo-Calc\2022b\python\						
	python.exe					
Linux	/home/UserName/Thermo-Calc/2022b/python/bin/					
	python3					
MacOS	/Applications/Thermo-Calc-2022b.app/Contents/					
	Resources/python/bin/python3					

5. Select your added interpreter and confirm.

1.2.4 Step 3: Run a TC-Python Example

Now you are ready to start working with TC-Python.

It is recommended that you open one or more of the included examples to both check that the installation has worked and to start familiarizing yourself with the code.

1.2.4.1 Open the TC-Python Project in PyCharm

When you first open the TC-Python project and examples, it can take a few moments for the Pycharm IDE to index before some of the options are available.

1. Open PyCharm and then choose **File**→**Open**. The first time you open the project you will need to navigate to the path of the TC-Python installation:

Operating sys-	Path to the TC-Python folder				
tem					
Windows	C:\Users\UserName\Documents\Thermo-Calc\2022b\SDK\				
	TC-Python				
Linux	/home/UserName/Thermo-Calc/2022b/SDK/TC-Python				
MacOS	/Users/Shared/Thermo-Calc/2022b/SDK/TC-Python				

- 2. Click on the Examples folder and then click OK.
- 3. From any subfolder:
 - Double-click to open an example file to examine the code.
 - Right-click an example and choose Run .

Note: If you are not following the recommended approach and create a *new* project (**File** \rightarrow **New Project...**), you need to consider that by default the options to choose the interpreter are hidden within the **Create Project** window. So click on **Project Interpreter:** New Virtual Environment and in most cases choose your *System Interpreter* containing the Python bundled to Thermo-Calc instead of the default *New Virtual Environment*.

1.3 Installing TC-Python into the Python-interpreter of your choice

1.3.1 Step 1: Install a Python Distribution

If you already have a Python distribution installation, version 3.5 or higher, skip this step.

These instructions are based on using the Anaconda platform for the Python distribution. Install version 3.5 or higher to be able to work with TC-Python, although it is recommended that you use the most recent version.

1.3.1.1 Install Anaconda

- 1. Navigate to the Anaconda website: https://www.anaconda.com/download/.
- 2. Click to choose your OS (operating system) and then click **Download**. Follow the instructions. It is recommended you keep all the defaults.

1.3.2 Step 2: Install Thermo-Calc and the TC-Python SDK

Note: TC-Python is available starting with Thermo-Calc version 2018a.

- 1. Install Thermo-Calc
- 2. When the installation is complete, open the TC-Python folder that includes the *.whl file needed for the next step. There is also an file:*Examples* folder with Python files you can use in the IDE to understand and work with TC-Python.

1.3.3 Step 3: Install TC-Python

On Windows, it is recommended that you use the Python distribution prompt (i.e. Anaconda, \dots), especially if you have other Python installations. Do not use Virtual Environments unless you have a good reason for that.

- 1. Open the command line. For example, in Anaconda on a Windows OS, go to **Start→Anaconda→Anaconda Prompt**.
- 2. At the command line, enter the following. Make sure there are no spaces at the end of the string or in the folder name or it will not run:

Tip: Note that on Linux depending on the interpreter usually *pip3* is used.

Operating sys-	Path to the TC-Python folder				
tem					
Windows	C:\Users\UserName\Documents\Thermo-Calc\2022b\SDK\				
	TC-Python				
Linux /home/UserName/Thermo-Calc/2022b/SDK/TC-Py					
MacOS	/Users/Shared/Thermo-Calc/2022b/SDK/TC-Python				

3. Press <Enter>. When the process is completed, there is a confirmation that TC-Python is installed.

Note: If your computer is located behind a proxy-server, the default *pip*-command will fail with a network connection error. In that case you need to install the dependencies of TC-Python in a special configuration:

pip install -proxy user:password@proxy_ip:port py4j jproperties

See "pip install" fails with "Failed to establish a new network connection" or similar for detailed information.

1.3.4 Step 4: Install an IDE (Integrated Development Environment)

Any editor can be used to write the Python code, but an IDE is recommended, e.g. PyCharm. These instructions are based on the use of PyCharm.

Use of an IDE will give you access to code completion, which is of great help when you use the API as it will give you the available methods on the objects you are working with.

- 1. Navigate to the PyCharm website: https://www.jetbrains.com/pycharm/download.
- 2. Click to choose your OS and then click **Download**. You can use the **Community** version of Py-Charm.
- 3. Follow the instructions. It is recommended you keep all the defaults.

Note: For Mac installations, you also need to set some environment variables as described below in *Mac OS: Setting Environment Variables.*

1.3.5 Step 5: Open the IDE and Run a TC-Python Example

After you complete all the software installations, you are ready to open the IDE to start working with TC-Python.

It is recommended that you open one or more of the included examples to both check that the installation has worked and to start familiarizing yourself with the code.

1.3.5.1 Open the TC-Python Project in PyCharm

When you first open the TC-Python project and examples, it can take a few moments for the Pycharm IDE to index before some of the options are available.

1. Open PyCharm and then choose **File**→**Open**. The first time you open the project you will need to navigate to the path of the TC-Python installation.

Operating sys-	Path to the TC-Python folder			
tem				
Windows	C:\Users\UserName\Documents\Thermo-Calc\2022b\SDK\			
	TC-Python			
Linux /home/UserName/Thermo-Calc/2022b/SDK/TC-Py				
MacOS /Users/Shared/Thermo-Calc/2022b/SDK/TC-Python				

- 2. Click on the Examples folder and then click OK.
- 3. From any subfolder:
 - Double-click to open an example file to examine the code.
 - Right-click an example and choose Run .

1.3.5.2 Fixing potential issues with the environment

In most cases you should run TC-Python within your **global** Python 3 interpreter and not use Virtual Environments unless you have a good reason to do so. A common problem on first usage of TC-Python is the error message "**No module named tc_python**". You can resolve this and other problems with the interpreter settings as follows:

- 1. Go the menu **File** \rightarrow **Settings**.
- 2. Navigate in the tree to **Project: YourProjectName** and choose **Project Interpreter**.
- 3. Click on the settings symbol close to the Project Interpreter dropdown menu and choose Add.
- 4. Now choose System Interpreter and add your existing Python 3 interpreter.
- 5. Select your added interpreter and confirm.

Note: If you are not following the recommended approach and create a *new* project (File \rightarrow New **Project...**), you need to consider that by default the options to choose the interpreter are hidden within the **Create Project** window. So click on **Project Interpreter:** New Virtual Environment and in most cases choose your *System Interpreter* instead of the default *New Virtual Environment*.

Note: If you really need to use a Virtual Environment, please consider the hints given in the *Python Virtual Environments* chapter.

1.4 Updating to a newer version

When updating to a newer version of Thermo-Calc, **you need to also install the latest version of TC-Python**. This is not necessary if are using the bundled Python-interpreter that has it automatically installed. It is not sufficient to run the installer of Thermo-Calc:

```
pip install <path to the TC-Python folder>/TC_Python-<version>-py3-none-any. {\hookrightarrow} whl
```

Tip: Note that on Linux depending on the interpreter usually *pip3* is used.

In case of problems you may wish to uninstall the previous version of TC-Python in advance:

However, that should normally not be required.

You can check the currently installed version of TC-Python by running:

pip show TC-Python

CHAPTER

TWO

MAC OS: SETTING ENVIRONMENT VARIABLES

In order to use TC-Python on Mac you need to set some environment variables.

TC22B_HOME=/Applications/Thermo-Calc-2022b.app/Contents/Resources

If you use a license server:

LSHOST=<name-of-the-license-server>

If you have a node-locked license:

LSHOST=NO-NET LSERVRC=/Users/Shared/Thermo-Calc/lservrc

In PyCharm, you can add environment variables in the configurations.

Select $Run \rightarrow Edit$ Configurations to open the Run/Debug Configurations window. Choose Templates and then Python. Enter the environment variable(s) by clicking the button to the right of the Environment Variables text field. Now the environment variables(s) will be set for each new configuration by default.

Note: Existing configurations need to be removed and recreated to obtain the environment variables in them.

The same way for configuring the environment variables can be used on other operating systems as if necessary.

CHAPTER

THREE

ARCHITECTURE OVERVIEW

TC-Python contains classes of these types:

- TCPython this is where you start with general settings.
- SystemBuilder and System where you choose database and elements etc.
- Calculation where you choose and configure the calculation.
- Result where you get the results from a calculation you have run.

3.1 TCPython

This is the starting point for all TC-Python usage.

You can think of this as the start of a "wizard".

You use it to select databases and elements. That will take you to the next step in the wizard, where you configure the system.

Example:

```
from tc_python import *
with TCPython() as start:
    start.select_database_and_elements(...
    # e.t.c
# after with clause
# or like this
with TCPython():
    SetUp().select_database_and_elements(...
    # e.t.c
# after with clause
```

Tip: If you use TC-Python from Jupyter Lab / Notebook, you should use TC-Python slightly different to be able to use multiple cells. See *Using TC-Python within a Jupyter Notebook or the Python console* for details.

Note: When your python script runs a row like this:

```
with TCPython() as start:
```

a process running a calculation server starts. Your code, via TC-Python, uses socket communication to send and receive messages to and from that server.

When your Python script has run as far as this row:

after with clause

the calculation server automatically shuts down, and all temporary files are deleted. It is important to ensure that this happens by structuring your Python code using a with () clause as in the above example.

Note: To re-use results from previous calculations, set a folder where TC-Python saves results, and looks for previous results.

This is done with the function set_cache_folder().

```
from tc_python import *
with TCPython() as start:
    start.set_cache_folder("cache")
```

This folder can be a network folder and shared by many users. If a previous TC-Python calculation has run with the same cache_folder and EXACTLY the same system and calculation settings, the calculation is not re-run. Instead the result is automatically loaded from disk.

It is also possible to explicitly save and load results.

```
from tc_python import *
with TCPython() as start:
    #... diffusion calculation (could be any calculation type)
    calculation_result.save_to_disk('path to folder')
    #...
    loaded_result = start.load_result_from_disk().diffusion('path to folder')
```

3.2 SystemBuilder and System

A SystemBuilder is returned when you have selected your database and elements in TCPython.

The **SystemBuilder** lets you further specify your system, for example the phases that should be part of your system.

Example:

```
from tc_python import *
with TCPython() as start:
    start.select_database_and_elements("ALDEMO", ["Al", "Sc"])
    # e.t.c
```

When all configuration is done, you call get_system() which returns an instance of a **System** class. The **System** class is fixed and cannot be changed. If you later want to change the database, elements or something else, change the **SystemBuilder** and call get_system() again, or create a new **SystemBuilder** and call get_system().

From the System you can create one or more calculations, which is the next step in the "wizard".

Note: You can use the same System object to create several calculations.

3.3 Calculation

The best way to see how a calculation can be used is in the TC-Python examples included with the Thermo-Calc installation.

Some calculations have many settings. Default values are used where it is applicable, and are overridden if you specify something different.

When you have configured your calculation you call calculate() to start the actual calculation. That returns a **Result**, which is the next step.

3.3.1 Single equilibrium calculations

In single equilibrium calculations you need to specify the correct number of conditions, depending on how many elements your **System** contains.

You do that by calling set_condition().

An important difference from other calculations is that single equilibrium calculations have two functions to get result values.

The calculate() method, which gives a **SingleEquilibriumTempResult**, is used to get actual values. This result is "temporary", meaning that if you run other calculations or rerun the current one, the resulting object no longer gives values corresponding to the first calculation.

This is different from how other calculations work. If you want a **Result** that you can use *after* running other calculations, you need to call calculate_with_state(), which returns a **SingleEquilibriumResult**.

Note: calculate() is the recommended function and works in almost all situations. Also it has *much* better performance than calculate_with_state().

3.3.2 Batch equilibrium calculations

Batch equilibrium calculations are used when you want to do many single equilibrium calculations and it is known from the beginning which result values are required from the equilibrium. This is a vectorized type of calculation that can reduce the overhead from Python and TC-Python similar to the approach used in *numpy*-functions for example.

Tip: The performance of batch equilibrium calculations can be significantly better than looping and using single equilibrium calculations **if the actual Thermo-Calc calculation is fast**. There is little advantage if the Thermo-Calc equilibrium calculations take a long time (typically for large systems and databases).

```
from tc_python import *
with TCPython() as start:
   calculation = (
        start
            .set_cache_folder(os.path.basename(__file__) + "_cache")
            .select_database_and_elements("NIDEMO", ["Ni", "Al", "Cr"])
            .get_system()
            .with_batch_equilibrium_calculation()
            .set_condition("T", 800.1)
            .set_condition("X(Al)", 1E-2)
            .set_condition("X(Cr)", 1E-2)
            .disable_global_minimization()
    )
    list_of_x_Al = [1, 2, 3, 4, 5, 6, 7, 8, 9, 10]
    list_of_x_Cr = [3, 5, 7, 9, 11, 13, 15]
    lists_of_conditions = []
    for x_Al in list_of_x_Al:
        for x_Cr in list_of_x_Cr:
            lists_of_conditions.append([
                ("X(Al)", x_Al / 100),
                ("X(Cr)", x_Cr / 100)])
    calculation.set_conditions_for_equilibria(lists_of_conditions)
    results = calculation.calculate(["BM", "VM"])
    masses = results.get_values_of("BM")
    volumes = results.get_values_of('VM')
print (masses)
print (volumes)
```

3.3.3 Precipitation calculations

All that can be configured in the *Precipitation Calculator* in Graphical Mode can also be done here in this calculation. However, you must at least enter a matrix phase, a precipitate phase, temperature, simulation time and compositions.

Example:

3.3.4 Scheil calculations

All Scheil calculations available in Graphical Mode or Console Mode can also be done here in this calculation. The minimum you need to specify are the elements and compositions. Everything else is set to a default value.

Example:

3.3.5 Property diagram calculations

For the property diagram (step) calculation, everything that you can configure in the *Equilibrium Calculator* when choosing *One axis* in Graphical Mode can also be configured in this calculation. In Console Mode the property diagram is created using the Step command. The minimum you need to specify are elements, conditions and the calculation axis. Everything else is set to default values, if you do not specify otherwise.

```
from tc python import *
with TCPython() as start:
   property_diagram = (
        start.
            select_database_and_elements("FEDEMO", ["Fe", "C"]).
            get_system().
            with_property_diagram_calculation().
                with_axis (CalculationAxis (ThermodynamicQuantity.temperature()).
                     set_min(500).
                     set_max(3000)).
                set_condition(ThermodynamicQuantity.mole_fraction_of_a_component("C"),
↔ 0.01).
                calculate().
                get_values_grouped_by_stable_phases_of(ThermodynamicQuantity.
\rightarrowtemperature(),
                                                         ThermodynamicQuantity.volume_

→ fraction_of_a_phase("ALL"))

        )
```

3.3.6 Phase diagram calculations

For the phase diagram (map) calculation, everything that you can configure in the *Equilibrium Calculator* when choosing *Phase diagram* in Graphical Mode can also be configured in this calculation. In Console Mode the phase diagram is created using the Map command. The minimum you need to specify are elements, conditions and two calculation axes. Everything else is set to default values, if you do not specify otherwise.

```
from tc_python import *
with TCPython() as start:
   phase_diagram = (
        start.
            select_database_and_elements("FEDEMO", ["Fe", "C"]).
            get_system().
            with_phase_diagram_calculation().
                with_first_axis(CalculationAxis(ThermodynamicQuantity.temperature()).
                    set_min(500).
                    set_max(3000)).
                with second axis (CalculationAxis (ThermodynamicQuantity.mole fraction
→of_a_component("C")).
                    set_min(0).
                    set_max(1)).
                set_condition(ThermodynamicQuantity.mole_fraction_of_a_component("C"),
↔ 0.01).
                calculate().
                get_values_grouped_by_stable_phases_of(ThermodynamicQuantity.mass_

→ fraction_of_a_component("C"),

                                                         ThermodynamicQuantity.
\rightarrowtemperature())
        )
```

3.3.7 Diffusion calculations

For diffusion calculations, everything that you can configure in the *Diffusion Calculator* can also be configured in this calculation. The minimum you need to specify are elements, temperature, simulation time, a region with a grid and width, a phase and an initial composition.

Example:

```
from tc_python import *
with TCPython() as start:
    diffusion_result = (
        start.
            select_thermodynamic_and_kinetic_databases_with_elements("FEDEMO",
↔ "MFEDEMO", ["Fe", "Ni"]).
            get_system().
            with_isothermal_diffusion_calculation().
                set_temperature(1400.0).
                set_simulation_time(108000.0).
                add_region(Region("Austenite").
                    set_width(1E-4).
                    with_grid(CalculatedGrid.linear().set_no_of_points(50)).
                    with_composition_profile(CompositionProfile().
                        add("Ni", ElementProfile.linear(10.0, 50.0))
                    ).
                add_phase("FCC_A1")).
        calculate())
    distance, ni_fraction = diffusion_result.get_mass_fraction_of_component_at_time(
→"Ni", 108000.0)
```

3.3.8 Property Model calculations

For Property Model calculations, everything that you can configure in the *Property Model Calculator* in Graphical Mode can also be configured in this calculation. The minimum you need to specify are elements, composition and which Property Model you want to use.

```
from tc_python import *
with TCPython() as start:
    print("Available Property Models: {}".format(start.get_property_models()))
    property_model = (
        start.
            select_database_and_elements("FEDEMO", ["Fe", "C"]).
            get_system().
            with_property_model_calculation("Driving force").
            set_composition("C", 1.0).
            set_argument("precipitate", "GRAPHITE"))

print("Available arguments: {}".format(property_model.get_arguments()))
result = property_model.calculate()

print("Available result quantities: {}".format(result.get_result_quantities()))
driving_force = result.get_value_of("normalizedDrivingForce")
```

3.3.9 Material to Material calculations

Material to Material calculations are generally regular single equilibrium, property diagram or phase diagram calculations but they are specialised to handle the mixture of two materials A and B. Everything that you can configure in the *Material to Material Calculator* in Graphical Mode can also be configured in this calculation. The minimum required configuration is shown below for a *Property diagram calculation for varying amount of material B*. The other calculators (single fraction of material B and phase diagram calculations) are configured in a similar way.

Example:

```
from tc_python import *
with TCPvthon() as start:
    material_to_material_property_diagram = (
        start.
            select_database_and_elements("FEDEMO", ["Fe", "Cr", "Ni", "C"]).
            qet_system().
            with_material_to_material().
            with_property_diagram_calculation().
            set material a({"Cr": 10.0, "Ni": 15.0}, "Fe").
            set_material_b({"Cr": 15.0, "Ni": 10.0}, "Fe").
            set_activities({"C": 0.1}).
            with_constant_condition(ConstantCondition.temperature(800 + 273.15)).
            with axis (MaterialToMaterialCalculationAxis.fraction_of_material_b(from_
\rightarrow fraction=0.0,
                                                                                   to
\rightarrow fraction=1.0,
                                                                                    start
\rightarrow fraction=0.5))
    )
    result = material_to_material_property_diagram.calculate()
    data = result.get_values_grouped_by_quantity_of (MATERIAL_B_FRACTION,
                                                       ThermodynamicQuantity.volume_
→ fraction_of_a_phase (ALL_PHASES))
    for group in data.values():
        fractions_of_b = group.x
        volume_fractions_of_phase = group.y
        phase_name = group.label
```

3.3.10 Process Metallurgy calculations

Process Metallurgy calculations are specialized to support the convenient handling of component-based additions (i.e., slag compositions such as 50% Al2O3 - 30% CaO - 20% SiO2), provide tailor-made result quantities, a framework for developing kinetic process simulations, and more useful features.

There are two distinct types of calculations:

- tc_python.process_metallurgy.equilibrium.EquilibriumCalculation: isothermal and adiabatic equilibrium calculations
- tc_python.process_metallurgy.process.ProcessSimulationCalculation: a kinetic process simulation framework, based on an *Effective Equilibrium Reaction Zone* (EERZ) approach

Equilibrium calculation example:

Equilibrium calculations are useful in a large range of situations when considering the kinetics of a process is unnecessary.

```
from tc_python import *
with TCPython() as session:
    metal = EquilibriumAddition({"Fe": None, "C": 4.5, "Si": 1.0}, 100e3,...
\rightarrowtemperature=1650 + 273.15)
    slag = EquilibriumAddition({"CaO": 75, "Al2O3": 25}, 3e3, temperature=1600 + 273.
→15)
    gas = EquilibriumGasAddition({"02": 100}, 1000, amount_unit=GasAmountUnit.NORM_
\hookrightarrow CUBIC_METER)
    calc = session.with_metallurgy().with_adiabatic_equilibrium_
→ calculation (ProcessDatabase.OXDEMO)
    (calc
     .add_addition(metal)
     .add_addition(slag)
     .add_addition(gas))
    result = calc.calculate()
    print(f"Stable phases: {result.get_stable_phases()}, temperature: {result.get_
\rightarrowtemperature() \} K")
```

Process simulation example:

TC-Python is providing a *framework* for modelling in principle any process in metallurgy, especially steel-making. It is up to the user to actually develop a concrete model for the process in question. The framework is in the current release limited to one reaction zone connecting two bulk zones. These bulk zones are typically the steel melt and the top slag, but not limited to that. The framework in its current version has proven to be useful to model industrial ladle furnaces, AOD- and VOD-converters and more. Process features such as heating and cooling, heat transfer between the bulk zones, inclusion formation and their flotation, etc., can be modelled.

This is a very simplified minimal but complete model mimicking a BOF process:

```
from tc_python import *
with TCPython() as session:
    calc = (session.with_metallurgy()
            .with_adiabatic_process_calculation(ProcessDatabase.OXDEMO)
            .set end time(15 * 60))
    steel_zone = MetalBulkZone(density=7800)
    slag_zone = SlagBulkZone(density=4500)
    steel_zone.add_addition(SingleTimeAddition({"Fe": None, "C": 4.5, "Si": 1.0},...
→120e3,
                                                temperature=1600 + 273.15), time=0)
    slag_zone.add_addition(SingleTimeAddition({"CaO": 75, "SiO2": 25}, 1.2e3,
                                               temperature=1500 + 273.15,
                                               composition_unit=CompositionUnit.MOLE_
\rightarrow PERCENT), time=0)
    steel_zone.add_continuous_addition(ContinuousGasAddition({"02": 100}, 1,
                                                               rate_unit=GasRateUnit.
→NORM_CUBIC_METER_PER_SEC))
```

```
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```

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3.4 Result

All calculations have a method called calculate() that starts the calculations and when finished, returns a **Result**.

The **Result** classes have very different methods, depending on the type of calculation.

The **Result** is used to get numerical values from a calculation that has run.

The **Result** can be saved to disk by the method save_to_disk().

Previously saved results can be loaded by the method load_result_from_disk() on the SetUp class.

Example:

```
# code above sets up the calculation
r = calculation.calculate()
time, meanRadius = r.get_mean_radius_of("AL3SC")
```

The **Result** objects are completely independent from calculations done before or after they are created. The objects return valid values corresponding to the calculation they were created from, for their lifetime. The only exception is if you call calculate() and not calculate_with_state() on a single equilibrium calculation.

As in the following example you can mix different calculations and results, and use old results after another calculation has run.

Example:

```
# ...
# some code to set up a single equilibrium calculation
# ...
single_eq_result = single_eq_calculation.calculate_with_state()
# ...
# some code to set up a precipitation calculation
# ...
prec_result = precipitation_calculation.calculate()
# ...
# some code to set up a Scheil calculation
# ...
```

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3.4.1 DiffusionResult

The DiffusionResult class, that is returned when calling calculate() on any DiffusionCalculation, has the possibility to create a ContinuedDiffusionCalculation, in addition to the "normal" functionality for results. This makes it possible to run a diffusion calculation and then, depending on the result, change some settings and continue.

Example:

```
# ...
# some code to set up a Diffusion calculation
# ...
first_diffusion_result = diffusion_calculation.calculate()
continued_calculation = first_diffusion_result.with_continued_calculation()
continued_calculation.set_simulation_time(110000.0)
continued_calculation.with_left_boundary_condition(BoundaryCondition.mixed_zero_flux_
...and_activity().set_activity_for_element('C', 1.0))
second_result = continued_calculation.calculate()
# ...
# Now you can use get second_result to get calculated values, just as normal.
# You can also use first_diffusion_result even after second_result is created.
# You can also use second_result (and even first_diffusion_result) to create a new_
...ContinuedDiffusionCalculation by calling with_continued_calculation.
```

3.5 Property Model Framework

The *Python Property Model SDK* extends the Thermo-Calc software to enable you to create your own Property Models. A *Property Model* is a Python-based calculation that can use any TC-Python functionality (including diffusion and precipitation calculations) but is usable through the Graphical User Interface (UI) of Thermo-Calc in a more simple way. It is typically used to model material properties but by no means limited to that. Examples of Property Models provided by Thermo-Calc include Martensite and Pearlite formation in steel.

The Property Model Framework uses standard Python 3 beginning with Thermo-Calc 2021a and can access all TC-Python functionality and any Python package including *numpy*, *scipy*, *tensorflow*, etc. The actual calculation code is nearly identical, regardless if called from within a Property Model or from standard Python.

This is a complete rewrite of the original version of the framework that was based on Jython 2.7 and therefore had a number of limitations. **Property models written with the old Property Model Framework before Thermo-Calc 2021a are not compatible with the new framework**. However, the migration should be relatively easy because the syntax was changed as little as possible.

3.5.1 Property models vs. TC-Python

The main difference between a *Property Model* and regular *TC-Python code* is that a Property Model is directly integrated into the UI of Thermo-Calc via a plugin architecture while TC-Python code can only be accessed by programs and scripts written in Python.

The user should develop a Property Model if the functionality needs to be available from the Thermo-Calc UI, especially if it should be applied by other users not familiar to programming languages. Otherwise it is preferable to implement the functionality directly in a TC-Python program. If required, Property Models can as well be accessed from within TC-Python.

3.5.2 Architecture

Every Property Model needs to contain a class that implements the interface *tc_python.propertymodel_sdk*. *PropertyModel*. There are naming conventions that must to be fulfilled: the file name is required to follow the the pattern XYPythonModel.py and the name of the class needs to match this. Additionally the file must be placed in a directory named XYPython within the Property Model directory. The content of the placeholder XY can be freely chosen.

A simple complete Property Model, saved in a file called SimplePythonModel.py in the directory SimplePython, looks like this:

```
from tc_python import *
class SimplePythonModel(PropertyModel):
   def provide_model_category(self) -> List[str]:
       return ["Demo"]
   def provide_model_name(self) -> str:
       return "My Demo Model"
   def provide_model_description(self) -> str:
       return "This is a demo model."
   def provide_ui_panel_components(self) -> List[UIComponent]:
       return [UIBooleanComponent("CHECKBOX", "Should this be checked?", "Simple_
def provide_calculation_result_quantities(self) -> List[ResultQuantity]:
       return [create_general_quantity("RESULT", "A result")]
   def evaluate_model(self, context: CalculationContext):
       if context.get_ui_boolean_value("CHECKBOX"):
           self.logger.info("The checkbox is checked")
        # obtain the entered values from the GUI
       composition_as_mass_fraction = context.get_mass_fractions()
       temp_in_k = context.get_temperature()
       calc = context.system.with_single_equilibrium_calculation()
        # continue with a TC-Python calculation now ...
       context.set_result_quantity_value("RESULT", 5.0) # the value would normally_
→ have been calculated
```

The basic building blocks of the Property Model API are:

- tc_python.propertymodel_sdk.ResultQuantity: Defines a calculation result of a Property Model that will be provided to the UI after each model evaluation
- tc_python.propertymodel_sdk.CalculationContext: Provides access to the data from the UI (such as the entered composition and temperature) and to the current TC-Python system object which is the entrypoint for using TC-Python from within the Property Model
- tc_python.propertymodel_sdk.UIComponent: These are the UI-components that create the user interface of the Property Model within the model panel of the Thermo-Calc application UI. Different components are available (for example checkboxes, text fields and lists).

3.5.3 Property Model directory

The Property Model *py*-files need to be located within subdirectories of the *Property Model directory*, e.g. PropertyModelDir/XYPython/XYPythonModel.py. The default Property Model directory can be changed in the menu *Tools -> Options* in the graphical user interface.

Operating system	Default Property Model directory
Windows	C:\Users\UserName\Documents\Thermo-Calc\2022b\PropertyModels
Linux	/home/UserName/Thermo-Calc/2022b/PropertyModels
MacOS	/Users/Shared/Thermo-Calc/2022b/PropertyModels

BEST PRACTICES

4.1 Re-use of the single equilibrium calculation state

The Thermo-Calc core keeps an internal state containing the data from previously performed calculations (such as composition of sublattices, previously formed phases, ...). This will be used for start values of future calculations (if not explicitly overwritten) and can strongly influence their convergence and calculation time. It can be useful to save and restore later the core-state **in advanced use cases**, these include:

- Improving the convergence speed in case of very complicated equilibria if a similar equilibrium had been calculated already before. Similarity refers here primarily to composition, temperature and entered phase set. This case can occur for example with the Nickel-database TCNi.
- Convenient and fast switching between states that have changed a lot (for example regarding suspended phases, numerical settings, ...)

The mechanism of saving and restoring the state is called bookmarking and is controlled with the two methods tc_python.single_equilibrium.SingleEquilibriumCalculation. bookmark_state() and tc_python.single_equilibrium.SingleEquilibriumCalculation. set_state_to_bookmark(). The following short example demonstrates how to switch between two different states easily in practice:

```
from tc_python import *
with TCPython() as session:
    calc = (session.
            select_database_and_elements("FEDEMO", ["Fe", "C"]).
            get_system().
            with_single_equilibrium_calculation().
            set_condition(ThermodynamicQuantity.temperature(), 2000.0).
            set_condition("X(C)", 0.01))
    calc.calculate()
   bookmark_temp_condition = calc.bookmark_state()
   calc.set_phase_to_fixed("BCC", 0.5)
    calc.remove_condition(ThermodynamicQuantity.temperature())
   bookmark_fixed_phase_condition = calc.bookmark_state()
    result_temp = calc.set_state_to_bookmark(bookmark_temp_condition)
    print ("Conditions do contain temperature: { }".format (result_temp.get_
\rightarrow conditions()))
    # this calculation had already been performed
    print("Stable phases (do not contain BCC): {}".format(result_temp.get_stable_
→phases()))
```

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```
result_fixed_phase = calc.set_state_to_bookmark(bookmark_fixed_phase_condition)
print("Conditions do not contain temperature: {}".format(result_fixed_phase.get_

conditions()))
# this calculation had **not yet** been performed
print("Stable phases (do contain BCC): {}".format(calc.calculate().get_stable_
cophases()))
```

4.2 Re-use and saving of results

Before a calculation is run in TC-Python, a check is made to see if the exact same calculation has run before, and if that is the case, the result from the calculation can be loaded from disk instead of being re-calculated.

This functionality is always enabled within a script running TC-Python, but you can make it work the same way when re-running a script, or even when running a completely different script.

To use results from previous calculations, set a folder where TC-Python saves results, and looks for previous results.

This is controlled by the method tc_python.server.SetUp.set_cache_folder().

```
from tc_python import *
with TCPython() as start:
    start.set_cache_folder("cache")
```

This folder can be a network folder and shared by many users. The calculation is not re-run if there is a previous TC-Python calculation with the same cache folder and exactly the same settings; the result is instead loaded from disk.

Another possibility is to explicitly save the result to disk and reload it later:

```
from tc_python import *
with TCPython() as start:
    # ... the system and calculator are set up and the calculation is performed
    result = calculator.calculate()
    result.save_to_disk("./result_dir")
```

You can then load the result again in another session:

```
from tc_python import *
with TCPython() as start:
    result = SetUp().load_result_from_disk().diffusion("./result_dir")
    x, frac = result.get_mole_fraction_of_component_at_time("Cr", 1000.0)
```

4.3 All TC-Python objects are non-copyable

Never create a copy of an instance of a class in TC-Python, neither by using the Python built-in function deepcopy() nor in any other way. All classes in TC-Python are proxies for classes in the underlying calculation server and normally hold references to result files. A copied class object in Python would consequently point to the same classes and result files in the calculation server.

Instead of making a copy, always create a new instance:

```
from tc_python import *
with TCPython() as start:
    system = start.select_database_and_elements("FEDEMO", ["Fe", "Cr"]).get_system()
    calculator = system.with_single_equilibrium_calculation()
    # *do not* copy the `calculator` object, create another one instead
    calculator_2 = system.with_single_equilibrium_calculation()
    # now you can use both calculators for different calculations ...
```

4.4 Python Virtual Environments

A Python installation can have several virtual environments. You can think of a virtual environment as a collection of third party packages that you have access to in your Python scripts. tc_python is such a package.

To run TC-Python, you need to **install it into the same virtual environment** as your Python scripts are running in. If your scripts fail on import tc_python, you need to execute the following command in the terminal of the same **Python environment** as your script is running in:

pip install TC_Python-<version>-py3-none-any.whl

If you use the PyCharm IDE, you should do that within the **Terminal** built into the IDE. This **Terminal** runs automatically within your actual (virtual) environment.

To prevent confusion, it is recommend in most cases to *install TC-Python within your global interpreter*, for example by running the pip install command within your default Anaconda prompt.

4.5 Using with TCPython() efficiently

Normally you should call with TCPython() only once within each process.

Note: When leaving the *with*-clause, the Java backend engine process is stopped and all temporary data is deleted. Finally when entering the next *with*-clause a new Java process is started. This can take several seconds.

If appropriate, it is safe to run *with TCPython()* in a loop. Due to the time it takes this only makes sense if the calculation time per iteration is longer than a minute.

To prevent calling with TCPython() multiple times and cleaning up temporary data, you can use the following pattern.

```
from tc python import *
# ...
def calculation(calculator):
   # you could also pass the `session` or `system` object if more appropriate
   calculator.set_condition("W(Cr)", 0.1)
    # further configuration ...
   result = calculator.calculate()
    # ...
    result.invalidate() # if the temporary data needs to be cleaned up immediately
if __name__ == '__main__':
    with TCPython() as session:
        system = session.select_database_and_elements("FEDEMO", ["Fe", "Cr"]).get_
\rightarrow system()
        calculator = system.with_single_equilibrium_calculation()
        for i in range(50):
            calculation (calculator)
```

4.6 Parallel calculations

It is possible to perform parallel calculations with TC-Python using multi-processing.

Note: Please note that **multi-threading is not suitable** for parallelization of computationally intensive tasks in Python. Additionally the Thermo-Calc core is not thread-safe. Using suitable Python-frameworks it is also possible to dispatch the calculations on different computers of a cluster.

A general pattern that can be applied is shown below. This code snippet shows how to perform single equilibrium calculations for different compositions in parallel. In the same way all other calculators of Thermo-Calc can be used or combined. For performance reasons in a real application, probably *numpy* arrays instead of Python arrays should be used.

Example:

```
import concurrent.futures
from tc_python import *

def do_perform(parameters):
    # this function runs within an own process
    with TCPython() as start:
        elements = ["Fe", "Cr", "Ni", "C"]
        calculation = (start.select_database_and_elements("FEDEMO", elements).
            get_system().
            with_single_equilibrium_calculation().
            set_condition("T", 1100).
            set_condition("W(C)", 0.1 / 100).
            set_condition("W(Ni)", 2.0 / 100))
```

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```
phase_fractions = []
        cr_contents = range(parameters["cr_min"],
                            parameters["cr_max"],
                            parameters["delta_cr"])
        for cr in cr_contents:
            result = (calculation.
                      set_condition("W(Cr)", cr / 100).
                      calculate())
            phase_fractions.append(result.get_value_of("NPM(BCC_A2)"))
    return phase_fractions
if __name__ == "__main__":
    parameters = [
        {"index": 0, "cr_min": 10, "cr_max": 15, "delta_cr": 1},
        {"index": 1, "cr_min": 15, "cr_max": 20, "delta_cr": 1}
    1
   bcc_phase_fraction = []
   num_processes = 2
    with concurrent.futures.ProcessPoolExecutor(num_processes) as executor:
        for result_from_process in zip(parameters, executor.map(do_perform,...
\rightarrow parameters)):
            # params can be used to identify the process and its parameters
            params, phase_fractions_from_process = result_from_process
            bcc_phase_fraction.extend(phase_fractions_from_process)
    # use the result in `bcc_phase_fraction`, for example for plotting
```

4.7 Handling crashes of the calculation engine

In some cases the Thermo-Calc calculation engine can crash. If batch calculations are performed, this brings down the complete batch. To handle this situation there is an exception you can use.

UnrecoverableCalculationException

That exception is thrown if the calculation server enters a state where no further calculations are possible. You should catch that exception outside of the *with TCPython()* clause and continue within a new *with*-clause.

Example:

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```
get_system().
                    with_isothermal_diffusion_calculation().
                        set_temperature(temperature).
                        set_simulation_time(108000.0).
                        add_region(Region("Austenite").
                            set_width(1E-4).
                            with_grid(CalculatedGrid.linear().set_no_of_points(50)).
                            with_composition_profile(CompositionProfile().
                                add("Ni", ElementProfile.linear(10.0, 50.0))
                            ).
                        add_phase("FCC_A1")).
               calculate())
           distance, ni_fraction = diffusion_result.get_mass_fraction_of_component_
→at_time("Ni", 108000.0)
           print(ni_fraction)
   except UnrecoverableCalculationException as e:
       print('Could not calculate. Continuing with next...')
```

4.8 Using TC-Python within a Jupyter Notebook or the Python console

TC-Python can also be used from within an interactive Jupyter Notebook and a Python console as well as similar products. The main difference from a regular Python program is that it is not recommended to use a *with*-clause to manage the TC-Python resources. That is only possible within a single Jupyter Notebook cell. Instead the standalone functions $tc_python.server.start_api_server()$ and $tc_python.server.stop_api_server()$ should be used for manually managing the resources.

Note: The *resources* of TC-Python are primarily the Java-process running on the backend side that performs the actual calculations and the temporary-directory of TC-Python that can grow to a large size over time, especially if precipitation calculations are performed. If a *with*-clause is used, these resources are automatically cleared after use.

You need to make sure that you execute the two functions tc_python.server.start_api_server() and tc_python.server.stop_api_server() exactly once within the Jupyter Notebook session. If not stopping TC-Python, extra Java-processes might be present and the temporary disk-space is not cleared. However, these issues can be resolved manually.

The temporary directories of TC-Python are named, for example, TC_TMP4747588488953835507 that has a random ID. The temporary directory on different operating systems varies according to the pattern shown in the table.

Operating sys-	Temporary directory				
tem					
Windows C:\Users{UserName}\AppData\Local\Temp\TC_TMP47475884889					
MacOS	/var/folders/g7/7du81ti_b7mm84n184fn3k910000lg/T/				
	TC_TMP4747588488953835507				
Linux	/tmp/TC_TMP4747588488953835507				

In a Jupyter Notebook some features of an IDE such as auto-completion (*TAB-key*), available method lookup (press. and then *TAB*) and parameter lookup (set the cursor within the method-parenthesis and press *SHIFT* + *TAB* or *SHIFT* + *TAB* + *TAB* for the whole docstring) are also available.

Example using TC-Python with a Jupyter Notebook:

Jupyter TC-Python with Jupyter Notebook Last Checkpoint: a few second	inds ago (autosaved)	Logout					
File Edit View Insert Cell Kernel Widgets Help	Trusted Python [conda env:An	aconda3] O					
E + ≥ 2 € ↑ ↓ NRun ■ C > Code							
In [1]: from tc_python import *							
<pre>In [2]: start_api_server()</pre>							
<pre>In [3]: system = SetUp().select_database_and_elements("FEDEMO",</pre>	["Fe", "Ni", "Cr"]).get_system()						
<pre>In [4]: temp = 825 # in K ni_conc = 10.0 # in wt-% cr_conc = 8.0 # in wt-%</pre>							
<pre>calc. \ set_condition(ThermodynamicQuantity.temperature(), set_condition(ThermodynamicQuantity.mass_fraction_of set_condition(ThermodynamicQuantity.mass_fraction_of result = calc.calculate()</pre>	f_a_component("Ni"), ni_conc / 100). \						
In [5]: result.get_value_of(ThermodynamicQuantity.mole_fraction]: result.get_value_of(ThermodynamicQuantity.mole_fraction_of_a_phase("FCC_A1"))						
Out[5]: 0.3345580340424432							
<pre>In [6]: stop_api_server()</pre>							

4.9 Property Model Framework

4.9.1 Debugging Property Model code

You can debug property models while running them from Thermo-Calc.

- Start Thermo-Calc and create a Property Model calculator.
- Select the model you want to debug and check the debug checkbox in the lower right corner of the Python code tab.

Dr	iving force				
Co	onfiguration	Description	Python code		
14				IIVI Matizeu_ui	TATHRTOLCE
75	ging mode	ι:			
76	c_python.	implementati	lonproperty	<pre>model_sdk import _PropertyModelP</pre>	roxy
77	rtyModelP	roxy(model=[)rivingForceP	ythonModel(), java_port=57334, p	<mark>ython_port=65247)</mark>
					🗹 Debug

Now the model that you want to debug has been updated with code needed to connect with Thermo-Calc.

• Start debugging the model in the IDE of your choice.

Note: You must use a Python interpreter where TC-Python is installed.

In PyCharm it looks like this:

_	🐉 DrivingForcePythonModel.py ×											
pe		1 from tc_python import *										
			Pcla					pertyModel):				
				RESL	JLT_QUANTITY		For					
				NORM	ALIZED_RESU	LT_QUANTITY		"normalizedDrivingForce				
				MATE	RIX_PHASE_LI	ST = "matrix						
				PREC	IPITATE_PHA	SE_LIST = "p	pre	cipitate"				
								Show Context Actions	Alt+Enter			
				def	provide_cal	culation_re	Ô.	<u>P</u> aste	Ctrl+V	ntity]:		
					results = [create_ener		Copy / Paste Special		ULT_QUANTITY,		
								Column Selection <u>M</u> ode	Alt+Shift+Insert	uantity.text")),		
						create_gene		Find <u>U</u> sages	Alt+F7	RMALIZED_RESULT_QUANTITY,		
								<u>R</u> efactor		izedResultQuantity.text"))]		
					return resu	lts		Folding				
								Go To	•			
				def	provide_mod			Generate	Alt+Insert			
					return [sel							
Ĩ								R <u>u</u> n 'DrivingForcePythonMo'	Ctrl+Shift+F10			
)		def	provide_mod			Debug 'DrivingForcePythonMo'	C			
	~ ~		Υ		1 70		V)	Run 'DrivingForcePythonMo' with	Co <u>v</u> erage			

Note: When your IDE and Thermo-Calc have successfully connected, you will see this in the Thermo-Calc log:

```
10:34:42,170 INFO Waiting for developer(!) to start Python process in debugger...
→DrivingForcePythonModel
10:34:42,171 INFO Connected successfully to the Python process for the model
→'DrivingForcePythonModel' in DEBUG mode
```

You can stop the debug session in your IDE, change the model code, and start debugging again. The changes you made will take effect in Thermo-Calc without the need to restart. If you for instance changed the method evaluate_model(), the change will take effect the next time you press *Perform*.

It is also possible to start the models from TC-Python. The workflow is exactly the same as described above, except instead of starting Thermo-Calc graphical user interface, you start a Python script and use the parameter debug_model=True when selecting your model.

```
from tc_python import *
```

```
with TCPython() as start:
    property_model = (
        start.
        select_database_and_elements("FEDEMO", ["Fe", "C"]).
        get_system().
        with_property_model_calculation("my own Driving Force", debug_model=True).
        set_composition("C", 1.0).
        )
    property_model.calculate()
....
```

4.9.2 Developing Property Models in several files

You can split your Property Model code in several .py files, and there are two ways of doing that:

- side-by-side modules
- common modules

Side-by-side modules are Python files located in the same folder as the Property Model.

Common modules are Python files located in a folder outside of the Property Model folder, which makes it possible to share them with several models as a common library.

4.9.2.1 side-by-side modules

You are required to:

- Add a __init__.py file to your Property Model folder
- Add all imports of *side-by-side* modules in your main Property Model Python file also to the __init__.py file

Example:

CriticalTemperaturesPythonModel.py (The main Property Model file):

```
from CriticalTemperaturesPython import CriticalTemperatures
from tc_python import *
import numpy as np
class CriticalTemperaturesPythonModel(PropertyModel):
    ...
```

__init__.py:

If you are using PyCharm, the package name of the Property Model might be highlighted as an error, in this case you can mark the Property Model directory (i.e. the root of the present model directory) by right-clicking on it in the project window of PyCharm and marking it as *Sources Root*:



critical_temperatures_library.py:

```
from tc_python import *
import numpy as np
from scipy import optimize
from enum import Enum
class CriticalTemperatures(object):
....
```

Note: Modules installed in the Python interpreter such as *numpy*, *scipy*, etc can be imported as normal. This only concerns files imported as *side-by-side* modules.

4.9.2.2 common modules

common modules work very similar to *side-by-side* modules except the import statements are done in the "main" _____init____.py file in *Property Model directory*.

You are required to:

- Add a _____init____.py file to your property model folder.
- Add all imports of *common* modules in your main property model python file also to both the __init__.py file in *Property Model directory* AND the __init__.py of the property model.

Example:

CriticalTemperaturesPythonModel.py (The main Property Model file):

```
from PropertyModels import Martensite
from tc_python import *
```

```
class CriticalTemperaturesPythonModel(PropertyModel):
    ...
```

___init___.py: (The init file located in the property model folder)

from PropertyModels import Martensite

___init___.py: (The init file located in *Property Model directory*)

from PropertyModels.common.martensite_library import Martensite

The file critical_temperatures_library.py should in this example be located in a folder called common in the *Property Model directory*.

critical_temperatures_library.py:

```
from tc_python import *
import numpy as np
from scipy import optimize
from enum import Enum
class CriticalTemperatures(object):
....
```

Note: *common* modules don't have to be located in folder called common. It can be any name, as long as the imports match the folder name.

Note: In this example the *Property Model directory* is called 'PropertyModels'. If you use a different directory for your property model, your imports have to match that.

4.9.3 Alternative Python for Property Models

4.9.3.1 Default bundled Python interpreter

Thermo-Calc is by default using a Python 3.7.2 interpreter bundled to the software for running the property models. It is containing the following major packages:

Package	Version
matplotlib	3.3.2
numpy	1.19.2
scikit-learn	0.23.2
scipy	1.5.2
TC-Python	2022b

Warning: Any changes to the interpreter packages can therefore break Thermo-Calc and should be avoided. If the installation gets broken, it can be fixed by reinstalling Thermo-Calc after having removed it.

Please contact the Thermo-Calc support if you think that further packages might be useful in future releases. If these packages are insufficient for you, it is possible to use another Python-interpreter: *Configuring another Python interpreter*.

The interpreter is located in different places depending on the platform:

Operating system	Path to the bundled Python-interpreter
Windows	C:\Program Files\Thermo-Calc\2022b\python\python.exe
Linux	/home/UserName/Thermo-Calc/2022b/python/bin/python3
MacOS	/Applications/Thermo-Calc-2022b.app/Contents/Resources/
	python/bin/python3

4.9.3.2 Configuring another Python interpreter

If you require additional Python-packages or prefer to use your own interpreter installed on your system, you can change the interpreter used by Thermo-Calc to run the property models. Select **Tools** \rightarrow **Options** in the Thermo-Calc GUI and modify the path to that of your Python 3 interpreter of choice:

Property model Python interpreter:	C:\Program Files\Thermo-Calc\2021a\python\python.exe	Modify

4.10 Process Metallurgy Calculations

4.10.1 Equilibrium calculations with changing elements between calculations

It is possible to add, change or remove additions after performing an equilibrium calculation using *tc_python*. *process_metallurgy.equilibrium.EquilibriumCalculation.calculate()*. This will change the elements being present in the system if the elements of the additions are differing. The Process Metallurgy Module will handle this situation by reloading the database with the latest set of elements. While this is an appropriate approach

in most cases, there can be some disadvantages: reloading the database takes some time and the internal engine state is lost, which may lead to successive calculations failures in some situations.

To avoid the database reload, it is possible to add the respective elements to additions being present in all calculations (with a zero-fraction):

```
from tc_python import *
with TCPython() as session:
    calc = session.with_metallurgy().with_adiabatic_equilibrium_
→ calculation (ProcessDatabase.OXDEMO)
    # add the element Al with zero-fraction already
    steel = EquilibriumAddition({'Fe': None, 'C': 4, 'Al': 0}, amount=100.0e3,__

→temperature=1700 + 273.15)

    slag = EquilibriumAddition({'CaO': 70, 'SiO2': 30}, amount=3.0e3,...
\rightarrowtemperature=1700 + 273.15)
    al_addition = EquilibriumAddition({'Al': 100}, amount=1.0e3)
    (calc
     .add_addition(steel)
     .add_addition(slag))
   result_1 = calc.calculate()
   calc.add_addition(al_addition)
    result_2 = calc.calculate()
    # evaluate the result as required ...
```

Or to add a later addition already before the first call to calculate() with a zero amount:

```
from tc_python import *
with TCPython() as session:
    calc = session.with_metallurgy().with_adiabatic_equilibrium_
→ calculation (ProcessDatabase.OXDEMO)
    steel = EquilibriumAddition({'Fe': None, 'C': 4}, amount=100.0e3,__

→temperature=1700 + 273.15)

    slag = EquilibriumAddition({'CaO': 70, 'SiO2': 30}, amount=3.0e3,_
\rightarrowtemperature=1700 + 273.15)
    # add the addition for now with zero-amount
    al_addition = EquilibriumAddition({'Al': 100}, amount=0)
    (calc
     .add_addition(al_addition)
     .add_addition(steel)
     .add_addition(slag))
   result_1 = calc.calculate()
   calc.update_addition(al_addition.set_amount(1.0e3))
   result_2 = calc.calculate()
    # evaluate the result as required ...
```

4.10.2 Zones

TC-Python is providing a *framework* for building time-dependent kinetic simulations of industrial and academic metallurgical processes where liquid phases are important. It is based on an *Effective Equilibrium Reaction Zone* (EERZ) approach which is separating a process into different zones. These zones have identical temperature and composition and are called *bulk zones*. Such zones can be in contact and react with each other by *reaction zones*. That means a *reaction zone* is modelling the interface between two *bulk zones*. One bulk zone is typically the steel melt and another bulk zone the top slag.

4.10.3 Applications

While this approach can in principle be extended to any number of zones, in the current release TC-Python is providing **only one reaction zone**. Practical work has however proven that this limitation is not critical for a lot of industrial processes, including ladle furnaces, AOD- and VOD-converters. Even more processes can be modelled with some limit of accuracy.

The reason for the power of the current implementation is that a number of important process features can be included:

- heating (tc_python.process_metallurgy.process.Zone.add_power())
- cooling (tc_python.process_metallurgy.process.Zone.add_power())
- heat transfer between bulk zones (tc_python.process_metallurgy.process.ReactionZone. add_heat_transfer())
- inclusion formation
- inclusion flotation and other transfer of phase groups between bulk zones (tc_python. process_metallurgy.process.ReactionZone.add_transfer_of_phase_group())
- addition of material and gas at any time in any zone (tc_python.process_metallurgy. process.Zone.add_addition() / tc_python.process_metallurgy.process.Zone. add_continuous_addition())
- an exhaust gas zone collecting all formed gas (tc_python.process_metallurgy.process. ProcessSimulationResult.get_exhaust_gas())
- time-dependent definition of most parameters (e.g., mass transfer coefficient, transfer of phase group, heating, etc.)

Please note that many of these features are called as well a *reaction zone* in other EERZ model implementations.

4.10.4 Implementation of practical process models

The Process Metallurgy Module has been successfully applied to a number of industrial processes.

Due to the broad range of industrial metallurgical processes, TC-Python is not providing ready-to-use models for certain processes. There are however examples available for common processes and this collection will be extended over time. The implementation of a model is an abstraction of the real process and should always be kept as simple as possible. Practical experience has proven that in many situations not more than one *reaction zone* is required.

The mass transfer coefficient is a fundamental parameter describing the kinetics in a *reaction zone* and is generally an empirical parameter. It depends however mostly on the geometry and stirring conditions in the process and not on the material compositions. Further on, the mass transfer coefficient has usually typical values for a given process - regardless of the actual furnace. That means that existing suggestions from the literature can be used as a starting point to derive the actual mass transfer coefficient for the process of interest.

CHAPTER

FIVE

API REFERENCE

5.1 Calculations

5.1.1 Module "single_equilibrium"

class tc_python.single_equilibrium.AbstractSingleEquilibriumCalculation(calculator)
 Bases: tc_python.abstract_base.AbstractCalculation

Abstract configuration required for a single equilibrium calculation.

Note: This is an abstract class that cannot be used directly.

```
disable_global_minimization()
```

Turns the global minimization completely off.

Returns This SingleEquilibriumCalculation object

enable_global_minimization()

Turns the global minimization on (using the default settings).

Returns This SingleEquilibriumCalculation object

get_components() \rightarrow List[str]

Returns a list of components in the system (including all components auto-selected by the database(s)).

Returns The components

$\texttt{get_gibbs_energy_addition_for}\ (\textit{phase: str}) \ \rightarrow \textit{float}$

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters phase – Specify the name of the (stoichiometric or solution) phase with the addition

Returns Gibbs energy addition to G per mole formula unit.

$\texttt{get_system_data()} \rightarrow \textit{tc_python.abstract_base.SystemData}$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

Parameters command - The Thermo-Calc Console Mode command

Returns This SingleEquilibriumCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

set_component_to_entered(component: str)

Sets the specified component to the status ENTERED, that is the default state.

Parameters component – The component name or ALL_COMPONENTS

Returns This SingleEquilibriumCalculation object

set_component_to_suspended(component: str, reset_conditions: bool = False)

Sets the specified component to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters

- **reset_conditions** if 'True' also remove composition conditions for the component if they are defined
- component The component name or ALL_COMPONENTS

Returns This SingleEquilibriumCalculation object

set_gibbs_energy_addition_for (phase: str, gibbs_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs_energy Addition to G per mole formula unit

Returns This SingleEquilibriumCalculation object

set_phase_to_dormant (phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This SingleEquilibriumCalculation object

set_phase_to_entered (phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

Parameters

- **phase** The phase name or *ALL_PHASES* for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

Returns This SingleEquilibriumCalculation object

set_phase_to_fixed (phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

Parameters

- phase The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

Returns This SingleEquilibriumCalculation object

set_phase_to_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This SingleEquilibriumCalculation object

with_options (*options*: tc_python.single_equilibrium.SingleEquilibriumOptions)

Sets the simulation options.

Parameters options – The simulation options

Returns This SingleEquilibriumCalculation object

with_reference_state (component: str, phase: str = 'SER', temperature: float = - 1.0, pressure:

float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

- component The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.

• **pressure** – The Pressure (in Pa) for the reference state.

Returns This SingleEquilibriumCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications) Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. user) databases loaded as a * .tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This SingleEquilibriumCalculation object

class tc_python.single_equilibrium.**SingleEquilibriumCalculation** (*calculator*) Bases: tc_python.single_equilibrium.AbstractSingleEquilibriumCalculation

Configuration for a single equilibrium calculation.

Note: Specify the conditions and possibly other settings, the calculation is performed with calculate().

bookmark_state (*bookmark_id: str* = '') \rightarrow str

Puts a "bookmark" on the current calculation-state of the calculator allowing the program to return to this state later as needed.

By bookmarking a state, you can simplify the convergence of equilibria when they strongly depend on the starting conditions (i.e. the state). Also use it to improve performance by running a calculation, then bookmarking it, and later returning to it for other equilibria whose conditions are "close" to the bookmarked equilibrium.

This method is used in combination with the method set_state_to_bookmark().

Parameters bookmark_id – The bookmark id. If omitted a generated id is used and returned

Returns The bookmark id

 $calculate() \rightarrow tc_python.single_equilibrium.SingleEquilibriumTempResult$

Performs the calculation and provides a temporary result object that is only valid until something gets changed in the calculation state. The method *calculate()* is the default approach and should be used in most cases.

Returns A new *SingleEquilibriumTempResult* object which can be used to get specific values from the calculated result. It is undefined behavior to use that object after the state of the calculation has been changed.

Warning: If the result object should be valid for the whole program lifetime, use calculate_with_state() instead.

 $calculate_with_state(timeout_in_minutes: float = 0.0) \rightarrow$

tc python.single equilibrium.SingleEquilibriumResult

Performs the calculation and provides a result object that reflects the present state of the calculation during the whole lifetime of the object.

Note: Because this method has performance and temporary disk space overhead (i.e. it is resource heavy), only use it when it is necessary to access the result object after the state is changed. In most cases you should use the method *calculate()*.

- **Parameters timeout_in_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.
- **Returns** A new *SingleEquilibriumResult* object which can be used later at any time to get specific values from the calculated result.

disable_global_minimization()

Turns the global minimization completely off.

Returns This SingleEquilibriumCalculation object

enable_global_minimization()

Turns the global minimization on (using the default settings).

Returns This SingleEquilibriumCalculation object

$\texttt{get_components()} \rightarrow List[str]$

Returns a list of components in the system (including all components auto-selected by the database(s)).

Returns The components

get_gibbs_energy_addition_for (*phase: str*) \rightarrow float

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters phase – Specify the name of the (stoichiometric or solution) phase with the addition

Returns Gibbs energy addition to G per mole formula unit.

get_interfacial_energy (*matrix_phase:* str, precipitate_phases: List[str], zero_volume_elements: List[str] = ['C', 'N']) \rightarrow Dict[str, float] Estimates the interfacial energy between a matrix phase and a precipitate phase using thermodynamic data from a CALPHAD database. The approximation model is based on Becker's bond energy approach.

Default: elements with no contribution to volume are C and N.

Parameters

- **matrix_phase** The matrix phase.
- **precipitate_phases** The list of precipitate phases for which interfacial energy between them and the matrix phase is to be calculated.
- **zero_volume_elements** The elements that are assumed to not contribute to the volume.

Returns A dictionary containing interfacial energy per precipitate phase.

$get_system_data() \rightarrow tc_python.abstract_base.SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

remove_all_conditions()

Removes all set conditions.

Returns This SingleEquilibriumCalculation object

- **remove_condition** (*quantity: Union*[tc_python.quantity_factory.ThermodynamicQuantity, *str*]) Removes the specified condition.
 - **Parameters quantity** the thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example "X(Cr)")

Returns This SingleEquilibriumCalculation object

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

Parameters command - The Thermo-Calc Console Mode command

Returns This SingleEquilibriumCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

set_component_to_entered(component: str)

Sets the specified component to the status ENTERED, that is the default state.

Parameters component – The component name or ALL_COMPONENTS

Returns This SingleEquilibriumCalculation object

set_component_to_suspended(component: str, reset_conditions: bool = False)

Sets the specified component to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters

- **reset_conditions** if 'True' also remove composition conditions for the component if they are defined
- component The component name or ALL_COMPONENTS

Returns This SingleEquilibriumCalculation object

Sets the specified condition.

Parameters

• **quantity** – The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example "X(Cr)")

• **value** – The value of the condition

Returns This SingleEquilibriumCalculation object

set_gibbs_energy_addition_for (phase: str, gibbs_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters

- **phase** Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs_energy Addition to G per mole formula unit

Returns This SingleEquilibriumCalculation object

set_phase_to_dormant (phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This SingleEquilibriumCalculation object

set_phase_to_entered (phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

Parameters

- **phase** The phase name or *ALL_PHASES* for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

Returns This SingleEquilibriumCalculation object

set_phase_to_fixed (phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

Parameters

- phase The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

Returns This SingleEquilibriumCalculation object

set_phase_to_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters phase – The phase name or ALL_PHASES for all phases

Returns This SingleEquilibriumCalculation object

set_state_to_bookmark (bookmark_id: str) \rightarrow tc_python.single_equilibrium.SingleEquilibriumTempResult Resets the calculation state to a previously bookmarked state.

After calling this method, the calculation behaves exactly as it would after the bookmarked calculation ran.

This method is used in combination with the method *bookmark_state()*.

Parameters bookmark_id – The bookmark id of the state to return to.

- **Returns** A new *SingleEquilibriumTempResult* object which can be used to get specific values from the calculated result. It is undefined behavior to use that object after the state of the calculation has been changed.
- with_options (options: tc_python.single_equilibrium.SingleEquilibriumOptions)
 Sets the simulation options.

Parameters options – The simulation options

Returns This SingleEquilibriumCalculation object

with_reference_state (component: str, phase: str = 'SER', temperature: float = - 1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

- **component** The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

Returns This SingleEquilibriumCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a * .tdb-file.

Parameters system_modifications - The system modification to be performed

Returns This SingleEquilibriumCalculation object

class tc_python.single_equilibrium.SingleEquilibriumOptions Bases: object

General simulation conditions for the thermodynamic calculations.

disable_approximate_driving_force_for_metastable_phases()

Disables the approximation of the driving force for metastable phases.

Default: Enabled

Note: When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use disable_approximate_driving_force_for_metastable_phases() to force the calculation to converge for the metastable phases.

Returns This *SingleEquilibriumOptions* object

disable_control_step_size_during_minimization()

Disables stepsize control during minimization (non-global).

Default: Enabled

Returns This SingleEquilibriumOptions object

disable_force_positive_definite_phase_hessian()

Disables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

Returns This SingleEquilibriumOptions object

enable_approximate_driving_force_for_metastable_phases()

Enables the approximation of the driving force for metastable phases.

Default: Enabled

Note: When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct. use disable approximate driving force for metastable phases () to force the calculation to converge for the metastable phases.

Returns This SingleEquilibriumOptions object

enable_control_step_size_during_minimization()

Enables stepsize control during normal minimization (non-global).

Default: Enabled

Returns This SingleEquilibriumOptions object

enable_force_positive_definite_phase_hessian()

Enables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

Returns This SingleEquilibriumOptions object

set_global_minimization_max_grid_points (max_grid_points: int = 2000)

Sets the maximum number of grid points in global minimization. Only applicable if global minimization is actually used.

Default: 2000 points

Parameters max_grid_points – The maximum number of grid points

Returns This SingleEquilibriumOptions object

set_max_no_of_iterations (max_no_of_iterations: int = 500)
Set the maximum number of iterations.

Default: max. 500 iterations

Note: As some models give computation times of more than 1 CPU second/iteration, this number is also used to check the CPU time and the calculation stops if 500 CPU seconds/iterations are used.

Parameters max_no_of_iterations – The max. number of iterations

Returns This SingleEquilibriumOptions object

set_required_accuracy (*accuracy: float = 1e-06*) Sets the required relative accuracy.

Default: 1.0E-6

Note: This is a relative accuracy, and the program requires that the relative difference in each variable must be lower than this value before it has converged. A larger value normally means fewer iterations but less accurate solutions. The value should be at least one order of magnitude larger than the machine precision.

Parameters accuracy – The required relative accuracy

Returns This *SingleEquilibriumOptions* object

set_smallest_fraction (smallest_fraction: float = 1e-12)

Sets the smallest fraction for constituents that are unstable.

It is normally only in the gas phase that you can find such low fractions.

The **default value** for the smallest site-fractions is 1E-12 for all phases except for IDEAL phase with one sublattice site (such as the GAS mixture phase in many databases) for which the default value is always as 1E-30.

Parameters smallest_fraction – The smallest fraction for constituents that are unstable

Returns This SingleEquilibriumOptions object

class tc_python.single_equilibrium.**SingleEquilibriumResult**(*result*) Bases: tc_python.abstract_base.AbstractResult

Result of a single equilibrium calculation, it can be evaluated using a Quantity or Console Mode syntax.

change_pressure (pressure: float)

Change the pressure and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with higher performance. The properties are calculated at the new pressure using the phase amount, temperature and composition of phases from the initial equilibrium. Use $get_value_of()$ to obtain them.

Parameters pressure – The pressure [Pa]

Returns This SingleEquilibriumCalculation object

change_temperature (temperature: float)

Change the temperature and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with high performance. The properties are calculated at the new temperature using the phase amount, pressure and composition of phases from the initial equilibrium. Use get_value_of() to obtain them.

Note: This is typically used when calculating room temperature properties (e.g. density) for a material when it is assumed that the equilibrium phase amount and composition freeze-in at a higher temperature during cooling.

Parameters temperature – The temperature [K]

Returns This SingleEquilibriumCalculation object

get_components() \rightarrow List[str]

Returns the names of the components selected in the system (including any components auto-selected by the database(s)).

Returns The names of the selected components

```
get_conditions () \rightarrow List[str]
Returns the conditions.
```

Returns The selected conditions

get_phases() \rightarrow List[str]

Returns the phases present in the system due to its configuration. It also contains all phases that have been automatically added during the calculation, this is the difference to the method System. get_phases_in_system().

Returns The names of the phases in the system including automatically added phases

get_stable_phases() \rightarrow List[str]

Returns the stable phases (i.e. the phases present in the current equilibrium).

Returns The names of the stable phases

get_value_of (quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str]) \rightarrow float Returns a value from a single equilibrium calculation.

Parameters quantity – The thermodynamic quantity to get the value of; a Console Mode syntax strings can be used as an alternative (for example "NPM(FCC_A1)")

Returns The requested value

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine. This affects only the state of the result object.

Parameters command - The Thermo-Calc Console Mode command

Returns This SingleEquilibriumCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

save_to_disk (path: str)

Saves the result to disk. Note that the result is a folder, containing potentially many files. The result can later be loaded with load_result_from_disk()

Parameters path – the path to the folder you want the result to be saved in. It can be relative or absolute.

Returns this SingleEquilibriumResult object

class tc_python.single_equilibrium.SingleEquilibriumTempResult(result)
 Bases: tc_python.abstract_base.AbstractResult

Result of a single equilibrium calculation that is only valid until something gets changed in the calculation state. It can be evaluated using a Quantity or Console Mode syntax.

Warning: Note that it is undefined behavior to use that object after something has been changed in the state of the calculation, this will result in an InvalidResultStateException exception being raised.

change_pressure (pressure: float)

Change the pressure and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with higher performance. The properties are calculated at the new pressure using the phase amount, temperature and composition of phases from the initial equilibrium. Use get_value_of() to obtain them.

Parameters pressure – The pressure [Pa]

Returns This SingleEquilibriumCalculation object

change_temperature (temperature: float)

Change the temperature and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with high performance. The properties are calculated at the new temperature using the phase amount, pressure and composition of phases from the initial equilibrium. Use get_value_of() to obtain them.

Note: This is typically used when calculating room temperature properties (e.g. density) for a material when it is assumed that the equilibrium phase amount and composition freeze-in at a higher temperature during cooling.

Parameters temperature – The temperature [K]

Returns This SingleEquilibriumCalculation object

get_components() \rightarrow List[str]

Returns the names of the components selected in the system (including any components auto-selected by the database(s)).

Returns The names of the selected components

Raises *InvalidResultStateException* – If something has been changed in the state of the calculation since that result object has been created

$\texttt{get_conditions}() \rightarrow List[str]$

Returns the conditions.

Returns List containing the selected conditions

Raises *InvalidResultStateException* – If something has been changed in the state of the calculation since that result object has been created

$\texttt{get_phases()} \rightarrow List[str]$

Returns the phases present in the system due to its configuration. It also contains all phases that have been automatically added during the calculation, this is the difference to the method System. get_phases_in_system().

Returns The names of the phases in the system including automatically added phases

Raises *InvalidResultStateException* – If something has been changed in the state of the calculation since that result object has been created

get_stable_phases() \rightarrow List[str]

Returns the stable phases (i.e. the phases present in the current equilibrium).

Returns The names of the stable phases

Raises *InvalidResultStateException* – If something has been changed in the state of the calculation since that result object has been created

- get_value_of (quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str]) \rightarrow float Returns a value from a single equilibrium calculation.
 - **Parameters quantity** The thermodynamic quantity to get the value of; a Console Mode syntax strings can be used as an alternative (for example "NPM(FCC_A1)")

Returns The requested value

Raises *InvalidResultStateException* – If something has been changed in the state of the calculation since that result object has been created

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

Parameters command – The Thermo-Calc Console Mode command

Returns This SingleEquilibriumCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

5.1.2 Module "batch_equilibrium"

class tc_python.batch_equilibrium.BatchEquilibriumCalculation(calculator)
 Bases: tc_python.abstract_base.AbstractCalculation

Configuration for a series of single equilibrium calculations performed in a vectorized fashion.

Note: Specify the conditions and call *calculate()*.

Tip: The performance of batch equilibrium calculations can be significantly better than looping and using SingleEquilibriumCalculation **if the actual Thermo-Calc calculation is fast**. There is little advantage if the Thermo-Calc equilibrium calculations take a long time (typically for large systems and databases).

calculate (quantities: List[Union[tc_python.quantity_factory.ThermodynamicQuantity, str]], logging_frequency: int = 10, timeout_in_minutes: float = 0.0) \rightarrow tc python.batch equilibrium.BatchEquilibriumResult

Runs the batch equilibrium calculation. The calculated *BatchEquilibriumResult* can then be queried for the values of the quantities specified.

Example:

>>> quantities = ['G', 'X(BCC)']

Parameters

- quantities A list of the quantities to be calculated.
- **logging_frequency** Determines how often logging should be done.
- timeout_in_minutes Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

Returns A *BatchEquilibriumResult* which later can be used to get specific values from the calculated result.

disable_global_minimization()

Turns the global minimization completely off.

Returns This BatchEquilibriumCalculation object

enable_global_minimization()

Turns the global minimization on (using the default settings).

Returns This BatchEquilibriumCalculation object

get_components() \rightarrow List[str]

Returns a list of components in the system (including all components auto-selected by the database(s)).

Returns The components

get_gibbs_energy_addition_for (*phase: str*) \rightarrow float

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters phase – Specify the name of the (stoichiometric or solution) phase with the addition

Returns Gibbs energy addition to G per mole formula unit.

$get_system_data() \rightarrow tc_python.abstract_base.SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

```
remove_all_conditions()
```

Removes all set conditions.

Returns This BatchEquilibriumCalculation object

remove_condition (*quantity: Union*[tc_python.quantity_factory.ThermodynamicQuantity, *str*]) Removes the specified condition.

Parameters quantity – the thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example "X(Cr)")

Returns This BatchEquilibriumCalculation object

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

Parameters command – The Thermo-Calc Console Mode command

Returns This BatchEquilibriumCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

set_component_to_entered(component: str)

Sets the specified component to the status ENTERED, that is the default state.

Parameters component - The component name or ALL_COMPONENTS

Returns This BatchEquilibriumCalculation object

set_component_to_suspended (component: str, reset_conditions: bool = False)

Sets the specified component to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters

- **reset_conditions** if 'True' also remove composition conditions for the component if they are defined
- component The component name or ALL_COMPONENTS

Returns This BatchEquilibriumCalculation object

Sets the specified condition.

Parameters

- **quantity** The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example "X(Cr)")
- **value** The value of the condition

Returns This BatchEquilibriumCalculation object

Set the conditions of the equilibria to be calculated.

This is done by sending a list of equilibria at once.

Each equilibrium itself is a list of conditions that will be changed for that equilibrium.

A condition is described by a tuple containing:

- 1. A Console Mode syntax string or a ThermodynamicQuantity instance,
- 2. A float value specifying the value of the condition.

Example:

>>> [[('T', 800), ('X(Cr)', 0.1)], [('T', 850), ('X(Cr)', 0.11)]]

You can use ThermodynamicQuantity instead of a Console Mode syntax string when specifying type of condition.

Example:

Parameters equilibria – The list of equilibria

Returns This BatchEquilibriumCalculation object

set_gibbs_energy_addition_for (phase: str, gibbs_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters

- **phase** Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs_energy Addition to G per mole formula unit

Returns This BatchEquilibriumCalculation object

set_phase_to_dormant (phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This BatchEquilibriumCalculation object

set_phase_to_entered (*phase: str, amount: float = 1.0*) Sets the phase to the status ENTERED, that is the default state.

Parameters

- **phase** The phase name or *ALL_PHASES* for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

Returns This BatchEquilibriumCalculation object

set_phase_to_fixed (phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

Parameters

• phase – The phase name

• **amount** – The fixed phase fraction (between 0.0 and 1.0)

Returns This BatchEquilibriumCalculation object

set_phase_to_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This BatchEquilibriumCalculation object

with_options (options: tc_python.single_equilibrium.SingleEquilibriumOptions)
 Sets the simulation options.

Parameters options – The simulation options

Returns This BatchEquilibriumCalculation object

with_reference_state (component: str, phase: str = 'SER', temperature: float = - 1.0, pressure:

float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

• **component** – The name of the element must be given.

- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

Returns This BatchEquilibriumCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications) Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a * .tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This BatchEquilibriumCalculation object

Bases: object

Result of a batch equilibrium calculation. This can be used to query for specific values.

 $\texttt{get_values_of}(\textit{quantity:} Union[tc_python.quantity_factory.ThermodynamicQuantity, str]) \rightarrow \texttt{figure}$

List[float] Returns values from a batch equilibrium calculation.

Warning: The quantity must be one of the quantities specified for the *BatchEquilibriumCalculation* object that created the result object.

Example:

```
>>> batch_result = batch_calculation.calculate(quantities = ['G', 'X(BCC)'])
>>> batch_result.get_values_of('G')
```

Parameters quantity – the thermodynamic quantity to get the value of; a Console Mode syntax strings can be used as an alternative (for example "NPM(FCC_A1)")

invalidate()

Invalidates the object and frees the disk space used by it.

Note: This is only required if the disk space occupied by the object needs to be released during the calculation. No data can be retrieved from the object afterwards.

5.1.3 Module "precipitation"

```
class tc_python.precipitation.FixedGrainSize(grain_radius: float = 0.0001)
Bases: tc_python.precipitation.GrainGrowthModel
```

set_grain_aspect_ratio (grain_aspect_ratio: float = 1.0)
Enter a numerical value. Default: 1.0.

Parameters grain_aspect_ratio – The grain aspect ratio [-]

class tc_python.precipitation.**GrainGrowth**(*grain_size_distribution:*

tc_python.precipitation.GrainSizeDistribution) Bases: tc_python.precipitation.GrainGrowthModel

disable_zener_pinning()

Disable Zener pinning to ignore the particle pinning effect on the grain growth. Zener pinning is by default disabled when no grain size distribution is defined, i.e. a single constant grain size is used. The setting is by default enabled when a grain size distribution is defined.

Returns This GrainSizeDistribution object

enable_zener_pinning()

Enable Zener pinning to simulate the particle pinning effect on the grain growth. The setting is by default enabled when a grain size distribution is defined.

Returns This GrainSizeDistribution object

set_grain_boundary_energy (energy: float = 0.5)

Set the energy of the grain bounday.

Parameters energy – The grain boundary energy [J/m2]

Returns This GrainSizeDistribution object

set_grain_boundary_mobility_activation_energy(activation_energy: float =

```
jioui
```

Set the grain boundary mobility activation energy where the mobility is defined by an Arrhenius type of equation.

242000.0)

Parameters activation_energy – The mobility activation energy [J/mol]

Returns This GrainSizeDistribution object

set_grain_boundary_mobility_pre_factor (pre_factor: float = 0.004)

Set the grain boundary mobility prefactor where the mobility is defined by an Arrhenius type of equation.

Parameters pre_factor – The grain boundary mobility pre factor [m⁴/(J s)]

Returns This GrainSizeDistribution object

class tc_python.precipitation.GrainGrowthModel

Bases: object

Factory class providing objects representing a grain growth model.

classmethod fixed_grain_size (*grain_radius: float = 0.0001*) Fixed grain radius size. **Default**: 1.0E-4 m

Parameters grain_radius – The grain radius / size [m]

classmethod grain_growth (*grain_size_distribution:* tc_python.precipitation.GrainSizeDistribution) Sets the initial grain size distribution for the matrix. **Default**: If the initial grain size distribution is not explicitly provided, a constant average grains size will be used and no grain growth evaluated during the simulation. Tip: Use this option if you want to study the further evolution of an existing microstructure.

Parameters grain_size_distribution – grain size distribution

class tc_python.precipitation.GrainSizeDistribution

Bases: object

Represents the grain size distribution at a certain time.

add_radius_and_number_density (*radius: float, number_density: float*) Adds a radius and number density pair to the grain size distribution.

Parameters

- **radius** The radius [m]
- number_density The number of grains per unit volume per unit length [m^-4]

Returns This GrainSizeDistribution object

class tc_python.precipitation.GrowthRateModel(value)

Bases: enum.Enum

Choice of the used growth rate model for a precipitate.

The most efficient model is the *Simplified model*, which is the default and applicable to most alloy systems under the assumption that either the supersaturation is small, or the alloying elements have comparable diffusivity. If all alloying elements are substitutional but they have remarkable diffusivity difference, e.g. in Al-Zr system, or if the diffusivity is strongly composition-dependent, the *General model* is preferred. If the supersaturation is high, and meanwhile there are fast-diffusing interstitial elements such as C, the *Advanced model* is more appropriate to capture the NPLE mechanism.

ADVANCED = 3

The **advanced model** has been proposed by *Chen, Jeppsson, and Ågren (CJA) (2008)* and calculates the velocity of a moving phase interface in multicomponent systems by identifying the operating tieline from the solution of the flux-balance equations. This model can treat both high supersaturation and cross-diffusion rigorously. Spontaneous transitions between different modes (LE and NPLE) of phase transformation can be captured without any ad-hoc treatment.

Note: Since it is not always possible to solve the flux-balance equations and it takes time, usage of a less rigorous but simple and efficient model is preferred if possible.

GENERAL = 5

The **general model** is based on the *Morral-Purdy* model, which follows the same quasi-steady state approximation as the *Simplified model*, but improves it by taking the cross-diffusion into account.

NPLE = 11

The **Non-Partitioning Local Equilibrium (NPLE) growth rate model** is only available for alloy systems where *Fe* is the major element and at least one interstitial element partitions into the precipitate phase. *This model is specifically designed to deal with the fast diffusion of interstitial elements (C, N, etc.) in Fe alloys.* Based on the *Simplified growth model*, it still holds a local equilibrium condition at the migrating interface. It chooses a tie-line under NPLE condition so that the u-fractions of all substitutional elements and minor interstitial elements in the precipitate phase are the same as those in the far-field matrix phase (i.e. the overall instantaneous matrix composition).

$PARA_EQ = 10$

The **para-equilibrium model** is only available for alloy systems where Fe is the major element and C is

the only interstitial element, which also partitions into the precipitate phase. The interstitial elements, e.g. C, N, etc., usually have remarkably faster diffusion rate than the substitutional elements. Meanwhile, they are assumed to have negligible volume contribution, and as a result the composition variables are replaced by u-fractions when interstitial elements are included in the system. This model is specifically designed to address the fast diffusion of C in Fe alloys. Based on the Simplified growth rate model it holds a paraequilibrium condition at the migrating interface. Contrary to the regular ortho-equilibrium condition state that assumes that all alloying elements are in equilibrium at the interface, the para-equilibrium assumes only equilibrium for C. The substitutional elements are immobile and thus have the same compositions (u-fractions) across the interface.

SIMPLIFIED = 2

The **simplified model** is based on the *advanced model* but avoids the difficulty of finding the operating tie-line and uses instead the tie-line across the bulk composition. **This is the default growth rate model**.

```
class tc_python.precipitation.MatrixPhase(matrix_phase_name: str)
```

```
Bases: object
```

The matrix phase in a precipitation calculation

add_precipitate_phase (*precipitate_phase*: tc_python.precipitation.PrecipitatePhase) Adds a precipitate phase.

```
Parameters precipitate_phase – The precipitate phase
```

set_dislocation_density (*dislocation_density: float = 5000000000000.0*) Enter a numerical value. **Default**: 5.0E12 m^-2.

Parameters dislocation_density – The dislocation density [m⁻²]

set_grain_aspect_ratio (grain_aspect_ratio: float = 1.0)
Enter a numerical value. Default: 1.0.

Note: Deprecated in version 2022a: Use *with_grain_growth_model()* instead. This method will be removed in release 2023a.

Parameters grain_aspect_ratio – The grain aspect ratio [-]

set_grain_radius (grain_radius: float = 0.0001)
Sets grain radius / size. Default: 1.0E-4 m

Note: Deprecated in version 2022a: Use *with_grain_growth_model()* instead. This method will be removed in release 2023a.

Parameters grain_radius - The grain radius / size [m]

A value that adds to the activation energy of mobility data from the database. Default: 0.0 J/mol

Parameters mobility_enhancement_activation_energy – The value that adds to the activation energy of mobility data from the database [J/mol].

set_mobility_enhancement_prefactor (*mobility_enhancement_prefactor: float = 1.0*) A parameter that multiplies to the mobility data from database. **Default**: 1.0

Parameters mobility_enhancement_prefactor – The mobility enhancement factor [-

set_molar_volume(volume: float)

Sets the molar volume of the phase.

Default: If not set, the molar volume is taken from the thermodynamic database (or set to 7.0e-6 m³/mol if the database contains no molar volume information).

Parameters volume – The molar volume [m³/mol]

with_elastic_properties_cubic (c11: float, c12: float, c44: float)

Sets the elastic properties to "cubic" and specifies the elastic stiffness tensor components. **Default**: if not chosen, the default is DISREGARD

Parameters

- c11 The stiffness tensor component c11 [GPa]
- c12 The stiffness tensor component c12 [GPa]
- c44 The stiffness tensor component c44 [GPa]

with_elastic_properties_disregard()

Set to disregard to ignore the elastic properties. Default: This is the default option

with_elastic_properties_isotropic (shear_modulus: float, poisson_ratio: float)
 Sets elastic properties to isotropic. Default: if not chosen, the default is DISREGARD

Parameters

- **shear_modulus** The shear modulus [GPa]
- poisson_ratio The Poisson's ratio [-]
- with_grain_growth_model (grain_growth_model: tc_python.precipitation.GrainGrowthModel)
 Sets the model for grain growth. Either fixed size or with a starting distribution

Default: Fixed grain radius size 1.0E-4 m

Parameters grain_growth_model - the grain growth model

class tc_python.precipitation.NumericalParameters

Bases: object

Numerical parameters

- set_max_overall_volume_change (max_overall_volume_change: float = 0.001)
 This defines the maximum absolute (not ratio) change of the volume fraction allowed during one time step.
 Default: 0.001
 - **Parameters max_overall_volume_change** The maximum absolute (not ratio) change of the volume fraction allowed during one time step [-]
- set_max_radius_points_per_magnitude (max_radius_points_per_magnitude: float = 200.0)
 Sets the maximum number of grid points over one order of magnitude in radius. Default: 200.0

Parameters max_radius_points_per_magnitude – The maximum number of grid points over one order of magnitude in radius [-]

set_max_rel_change_critical_radius (max_rel_change_critical_radius: float = 0.1)
Used to place a constraint on how fast the critical radium can vary, and thus put a limit on time step.
Default: 0.1

Parameters max_rel_change_critical_radius – The maximum relative change of the critical radius [-]

This parameter ensures accuracy for the evolution of effective nucleation rate. Default: 0.5

Parameters max_rel_change_nucleation_rate_log – The maximum logarithmic relative change of the nucleation rate [-]

set_max_rel_radius_change (max_rel_radius_change: float = 0.01)
The maximum value allowed for relative radius change in one time step. Default: 0.01

Parameters max_rel_radius_change - The maximum relative radius change in one time
 step [-]

Set a limit on the time step by controlling solute depletion or saturation, especially at isothermal stage. **Default**: 0.01

Parameters max_rel_solute_composition_change – The limit for the relative solute composition change [-]

set_max_time_step (max_time_step: float = 0.1)

The maximum time step allowed for time integration as fraction of the simulation time. Default: 0.1

Parameters max_time_step - The maximum time step as fraction of the simulation time [-]

set_max_time_step_during_heating (*max_time_step_during_heating: float = 1.0*) The upper limit of the time step that has been enforced in the heating stages. **Default**: 1.0 s

> Parameters max_time_step_during_heating - The maximum time step during heating [s]

Sets the maximum volume fraction of subcritical particles allowed to dissolve in one time step. **Default**: 0.01

Parameters max_volume_fraction_dissolve_time_step – The maximum volume fraction of subcritical particles allowed to dissolve in one time step [-]

set_min_radius_nucleus_as_particle (*min_radius_nucleus_as_particle: float = 5e-10*) The cut-off lower limit of precipitate radius. **Default**: 5.0E-10 m

Parameters min_radius_nucleus_as_particle – The minimum radius of a nucleus to be considered as a particle [m]

set_min_radius_points_per_magnitude (min_radius_points_per_magnitude: float = 100.0)
Sets the minimum number of grid points over one order of magnitude in radius. Default: 100.0

Parameters min_radius_points_per_magnitude – The minimum number of grid points over one order of magnitude in radius [-]

set_radius_points_per_magnitude (radius_points_per_magnitude: float = 150.0)
Sets the number of grid points over one order of magnitude in radius. Default: 150.0

Parameters radius_points_per_magnitude – The number of grid points over one order of magnitude in radius [-]

- **set_rel_radius_change_class_collision** (*rel_radius_change_class_collision: float = 0.5*) Sets the relative radius change for avoiding class collision. **Default**: 0.5
 - **Parameters rel_radius_change_class_collision** The relative radius change for avoiding class collision [-]

class tc_python.precipitation.ParticleSizeDistribution Bases: object

Represents the state of a microstructure evolution at a certain time including its particle size distribution, composition and overall phase fraction.

add_radius_and_number_density (radius: float, number_density: float)

Adds a radius and number density pair to the particle size distribution.

Parameters

- radius The radius [m]
- number_density The number of particles per unit volume per unit length [m^-4]

Returns This ParticleSizeDistribution object

set_initial_composition (*element_name: str*, *composition_value: float*) Sets the initial precipitate composition.

Parameters

- element_name The name of the element
- composition_value The composition value [composition unit defined for the calculation]

Returns This *ParticleSizeDistribution* object

set_volume_fraction_of_phase_type (volume_fraction_of_phase_type_enum:

tc_python.precipitation.VolumeFractionOfPhaseType)

Sets the type of the phase fraction or percentage. Default: By default volume fraction is used.

Parameters volume_fraction_of_phase_type_enum – Specifies if volume percent or fraction is used

Returns This ParticleSizeDistribution object

set_volume_fraction_of_phase_value(value: float)

Sets the overall volume fraction of the phase (unit based on the setting of set_volume_fraction_of_phase_type()).

Parameters value – The volume fraction 0.0 - 1.0 or percent value 0 - 100

Returns This *ParticleSizeDistribution* object

class tc_python.precipitation.PrecipitateElasticProperties Bases: object

Represents the elastic transformation strain of a certain precipitate class.

Note: This class is only relevant if the option *TransformationStrainCalculationOption*. *USER_DEFINED* has been chosen using *PrecipitatePhase.set_transformation_strain_calculation_opti* The elastic strain can only be considered for non-spherical precipitates.

```
set_e11 (e11: float)
```

Sets the elastic strain tensor component e11. Default: 0.0

Parameters e11 – The elastic strain tensor component e11

Returns This *PrecipitateElasticProperties* object

set_e12 (*e12: float*)

Sets the strain tensor component e12. Default: 0.0

Parameters e12 – The elastic strain tensor component e12

Returns This *PrecipitateElasticProperties* object

set_e13 (*e13: float*)

Sets the elastic strain tensor component e13. Default: 0.0

Parameters e13 – The elastic strain tensor component e13

Returns This *PrecipitateElasticProperties* object

set_e22 (*e22: float*)

Sets the elastic strain tensor component e22. Default: 0.0

Parameters e22 – The elastic strain tensor component e22

Returns This *PrecipitateElasticProperties* object

set_e23 (*e23: float*)

Sets the elastic strain tensor component e23. Default: 0.0

Parameters e23 – The elastic strain tensor component e23

Returns This PrecipitateElasticProperties object

set_e33 (*e33: float*)

Sets the elastic strain tensor component e33. Default: 0.0

Parameters e33 – The elastic strain tensor component e33

Returns This *PrecipitateElasticProperties* object

class tc_python.precipitation.**PrecipitateMorphology**(*value*) Bases: enum.Enum

Available precipitate morphologies.

CUBOID = 3

Cuboidal precipitates, only available for bulk nucleation.

NEEDLE = 1

Needle-like precipitates, only available for bulk nucleation.

PLATE = 2

Plate-like precipitates, only available for bulk nucleation.

SPHERE = 0

Spherical precipitates, this is the default morphology.

class tc_python.precipitation.PrecipitatePhase(precipitate_phase_name: str)

Bases: object

Represents a certain precipitate class (i.e. a group of precipitates with the same phase and settings).

disable_calculate_aspect_ratio_from_elastic_energy()

Disables the automatic calculation of the aspect ratio from the elastic energy of the phase.

Returns This PrecipitatePhase object

Note: If you use this method, you are required to set the aspect ratio explicitly using the method set_aspect_ratio_value().

Default: This is the default setting (with an aspect ratio of 1.0).

disable_driving_force_approximation()

Disables driving force approximation for this precipitate class. **Default**: Driving force approximation is disabled.

Returns This *PrecipitatePhase* object

enable_calculate_aspect_ratio_from_elastic_energy()

Enables the automatic calculation of the aspect ratio from the elastic energy of the phase. **Default**: The aspect ratio is set to a value of 1.0.

Returns This *PrecipitatePhase* object

enable_driving_force_approximation()

Enables driving force approximation for this precipitate class. This approximation is often required when simulating precipitation of multiple particles that use the same phase description. E.g. simultaneous precipitation of a Metal-Carbide(MC) and Metal-Nitride(MN) if configured as different composition sets of the same phase FCC_A1. **Default**: Driving force approximation is disabled.

Returns This *PrecipitatePhase* object

Tip: Use this if simulations with several compositions sets of the same phase cause problems.

set_alias (alias: str)

Sets an alias string that can later be used to get values from a calculated result. Typically used when having the same phase for several precipitates, but with different nucleation sites. For example two precipitates of the phase M7C3 with nucleation sites in 'Bulk' and at 'Dislocations'. The alias can be used instead of the phase name when retrieving simulated results.

Parameters alias – The alias string for this class of precipitates

Returns This *PrecipitatePhase* object

Note: Typically used when having using the same precipitate phase, but with different settings in the same calculation.

```
set_aspect_ratio_value(aspect_ratio_value: float)
```

Sets the aspect ratio of the phase. Default: An aspect ratio of 1.0.

Parameters aspect_ratio_value – The aspect ratio value

Returns This *PrecipitatePhase* object

Note: Only relevant if *disable_calculate_aspect_ratio_from_elastic_energy()* is used (which is the default).

set_gibbs_energy_addition(gibbs_energy_addition: float)

Sets a Gibbs energy addition to the Gibbs energy of the phase. Default: 0,0 J/mol

Parameters gibbs_energy_addition - The Gibbs energy addition [J/mol]

Returns This *PrecipitatePhase* object

set_interfacial_energy(interfacial_energy: float)

Sets the interfacial energy. **Default**: If the interfacial energy is not set, it is automatically calculated using a broken-bond model.

```
Parameters interfacial_energy – The interfacial energy [J/m^2]
```

Returns This PrecipitatePhase object

Note: The calculation of the interfacial energy using a broken-bond model is based on the assumption of an interface between a bcc- and a fcc-crystal structure with (110) and (111) lattice planes regardless of the actual phases.

set_interfacial_energy_estimation_prefactor(interfacial_energy_estimation_prefactor:

float)

Sets the interfacial energy prefactor. **Default**: Prefactor of 1.0 (only relevant if the interfacial energy is automatically calculated).

Parameters interfacial_energy_estimation_prefactor – The prefactor for the calculated interfacial energy

Returns This *PrecipitatePhase* object

Note: The interfacial energy prefactor is an amplification factor for the automatically calculated interfacial energy. Example: *interfacial_energy_estimation_prefactor* = $2.5 \Rightarrow 2.5 \approx$ calculated interfacial energy

set_molar_volume (volume: float)

Sets the molar volume of the precipitate phase. **Default**: The molar volume obtained from the database. If no molar volume information is present in the database, a value of 7.0e-6 m^3/mol is used.

Parameters volume – The molar volume [m³/mol]

Returns This *PrecipitatePhase* object

set_nucleation_at_dislocations (number_density=- 1)

Activates nucleation at dislocations for this class of precipitates. Calling the method overrides any other nucleation setting for this class of precipitates. **Default**: If not set, by default bulk nucleation is chosen.

Parameters number_density – Number density of nucleation sites. If not set, the value is calculated based on the matrix settings (grain size, dislocation density) [m^-3].

Returns This *PrecipitatePhase* object

set_nucleation_at_grain_boundaries (wetting_angle: float = 90.0, number_density: float =

Activates nucleation at grain boundaries for this class of precipitates. Calling the method overrides any other nucleation setting for this class of precipitates. **Default**: If not set, by default bulk nucleation is chosen.

Parameters

- wetting_angle If not set, a default value of 90 degrees is used
- **number_density** Number density of nucleation sites. If not set, the value is calculated based on the matrix settings (grain size) [m⁻³].

Returns This *PrecipitatePhase* object

set_nucleation_at_grain_corners (wetting_angle: float = 90, number_density: float = -1)
Activates nucleation at grain corners for this class of precipitates. Calling the method overrides any other
nucleation setting for this class of precipitates. Default: If not set, by default bulk nucleation is chosen.

Parameters

• wetting_angle – If not set, a default value of 90 degrees is used]

• **number_density** – Number density of nucleation sites. If not set, the value is calculated based on the matrix settings (grain size) [m⁻³].

Returns This PrecipitatePhase object

set_nucleation_at_grain_edges (wetting_angle=90, number_density=-1)

Activates nucleation at the grain edges for this class of precipitates. Calling the method overrides any other nucleation setting for this class of precipitates. **Default**: If not set, by default bulk nucleation is chosen.

Parameters

- wetting_angle If not set, a default value of 90 degrees is used
- **number_density** Number density of nucleation sites. If not set, the value is calculated based on the matrix settings (grain size) [m⁻³].

Returns This *PrecipitatePhase* object

set_nucleation_in_bulk (number_density: float = - 1.0)

Activates nucleation in the bulk for this class of precipitates. Calling the method overrides any other nucleation setting for this class of precipitates. **Default**: This is the default setting (with an automatically calculated number density).

Parameters number_density – Number density of nucleation sites. If not set, the value is calculated based on the matrix settings (molar volume) [m⁻³]

Returns This *PrecipitatePhase* object

set_phase_boundary_mobility (*phase_boundary_mobility: float*) Sets the phase boundary mobility. **Default**: 10.0 m^4/(Js).

Parameters phase_boundary_mobility – The phase boundary mobility [m⁴/(Js)]

Returns This *PrecipitatePhase* object

set_precipitate_morphology (precipitate_morphology_enum: tc_python.precipitateMorphology)
Sets the precipitate morphology. Default: PrecipitateMorphology.SPHERE

Parameters precipitate_morphology_enum – The precipitate morphology

Returns This PrecipitatePhase object

set_transformation_strain_calculation_option (transformation_strain_calculation_option_enum:

tc_python.precipitation.TransformationStrainCalculationOption Sets the transformation strain calculation option. **Default**: *TransformationStrainCalculationOption*. *DISREGARD*.

Parameters transformation_strain_calculation_option_enum - The chosen option

Returns This *PrecipitatePhase* object

with_elastic_properties (elastic_properties: tc_python.precipitation.PrecipitateElasticProperties)
Sets the elastic properties. Default: The elastic transformation strain is disregarded by default.

Parameters elastic_properties – The elastic properties object

Returns This PrecipitatePhase object

Note: This method has only an effect if the option *TransformationStrainCalculationOption*. *USER_DEFINED* is chosen using the method *set_transformation_strain_calculation_option()*.

with_growth_rate_model (growth_rate_model_enum: tc_python.precipitation.GrowthRateModel)
Sets the growth rate model for the class of precipitates. Default: GrowthRateModel.SIMPLIFIED

Parameters growth_rate_model_enum - The growth rate model

Returns This *PrecipitatePhase* object

with_particle_size_distribution (particle_size_distribution:

tc_python.precipitation.ParticleSizeDistribution)

Sets the initial particle size distribution for this class of precipitates. **Default**: If the initial particle size distribution is not explicitly provided, the simulation will start from a supersaturated matrix.

Parameters particle_size_distribution - The initial particle size distribution object

Returns This *PrecipitatePhase* object

Tip: Use this option if you want to study the further evolution of an existing microstructure.

```
class tc_python.precipitation.PrecipitationCCTCalculation(calculation)
Bases: tc_python.abstract_base.AbstractCalculation
```

Configuration for a Continuous-Cooling-Time (CCT) precipitation calculation.

calculate (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.precipitation.PrecipitationCalculationTTTorCCTResult* Runs the CCT diagram calculation.

Parameters timeout_in_minutes – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

Returns A *PrecipitationCalculationTTTorCCTResult* which later can be used to get specific values from the calculated result

get_system_data() \rightarrow tc_python.abstract_base.SystemData

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with_system_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

set_composition (element_name: str, value: float)

Sets the composition of the elements. The unit for the composition can be changed using set_composition_unit(). Default: Mole percent (CompositionUnit.MOLE_PERCENT)

Parameters

- element_name The element
- value The composition (fraction or percent depending on the composition unit)

Returns This *PrecipitationCCTCalculation* object

set_composition_unit (unit_enum: tc_python.utils.CompositionUnit)

Sets the composition unit. Default: Mole percent (CompositionUnit.MOLE_PERCENT).

Parameters unit_enum – The new composition unit

Returns This PrecipitationCCTCalculation object

set_cooling_rates (cooling_rates: List[float])

Sets all cooling rates for which the CCT diagram should be calculated.

Parameters cooling_rates – A list of cooling rates [K/s]

Returns This *PrecipitationCCTCalculation* object

set_max_temperature (*max_temperature: float*) Sets maximum temperature of the CCT diagram.

Parameters max_temperature – the maximum temperature [K]

Returns This PrecipitationCCTCalculation object

set_min_temperature (min_temperature: float)
 Sets the minimum temperature of the CCT diagram.

Parameters min_temperature – the minimum temperature [K]

Returns This PrecipitationCCTCalculation object

- stop_at_volume_fraction_of_phase (stop_criterion_value: float)
 Sets the stop criterion as a volume fraction of the phase. This setting is applied to all phases.
 - **Parameters** stop_criterion_value the volume fraction of the phase (a value between 0 and 1)

Returns This PrecipitationCCTCalculation object

with_matrix_phase (matrix_phase: tc_python.precipitation.MatrixPhase)
 Sets the matrix phase.

Parameters matrix_phase – The matrix phase

Returns This PrecipitationCCTCalculation object

with_numerical_parameters (*numerical_parameters*: tc_python.precipitation.NumericalParameters) Sets the numerical parameters. If not specified, reasonable defaults are be used.

Parameters numerical_parameters – The parameters

Returns This *PrecipitationCCTCalculation* object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)
 Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. user) databases loaded as a *.tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This *PrecipitationCCTCalculation* object

class tc_python.precipitation.PrecipitationCalculationResult(result)
 Bases: tc_python.abstract_base.AbstractResult

Result of a precipitation calculation. This can be used to query for specific values.

save_to_disk (path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load_result_from_disk()

Parameters path – the path to the folder you want the result to be saved in. It can be relative or absolute.

Returns this PrecipitationCalculationResult object

class tc_python.precipitation.PrecipitationCalculationSingleResult(result)
 Bases: tc python.precipitation.PrecipitationCalculationResult

Result of a isothermal or non-isothermal precipitation calculation. This can be used to query for specific values.

Search the Thermo-Calc help for definitions of the axis variables, e.g. search *isothermal variables* or *non-isothermal variables*.

Returns the aspect ratio distribution of a precipitate in dependency of its mean particle length at a certain time.

Only available if the morphology is set to *PrecipitateMorphology.NEEDLE* or *PrecipitateMorphology.PLATE*.

Parameters

- time The time [s]
- precipitate_id The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (mean particle length [m], aspect ratio)

get_aspect_ratio_distribution_for_radius_of (*precipitate_id: str, time: float*) → [typing.List[float], typing.List[float]]

Returns the aspect ratio distribution of a precipitate in dependency of its mean radius at a certain time.

Only available if the morphology is set to *PrecipitateMorphology.NEEDLE* or *PrecipitateMorphology.PLATE*.

Parameters

- time The time [s]
- **precipitate_id** The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (mean radius [m], aspect ratio)

get_critical_radius_of (*precipitate_id: str*) \rightarrow [typing.List[float], typing.List[float]] Returns the critical radius of a precipitate in dependency of the time.

Parameters precipitate_id – The id of a precipitate can either be phase name or alias

Returns A tuple of two lists of floats (time [s], critical radius [m])

<pre>get_cubic_factor_distribution_for_particle_length_of (precipitate_id:</pre>		str,	
time: float) -	\rightarrow	[typ-	
ing.List[float],		typ-	
ing.List[float]]			

Returns the cubic factor distribution of a precipitate in dependency of its mean particle length at a certain time.

Only available if the morphology is set to *PrecipitateMorphology*. CUBOID.

Parameters

- time The time in seconds
- precipitate_id The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (particle length [m], cubic factor)

get_cubic_factor_distribution_for_radius_of (precipitate_id: str, time: float) → [typing.List[float], typing.List[float]]

Returns the cubic factor distribution of a precipitate in dependency of its mean radius at a certain time. Only available if the morphology is set to *PrecipitateMorphology*.*CUBOID*.

Parameters

• time – The time [s]

• precipitate_id - The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (radius [m], cubic factor)

get_driving_force_of (*precipitate_id: str*) \rightarrow [typing.List[float], typing.List[float]] Returns the (by R * T) normalized driving force of a precipitate in dependency of the time.

Parameters precipitate_id – The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (time [s], normalized driving force)

get_grain_critical_radius () \rightarrow [typing.List[float], typing.List[float]] Returns the critical radius of grains in dependency of the time.

Returns A tuple of two lists of floats (time [s], critical radius [m])

get_grain_mean_radius () \rightarrow [typing.List[float], typing.List[float]] Returns the mean grain size of the matrix phase in dependency of the time.

Returns A tuple of two lists of floats (time [s], mean radius [m])

get_grain_number_density() \rightarrow [typing.List[float], typing.List[float]] Returns the grain number density in dependency of the time.

Returns A tuple of two lists of floats (time [s], grain number density [m⁻³])

get_grain_number_density_distribution_for_length(time: float) → [typing.List[float], typing.List[float]]

Returns the number density distribution of grains in dependency of its mean particle length at a certain time.

Parameters time – The time [s]

Returns A tuple of two lists of floats (grain length[m], number of grains per unit volume per unit length [m^-4])

<pre>get_grain_number_density_distribution_for_radius(time:</pre>	float)	\rightarrow	[typ-
ing.List[float],		typ-
ingList	float]]		

Returns the number density distribution of a grains in dependency of its mean radius at a certain time.

Parameters time – The time [s]

Returns A tuple of two lists of floats (radius [m], number of grains per unit volume per unit length [m⁻⁴])

```
get_grain_size_distribution (time: float) \rightarrow [typing.List[float], typing.List[float]]
Returns the size distribution of the matrix phase in dependency of its grain radius length at a certain time.
```

Parameters time – The time [s]

Returns A tuple of two lists of floats (grain radius[m], number density of grains[m^-3])

get_matrix_composition_in_mole_fraction_of(*element_name:* str) → [typ-ing.List[float], typing.List[float]]

Returns the matrix composition (as mole fractions) of a certain element in dependency of the time.

Parameters element_name - The element

Returns A tuple of two lists of floats (time [s], mole fraction)

get_matrix_composition_in_weight_fraction_of (*element_name:* str) \rightarrow [typing.List[float], typing.List[float]]

Returns the matrix composition (as weight fraction) of a certain element in dependency of the time.

Parameters element_name – The element

Returns A tuple of two lists of floats (time [s], weight fraction)

get_mean_aspect_ratio_of (*precipitate_id: str*) \rightarrow [typing.List[float], typing.List[float]] Returns the mean aspect ratio of a precipitate in dependency of the time.

Only available if the morphology is set to *PrecipitateMorphology.NEEDLE* or *PrecipitateMorphology.PLATE*.

Parameters precipitate_id – The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (time [s], mean aspect ratio)

- get_mean_cubic_factor_of (precipitate_id: str) → [typing.List[float], typing.List[float]]
 Returns the mean cubic factor of a precipitate in dependency of the time. Only available if the morphology
 is set to PrecipitateMorphology.CUBOID.
 - **Parameters** precipitate_id The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (time [s], mean cubic factor)

- get_mean_particle_length_of (*precipitate_id: str*) \rightarrow [typing.List[float], typing.List[float]] Returns the mean particle length of a precipitate in dependency of the time.
 - Only available if the morphology is set to *PrecipitateMorphology.NEEDLE* or *PrecipitateMorphology.PLATE*.
 - **Parameters** precipitate_id The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (time [s], mean particle length [m])

get_mean_radius_of (*precipitate_id: str*) \rightarrow [typing.List[float], typing.List[float]] Returns the mean radius of a precipitate in dependency of the time.

Parameters precipitate_id – The id of a precipitate can either be phase name or alias

Returns A tuple of two lists of floats (time [s], mean radius [m])

 $\begin{array}{rcl} \texttt{get_normalized_grain_size_distribution} (\textit{time: float}) & \rightarrow & [\texttt{typing.List[float]}, & \texttt{typing.List[float]}] \end{array}$

Returns the normalized number density distribution of a grains at a certain time.

Parameters time – The time [s]

Returns A tuple of two lists of floats (Normalized size, Frequency)

Returns the normalized number density distribution of a precipitate at a certain time.

Parameters

- time The time [s]
- precipitate_id The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (Normalized size, Frequency)

- **get_nucleation_rate_of** (*precipitate_id: str*) \rightarrow [typing.List[float], typing.List[float]] Returns the nucleation rate of a precipitate in dependency of the time.
 - **Parameters precipitate_id** The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (time [s], nucleation rate [m⁻³ s⁻¹)

get_number_density_distribution_for_particle_length_of (precipitate_id: str,

```
time: float) \rightarrow
```

```
[typing.List[float],
```

typing.List[float]]

Returns the number density distribution of a precipitate in dependency of its mean particle length at a certain time.

Parameters

- time The time [s]
- precipitate_id The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (particle length[m], number of particles per unit volume per unit length [m^-4])

Returns the number density distribution of a precipitate in dependency of its mean radius at a certain time.

Parameters

- time The time [s]
- precipitate_id The id of a precipitate can either be the phase name or an alias
- **Returns** A tuple of two lists of floats (radius [m], number of particles per unit volume per unit length [m^-4])

```
get_number_density_of (precipitate_id: str) \rightarrow [typing.List[float], typing.List[float]]
Returns the particle number density of a precipitate in dependency of the time.
```

Parameters precipitate_id – The id of a precipitate can either be phase name or alias

Returns A tuple of two lists of floats (time [s], particle number density [m^-3])

```
get_precipitate_composition_in_mole_fraction_of (precipitate_id: str, ele-
```

ment_name: str) \rightarrow [typ-

ing.List[float], typing.List[float]]

Returns the precipitate composition (as mole fractions) of a certain element in dependency of the time.

Parameters

- precipitate_id The id of a precipitate can either be phase name or alias
- element_name The element

Returns A tuple of two lists of floats (time [s], mole fraction)

<pre>get_precipitate_composition_in_weight_fraction_of(precipitate_id:</pre>	str,	el-
ement_name:	str)	\rightarrow
[typing.List[float],		typ-
ing.List[float]]		

Returns the precipitate composition (as weight fraction) of a certain element in dependency of the time.

Parameters

- **precipitate_id** The id of a precipitate can either be phase name or alias
- **element_name** The element

Returns A tuple of two lists of floats (time [s], weight fraction)

get_size_distribution_for_particle_length_of (precipitate_id: str, time: float) → [typing.List[float], typing.List[float]]

Returns the size distribution of a precipitate in dependency of its mean particle length at a certain time.

Parameters

- time The time [s]
- precipitate_id The id of a precipitate can either be the phase name or an alias
- **Returns** A tuple of two lists of floats (particle length[m], number of particles per unit volume per unit length [m^-4])
- $\label{eq:get_size_distribution_for_radius_of} (precipitate_id: str, time: float) \rightarrow [typ-ing.List[float], typing.List[float]]$

Returns the size distribution of a precipitate in dependency of its mean radius at a certain time.

Parameters

- time The time [s]
- precipitate_id The id of a precipitate can either be the phase name or an alias
- **Returns** A tuple of two lists of floats (radius [m], number of particles per unit volume per unit length [m^-4])
- **get_volume_fraction_of** (*precipitate_id: str*) \rightarrow [typing.List[float], typing.List[float]] Returns the volume fraction of a precipitate in dependency of the time.
 - **Parameters** precipitate_id The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (time [s], volume fraction)

class tc_python.precipitation.PrecipitationCalculationTTTorCCTResult (result)
 Bases: tc_python.precipitation.PrecipitationCalculationResult

Result of a TTT or CCT precipitation calculation.

- get_result_for_precipitate (*precipitate_id: str*) \rightarrow [typing.List[float], typing.List[float]] Returns the calculated data of a TTT or CCT diagram for a certain precipitate.
 - **Parameters** precipitate_id The id of a precipitate can either be the phase name or an alias

Returns A tuple of two lists of floats (time [s], temp [K])

class tc_python.precipitation.**PrecipitationIsoThermalCalculation**(*calculation*) Bases: tc_python.abstract_base.AbstractCalculation

Configuration for an isothermal precipitation calculation.

- **calculate** (*timeout_in_minutes:* float = 0.0) \rightarrow tc_python.precipitation.PrecipitationCalculationSingleResult Runs the isothermal precipitation calculation.
 - **Parameters timeout_in_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.
 - **Returns** A *PrecipitationCalculationSingleResult* which later can be used to get specific values from the calculated result

$get_system_data() \rightarrow tc_python.abstract_base.SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

set_composition (element_name: str, value: float)

Sets the composition of the elements. The unit for the composition can be changed using set_composition_unit(). Default: Mole percent (CompositionUnit.MOLE_PERCENT)

Parameters

- element_name The element
- value The composition (fraction or percent depending on the composition unit)

Returns This *PrecipitationIsoThermalCalculation* object

set_composition_unit (unit_enum: tc_python.utils.CompositionUnit = <CompositionUnit.MOLE_PERCENT: 2>)

Sets the composition unit. Default: Mole percent (CompositionUnit.MOLE_PERCENT).

Parameters unit_enum – The new composition unit

Returns This *PrecipitationIsoThermalCalculation* object

set_simulation_time (simulation_time: float)

Sets the simulation time.

Parameters simulation_time – The simulation time [s]

 $Returns \ This \ {\it Precipitation} IsoThermalCalculation \ object$

set_temperature(temperature: float)

Sets the temperature for the isothermal simulation.

Parameters temperature – the temperature [K]

Returns This PrecipitationIsoThermalCalculation object

with_matrix_phase (matrix_phase: tc_python.precipitation.MatrixPhase)
 Sets the matrix phase.

Parameters matrix_phase – The matrix phase

Returns This PrecipitationIsoThermalCalculation object

with_numerical_parameters (*numerical_parameters*: tc_python.precipitation.NumericalParameters) Sets the numerical parameters. If not specified, reasonable defaults are be used.

Parameters numerical_parameters – The parameters

Returns This PrecipitationIsoThermalCalculation object

with_system_modifications (*system_modifications:* tc_python.abstract_base.SystemModifications) Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a * .tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This PrecipitationIsoThermalCalculation object

class tc_python.precipitation.**PrecipitationNonIsoThermalCalculation**(*calculation*) Bases: tc_python.abstract_base.AbstractCalculation

Configuration for a non-isothermal precipitation calculation.

- **calculate** (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.precipitation.PrecipitationCalculationSingleResult* Runs the non-isothermal precipitation calculation.
 - **Parameters timeout_in_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.
 - **Returns** A *PrecipitationCalculationSingleResult* which later can be used to get specific values from the calculated result

 $get_system_data() \rightarrow tc_python.abstract_base.SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

set_composition (element_name: str, value: float)

Sets the composition of the elements. The unit for the composition can be changed using set_composition_unit(). Default: Mole percent (CompositionUnit.MOLE_PERCENT)

Parameters

- element_name The element
- value The composition (fraction or percent depending on the composition unit)

Returns This PrecipitationIsoThermalCalculation object

set_composition_unit (unit_enum: tc_python.utils.CompositionUnit)

Sets the composition unit. Default: Mole percent (CompositionUnit.MOLE_PERCENT).

Parameters unit_enum – The new composition unit

Returns This PrecipitationIsoThermalCalculation object

set_simulation_time (simulation_time: float)
Sets the simulation time.

Parameters simulation_time – The simulation time [s]

Returns This PrecipitationNonThermalCalculation object

with_matrix_phase (matrix_phase: tc_python.precipitation.MatrixPhase)
 Sets the matrix phase.

Parameters matrix_phase – The matrix phase

Returns This PrecipitationIsoThermalCalculation object

with_numerical_parameters (*numerical_parameters*: tc_python.precipitation.NumericalParameters) Sets the numerical parameters. If not specified, reasonable defaults are be used.

Parameters numerical_parameters - The parameters

Returns This PrecipitationIsoThermalCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications) Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a * .tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This PrecipitationNonThermalCalculation object

- with_temperature_profile (temperature_profile: tc_python.utils.TemperatureProfile)
 Sets the temperature profile to use with this calculation.
 - **Parameters temperature_profile** the temperature profile object (specifying time / temperature points)

Returns This PrecipitationNonThermalCalculation object

class tc_python.precipitation.**PrecipitationTTTCalculation**(*calculation*) Bases: tc_python.abstract_base.AbstractCalculation

Configuration for a TTT (Time-Temperature-Transformation) precipitation calculation.

- **calculate** (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.precipitation.PrecipitationCalculationTTTorCCTResult* Runs the TTT diagram calculation.
 - **Parameters timeout_in_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

Returns A *PrecipitationCalculationTTTorCCTResult* which later can be used to get specific values from the calculated result.

get_system_data() \rightarrow tc_python.abstract_base.SystemData

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

set_composition (element_name: str, value: float)

Sets the composition of the elements. The unit for the composition can be changed using set_composition_unit(). Default: Mole percent (CompositionUnit.MOLE_PERCENT)

Parameters

• element_name - The element

• value – The composition (fraction or percent depending on the composition unit)

Returns This PrecipitationTTTCalculation object

set_composition_unit (unit_enum: tc_python.utils.CompositionUnit)

Sets the composition unit. Default: Mole percent (CompositionUnit.MOLE_PERCENT).

Parameters unit_enum – The new composition unit

Returns This PrecipitationTTTCalculation object

set_max_annealing_time (max_annealing_time: float)

Sets the maximum annealing time, i.e. the maximum time of the simulation if the stopping criterion is not reached.

Parameters max_annealing_time – the maximum annealing time [s]

Returns This PrecipitationTTTCalculation object

set_max_temperature (max_temperature: float)

Sets the maximum temperature for the TTT diagram.

Parameters max_temperature – the maximum temperature [K]

Returns This PrecipitationTTTCalculation object

set_min_temperature (min_temperature: float)

Sets the minimum temperature for the TTT diagram.

Parameters min_temperature – the minimum temperature [K]

Returns This PrecipitationTTTCalculation object

set_temperature_step (temperature_step: float)

Sets the temperature step for the TTT diagram. If not set, the default value is 10 K.

Parameters temperature_step – the temperature step [K]

Returns This PrecipitationTTTCalculation object

stop_at_percent_of_equilibrium_fraction (percentage: float)

Sets the stop criterion to a percentage of the overall equilibrium phase fraction, alternatively a required volume fraction can be specified (using *stop_at_volume_fraction_of_phase()*).

Parameters percentage – the percentage to stop at (value between 0 and 100)

Returns This PrecipitationTTTCalculation object

stop_at_volume_fraction_of_phase(volume_fraction: float)

Sets the stop criterion as a volume fraction of the phase, alternatively a required percentage of the equilibrium phase fraction can be specified (using stop_at_percent_of_equilibria_fraction()). Stopping at a specified volume fraction is the default setting.

This setting is applied to all phases.

Parameters volume_fraction – the volume fraction to stop at (a value between 0 and 1)

Returns This *PrecipitationTTTCalculation* object

with_matrix_phase (matrix_phase: tc_python.precipitation.MatrixPhase)
 Sets the matrix phase.

Parameters matrix_phase – The matrix phase

Returns This PrecipitationTTTCalculation object

with_numerical_parameters (*numerical_parameters*: tc_python.precipitation.NumericalParameters) Sets the numerical parameters. If not specified, reasonable defaults are be used.

Parameters numerical_parameters – The parameters

Returns This PrecipitationTTTCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)
 Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. user) databases loaded as a *.tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This PrecipitationTTTCalculation object

class tc_python.precipitation.TransformationStrainCalculationOption(value)
 Bases: enum.Enum

Options for calculating the transformation strain.

CALCULATE_FROM_MOLAR_VOLUME = 2

Calculates the transformation strain from the molar volume, obtains a purely dilatational strain.

DISREGARD = 1

Ignores the transformation strain, **this is the default setting**.

```
USER_DEFINED = 3
```

Transformation strain to be specified by the user.

class tc_python.precipitation.VolumeFractionOfPhaseType(value)

Bases: enum.Enum

Unit of the volume fraction of a phase.

```
VOLUME_FRACTION = 6
Volume fraction (0 - 1), this is the default.
```

VOLUME_PERCENT = 5

Volume percent (0% - 100%).

5.1.4 Module "scheil"

```
class tc_python.scheil.CalculateSecondaryDendriteArmSpacing
    Bases: tc_python.scheil.ScheilBackDiffusion
```

Configures a secondary dendrite arm spacing calculation used by Scheil *with back diffusion*. The used equation is $c * cooling_rate^{(-n)}$ with c and n being provided either by the user or taken from the defaults.

set_c (*c*: *float* = 5*e*-05)

Sets the scaling factor c in the governing equation c * cooling_rate^(-n).

Default: 50 µm

Parameters c – The scaling factor [m]

Returns This CalculateSecondaryDendriteArmSpacing object

set_cooling_rate (*cooling_rate: float* = 1.0) Sets the cooling rate.

Default: 1.0 K/s

An increased value moves the result from equilibrium toward a Scheil-Gulliver calculation.

Parameters cooling_rate - The cooling rate [K/s]

Returns This CalculateSecondaryDendriteArmSpacing object

set_fast_diffusing_elements(element_names: List[str])

Sets elements as fast diffusing. This allows redistribution of these elements in both the solid and liquid parts of the alloy.

Default: No fast-diffusing elements.

Parameters element_names - The elements

Returns This CalculateSecondaryDendriteArmSpacing object

$set_n(n: float = 0.33)$

Sets the exponent n in the governing equation c * cooling_rate^(-n).

Default: 0.33

Parameters n – The exponent [-]

Returns This CalculateSecondaryDendriteArmSpacing object

set_primary_phasename (primary_phase_name: str = 'AUTOMATIC')

Sets the name of the primary phase.

The primary phase is the phase where the back diffusion takes place. If *AUTOMATIC* is selected, the program tries to find the phase which will give the most back diffusion. That behavior can be overridden by selecting a specific primary phase.

Default: AUTOMATIC

Parameters primary_phase_name – The phase name (or AUTOMATIC)

Returns This CalculateSecondaryDendriteArmSpacing object

class tc_python.scheil.ConstantSecondaryDendriteArmSpacing(secondary_dendrite_arm_spacing:

float = 5e-05)

Bases: tc_python.scheil.ScheilBackDiffusion

Configures a constant secondary dendrite arm spacing used by Scheil *with back diffusion*. The secondary dendrite arm spacing can either be provided by the user or taken from the defaults.

set_cooling_rate (*cooling_rate: float = 1.0*) Sets the cooling rate.

Default: 1.0 K/s

An increased value moves the result from equilibrium toward a Scheil-Gulliver calculation.

Parameters cooling_rate – The cooling rate [K/s]

Returns This ConstantSecondaryDendriteArmSpacing object

set_fast_diffusing_elements (element_names: List[str])

Sets elements as fast diffusing. This allows redistribution of these elements in both the solid and liquid parts of the alloy.

Default: No fast-diffusing elements.

Parameters element_names - The elements

Returns This ConstantSecondaryDendriteArmSpacing object

set_primary_phasename (*primary_phase_name: str = 'AUTOMATIC'*) Sets the name of the primary phase.

The primary phase is the phase where the back diffusion takes place. If *AUTOMATIC* is selected, the program tries to find the phase which will give the most back diffusion. That behavior can be overridden by selecting a specific primary phase.

Default: AUTOMATIC

Parameters primary_phase_name – The phase name (or AUTOMATIC)

Returns This ConstantSecondaryDendriteArmSpacing object

class tc_python.scheil.ScheilBackDiffusion

Bases: tc_python.scheil.ScheilCalculationType

Configuration for back diffusion in the solid primary phase.

Warning: This feature has only effect on systems with diffusion data (typically a mobility database). If used for a system without diffusion data, a normal Scheil calculation is done.

classmethod calculate_secondary_dendrite_arm_spacing()

Calculate the secondary dendrite arm spacing based on the following equation: $c * cooling_rate^{(-n)}$ with c and n being provided either by the user or taken from the defaults.

Use the methods provide by CalculateSecondaryDendriteArmSpacing to configure the parameters.

Returns A CalculateSecondaryDendriteArmSpacing

classmethod constant_secondary_dendrite_arm_spacing(secondary_dendrite_arm_spacing:

float = 5e-05)

Assuming constant secondary dendrite arm spacing, provided either by the user or taken from the defaults.

Default: 50 µm

Parameters secondary_dendrite_arm_spacing - The dendrite arm spacing [m]

Returns A ConstantSecondaryDendriteArmSpacing

class tc_python.scheil.ScheilCalculation(calculator)

Bases: tc_python.abstract_base.AbstractCalculation

Configuration for a Scheil solidification calculation.

Note: Specify the settings, the calculation is performed with *calculate()*.

calculate (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.scheil.ScheilCalculationResult* Runs the Scheil calculation.

Warning: Scheil calculations do not support the GAS phase being selected, this means the *GAS phase must always be deselected in the system* if it is present in the database

- **Parameters timeout_in_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.
- **Returns** A *ScheilCalculationResult* which later can be used to get specific values from the simulation.

disable_global_minimization()

Disables global minimization.

Default: Enabled

Note: When enabled, a global minimization test is performed when an equilibrium is reached. This costs more computer time but the calculations are more robust.

Returns This ScheilCalculation object

enable_global_minimization()

Enables global minimization.

Default: Enabled

Note: When enabled, a global minimization test is performed when an equilibrium is reached. This costs more computer time but the calculations are more robust.

Returns This ScheilCalculation object

get_system_data() \rightarrow tc_python.abstract_base.SystemData

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with_system_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

set_composition (component_name: str, value: float)

Sets the composition of a component. The unit for the composition can be changed using set_composition_unit().

Default: Mole percent (CompositionUnit.MOLE_PERCENT)

Parameters

• component_name - The component

• **value** – The composition value [composition unit defined for the calculation]

Returns This ScheilCalculation object

set_composition_unit (unit_enum: tc_python.utils.CompositionUnit = <CompositionUnit.MOLE_PERCENT: 2>)

Sets the composition unit.

Default: Mole percent (CompositionUnit.MOLE_PERCENT).

Parameters unit_enum – The new composition unit

Returns This ScheilCalculation object

set_start_temperature (*temperature_in_kelvin: float = 2500.0*) Sets the start temperature.

Warning: The start temperature needs to be higher than the liquidus temperature of the alloy.

Default: 2500.0 K

Parameters temperature_in_kelvin – The temperature [K]

Returns This ScheilCalculation object

with_calculation_type (scheil_calculation_type: tc_python.scheil.ScheilCalculationType)

Chooses a specific Scheil calculation. ClassicScheil for only setting fast diffusers, ScheilBackDiffusion enables back diffusion in the solid primary phase and optionally fast diffusers in all solid phases, and ScheilSoluteTrapping enables solute trapping in the solid primary phase. :param scheil_type: Type of Scheil calculation, either ScheilClassic, ScheilBackDiffusion or ScheilSoluteTrapping :return: This ScheilCalculation object

with_options (options: tc_python.scheil.ScheilOptions)

Sets the Scheil simulation options.

Parameters options – The Scheil simulation options

Returns This ScheilCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)
 Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. user) databases loaded as a *.tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This ScheilCalculation object

class tc_python.scheil.ScheilCalculationResult (result) Bases: tc python.abstract base.AbstractResult

Result of a Scheil calculation.

get_values_grouped_by_quantity_of (x_quantity: Union[tc_python.quantity_factory.ScheilQuantity, str], y_quantity: Union[tc_python.quantity_factory.ScheilQuantity,

str], $sort_and_merge: bool = True) \rightarrow Dict[str,$ $tc_python.utils.ResultValueGroup]$ Returns x-y-line data grouped by the multiple datasets of the specified quantities (for example in de-

pendency of phases or components). Use $get_values_of()$ instead if you need no separation. The available quantities can be found in the documentation of the factory class ScheilQuantity.

Note: The different datasets might contain *NaN*-values between different subsections and might not be sorted **even if the flag `sort_and_merge` has been set** (because they might be unsortable due to their nature).

Parameters

- **x_quantity** The first Scheil quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example "T")
- **y_quantity** The second Scheil quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example "NV")
- **sort_and_merge** If *True*, the data is sorted and merged into as few subsections as possible (divided by *NaN*)
- **Returns** Containing the ResultValueGroup dataset objects with their *quantity labels* as keys

Returns x-y-line data grouped by the sets of "stable phases" (for example "LIQUID" or "LIQUID + FCC_A1"). Use get_values_of() instead if you need no separation. The available quantities can be found in the documentation of the factory class ScheilQuantity.

Note: The different datasets might contain *NaN*-values between different subsections and might not be sorted **even if the flag `sort_and_merge` has been set** (because they might be unsortable due to their nature).

Parameters

- **x_quantity** The first Scheil quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example "T")
- **y_quantity** The second Scheil quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example "NV")
- **sort_and_merge** If *True*, the data will be sorted and merged into as few subsections as possible (divided by *NaN*)

Returns Containing the ResultValueGroup dataset objects with their "stable phases" labels as keys

 $\texttt{get_values_of} (x_quantity: Union[tc_python.quantity_factory.ScheilQuantity, str], y_quantity: Union[tc_python.quantity_factory.ScheilQuantity, str]) \rightarrow [typing.List[float], typing.List[float]]$

Returns sorted x-y-line data without any separation. Use get_values_grouped_by_quantity_of() or get_values_grouped_by_stable_phases_of() instead if you need such a separation. The available quantities can be found in the documentation of the factory class ScheilQuantity.

Note: This method will always return sorted data without any *NaN*-values. In case of ambiguous quantities (for example: CompositionOfPhaseAsWeightFraction("FCC_A1", "All")) that can give data that is hard to interpret. In such a case you need to choose the quantity in another way or use one of the other methods.

Parameters

- **x_quantity** The first Scheil quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example "T")
- **y_quantity** The second Scheil quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example "NV")

Returns A tuple containing the x- and y-data in lists

save_to_disk (path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load_result_from_disk()

Parameters path – the path to the folder you want the result to be saved in.

Returns this *ScheilCalculationResult* object

class tc_python.scheil.ScheilCalculationType

Bases: object

Specific configuration for the different Scheil calculation types

classmethod scheil_back_diffusion()

Configuration for *back diffusion in the solid primary phase*.

Warning: This feature has only effect on systems with diffusion data (typically a mobility database). If used for a system without diffusion data, a normal Scheil calculation is done. :return: A *ScheilBackDiffusion*

classmethod scheil_classic()

Configuration for Classic Scheil with fast diffusers. :return: A ScheilClassic

classmethod scheil_solute_trapping()

Configures the Scheil solute trapping settings. The used solidification speed equation is *Scanning speed* * *cos(angle)* with *Scanning speed* and *angle* being provided either by the user or taken from the defaults. :return: A *ScheilSoluteTrapping*

class tc_python.scheil.ScheilClassic

Bases: tc_python.scheil.ScheilCalculationType

Configuration for Classic Scheil with fast diffusers.

set_fast_diffusing_elements(element_names: List[str])

Sets elements as fast diffusing. This allows redistribution of these elements in both the solid and liquid parts of the alloy.

Default: No fast-diffusing elements.

Parameters element_names - The elements

Returns This *ScheilClassic* object

class tc_python.scheil.ScheilOptions

Bases: object

Options for the Scheil simulation.

calculate_from_gas()

Calculates the evaporation temperature if a gas phase is selected in the system, and then calculates equilibria in the gas+liquid and liquid regions until liquidus temperature is reached.

Default: Calculation starts from liquidus temperature.

Returns This ScheilOptions object

calculate_from_liquidus()

Solidification calculation starting from the liquidus temperature. Liquid properties between start temperature and liquidus are not obtainable.

Default: Calculation starts from liquidus temperature.

Returns This ScheilOptions object

calculate_from_start_temperature()

Calculation of equilibria from start temperature at 50 K intervals until liquidus temperature is reached. This option makes it possible to obtain properties of the liquid phase before the solidification starts.

Default: Calculation starts from liquidus temperature.

Returns This ScheilOptions object

disable_approximate_driving_force_for_metastable_phases()

Disables the approximation of the driving force for metastable phases.

Default: Enabled

Note: When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct. use disable approximate driving force for metastable phases () to force the calculation to converge for the metastable phases.

Returns This ScheilOptions object

disable_control_step_size_during_minimization()

Disables stepsize control during minimization (non-global).

Default: Enabled

Returns This ScheilOptions object

disable_equilibrium_solidification_calculation()

Skips the property (one axis) diagram calculation of solidification under equilibrium conditions, before the Scheil solidification calculation starts.

In general it is not necessary to perform this calculation.

Default: Disabled. The equilibrium solidification calculation is skipped.

Returns This ScheilOptions object

disable_force_positive_definite_phase_hessian()

Disables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

Returns This ScheilOptions object

enable_approximate_driving_force_for_metastable_phases()

Enables the approximation of the driving force for metastable phases.

Default: Enabled

Note: When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct. use disable approximate driving force for metastable phases () to force the calculation to converge for the metastable phases.

Returns This ScheilOptions object

enable_control_step_size_during_minimization()

Enables stepsize control during normal minimization (non-global).

Default: Enabled

Returns This ScheilOptions object

enable_equilibrium_solidification_calculation()

Performs a property (one axis) diagram calculation of solidification under equilibrium conditions, before the Scheil solidification calculation starts, in the same way as is typically done in graphical and console mode.

In general it is not necessary to perform this calculation.

Default: Disabled. The equilibrium solidification calculation is skipped.

Returns This *ScheilOptions* object

enable_force_positive_definite_phase_hessian()

Enables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

Returns This ScheilOptions object

set_gas_phase (phase_name: str = 'GAS')
Sets the phase used as the gas phase.

Default: The phase "GAS".

Parameters phase_name - The phase name

Returns This ScheilOptions object

set_global_minimization_max_grid_points (max_grid_points: int = 2000)

Sets the maximum number of grid points in global minimization. ** Only applicable if global minimization is actually used**.

Default: 2000 points

Parameters max_grid_points - The maximum number of grid points

Returns This ScheilOptions object

set_global_minimization_test_interval (global_test_interval: int = 10)
Sets the interval for the global test.

Default: 10

Parameters global_test_interval – The global test interval

Returns This ScheilOptions object

set_liquid_phase (phase_name: str = 'LIQUID')
Sets the phase used as the liquid phase.

Default: The phase "LIQUID".

Parameters phase_name – The phase name

Returns This ScheilOptions object

set_max_no_of_iterations (max_no_of_iterations: int = 500)
Set the maximum number of iterations.

Default: max. 500 iterations

Note: As some models give computation times of more than 1 CPU second/iteration, this number is also used to check the CPU time and the calculation stops if 500 CPU seconds/iterations are used.

Parameters max_no_of_iterations – The max. number of iterations

Returns This ScheilOptions object

set_required_accuracy (accuracy: float = 1e-06)

Sets the required relative accuracy.

Default: 1.0E-6

Note: This is a relative accuracy, and the program requires that the relative difference in each variable must be lower than this value before it has converged. A larger value normally means fewer iterations but less accurate solutions. The value should be at least one order of magnitude larger than the machine precision.

Parameters accuracy – The required relative accuracy

Returns This ScheilOptions object

set_smallest_fraction (smallest_fraction: float = 1e-12)

Sets the smallest fraction for constituents that are unstable.

It is normally only in the gas phase that you can find such low fractions.

The **default value** for the smallest site-fractions is 1E-12 for all phases except for IDEAL phase with one sublattice site (such as the GAS mixture phase in many databases) for which the default value is always as 1E-30.

Parameters smallest_fraction – The smallest fraction for constituents that are unstable

Returns This ScheilOptions object

set_temperature_step (temperature_step_in_kelvin: float = 1.0)

Sets the temperature step. Decreasing the temperature step increases the accuracy, but the default value is usually adequate.

Default step: 1.0 K

Parameters temperature_step_in_kelvin – The temperature step [K]

Returns This ScheilOptions object

terminate_on_fraction_of_liquid_phase (*fraction_to_terminate_at: float = 0.01*) Sets the termination condition to a specified remaining fraction of liquid phase.

Default: Terminates at 0.01 fraction of liquid phase.

Note: Either the termination criterion is set to a temperature or fraction of liquid limit, both together are not possible.

Parameters fraction_to_terminate_at – the termination fraction of liquid phase (value between 0 and 1)

Returns This ScheilOptions object

terminate_on_temperature(temperature_in_kelvin: float)

Sets the termination condition to a specified temperature.

Default: Terminates at 0.01 fraction of liquid phase, i.e. not at a specified temperature.

Note: Either the termination criterion is set to a temperature or fraction of liquid limit, both together are not possible.

Parameters temperature_in_kelvin – the termination temperature [K]

Returns This ScheilOptions object

class tc_python.scheil.ScheilSoluteTrapping

Bases: tc_python.scheil.ScheilCalculationType

Configures the Scheil solute trapping settings. The used solidification speed equation is *Scanning speed* * *cos(angle)* with *Scanning speed* and *angle* being provided either by the user or taken from the defaults.

 $set_angle (alpha: float = 45.0)$

Sets the transformation angle alpha between the solid/liquid boundary and laser scanning direction.

Default: 45.0

Parameters alpha – The transformation angle [degree]

Returns This ScheilSoluteTrapping object

set_primary_phasename (primary_phase_name: str = 'AUTOMATIC')

Sets the name of the primary phase.

The primary phase is the phase where solute trapping takes place. A necessary condition for this phase is that the phase definition contains all of the elements that are chosen in the system. When *AUTOMATIC* is selected, the program tries to find a suitable primary phase that fills this condition.

Default: AUTOMATIC

Parameters primary_phase_name – The phase name (or AUTOMATIC)

Returns This ScheilSoluteTrapping object

set_scanning_speed (scanning_speed: float = 1.0)
Sets the scanning speed.

Default: 1 m/s

Parameters scanning_speed – The scaling factor [m/s]

Returns This ScheilSoluteTrapping object

5.1.5 Module "step_or_map_diagrams"

class tc_python.step_or_map_diagrams.AbstractAxisType

Bases: object

The abstract base class for all axis types.

class tc_python.step_or_map_diagrams.**AbstractPhaseDiagramCalculation**(*calculator*) **Bases:** tc_python.abstract_base.AbstractCalculation

Abstract configuration required for a property diagram calculation.

Note: This is an abstract class that cannot be used directly.

add_initial_equilibrium (*initial_equilibrium*: tc_python.step_or_map_diagrams.InitialEquilibrium) Add initial equilibrium start points from which a phase diagram is calculated.

Scans along the axis variables and generates start points when the scan procedure crosses a phase boundary.

It may take a little longer to execute than using the minimum number of start points, as some lines may be calculated more than once. But the core remembers all node points and subsequently stops calculations along a line when it finds a known node point.

It is also possible to create a sequence of start points from one initial equilibria.

Parameters initial_equilibrium – The initial equilibrium

Returns This *PhaseDiagramCalculation* object

abstract calculate (keep_previous_results: bool = False, timeout_in_minutes: float = 0.0) \rightarrow tc_python.step_or_map_diagrams.PhaseDiagramResult

disable_global_minimization()

Disables global minimization.

Default: Enabled

Returns This *PhaseDiagramCalculation* object

dont_keep_default_equilibria()

Do not keep the initial equilibria added by default.

This is only relevant in combination with add_initial_equilibrium().

This is the default behavior.

Returns This PhaseDiagramCalculation object

enable_global_minimization()

Enables global minimization.

Default: Enabled

Returns This PhaseDiagramCalculation object

get_components() \rightarrow List[str]

Returns the names of the components in the system (including all components auto-selected by the database(s)).

Returns The component names

$get_gibbs_energy_addition_for(phase: str) \rightarrow float$

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters phase – Specify the name of the (stoichiometric or solution) phase with the addition

Returns Gibbs energy addition to G per mole formula unit.

$get_system_data() \rightarrow tc_python.abstract_base.SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. *user*) databases loaded as **.tdb*-file.

Returns The system data

keep_default_equilibria()

Keep the initial equilibria added by default. This is only relevant in combination with *add_initial_equilibrium()*.

Default behavior is to not keep default equilibria.

Returns This PhaseDiagramCalculation object

remove_all_initial_equilibria()

Removes all previously added initial equilibria.

Returns This PhaseDiagramCalculation object

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

Parameters command - The Thermo-Calc Console Mode command

Returns This PhaseDiagramCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

set_gibbs_energy_addition_for (phase: str, gibbs_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs_energy Addition to G per mole formula unit

Returns This PhaseDiagramCalculation object

set_phase_to_dormant (phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This PhaseDiagramCalculation object

set_phase_to_entered (phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

Parameters

- phase The phase name or ALL_PHASES for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

Returns This PhaseDiagramCalculation object

set_phase_to_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

Parameters

- **phase** The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

Returns This PhaseDiagramCalculation object

set_phase_to_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This PhaseDiagramCalculation object

with_options (*options:* tc_python.step_or_map_diagrams.PhaseDiagramOptions) Sets the simulation options.

Parameters options – The simulation options

Returns This PhaseDiagramCalculation object

with_reference_state (component: str, phase: str = 'SER', temperature: float = - 1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

- **component** The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

Returns This PhaseDiagramCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. user) databases loaded as a *.tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This PhaseDiagramCalculation object

class tc_python.step_or_map_diagrams.**AbstractPropertyDiagramCalculation**(*calculator*) **Bases:** tc_python.abstract_base.AbstractCalculation

Abstract configuration required for a property diagram calculation.

Note: This is an abstract class that cannot be used directly.

abstract calculate (keep_previous_results: bool = False, timeout_in_minutes: float = 0.0)

disable_global_minimization()

Disables global minimization.

Default: Enabled

Returns This PropertyDiagramCalculation object

disable_step_separate_phases()

Disables step separate phases. This is the **default** setting.

Returns This PropertyDiagramCalculation object

enable_global_minimization()

Enables global minimization.

Default: Enabled

Returns This PropertyDiagramCalculation object

enable_step_separate_phases()

Enables step separate phases.

Default: By default separate phase stepping is disabled

Note: This is an advanced option, it is used mostly to calculate how the Gibbs energy for a number of phases varies for different compositions. This is particularly useful to calculate Gibbs energies for complex phases with miscibility gaps and for an ordered phase that is never disordered (e.g. SIGMA-phase, G-phase, MU-phase, etc.).

Returns This PropertyDiagramCalculation object

get_components() \rightarrow List[str]

Returns the names of the components in the system (including all components auto-selected by the database(s)).

Returns The component names

get_gibbs_energy_addition_for (*phase: str*) \rightarrow float

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters phase – Specify the name of the (stoichiometric or solution) phase with the addition

Returns Gibbs energy addition to G per mole formula unit.

$get_system_data() \rightarrow tc_python.abstract_base.SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

Parameters command - The Thermo-Calc Console Mode command

Returns This PropertyDiagramCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

set_gibbs_energy_addition_for (phase: str, gibbs_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs_energy Addition to G per mole formula unit

Returns This PropertyDiagramCalculation object

set_phase_to_dormant (phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters phase – The phase name or ALL_PHASES for all phases

Returns This PropertyDiagramCalculation object

set_phase_to_entered (phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

Parameters

- **phase** The phase name or *ALL_PHASES* for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

Returns This PropertyDiagramCalculation object

set_phase_to_fixed (phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

Parameters

- phase The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

Returns This PropertyDiagramCalculation object

set_phase_to_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This PropertyDiagramCalculation object

with_options (options: tc_python.step_or_map_diagrams.PropertyDiagramOptions)
 Sets the simulation options.

Parameters options – The simulation options

Returns This PropertyDiagramCalculation object

with_reference_state (component: str, phase: str = 'SER', temperature: float = - 1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

- **component** The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

Returns This PropertyDiagramCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)
 Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a * .tdb-file.

Parameters system_modifications - The system modification to be performed

Returns This PropertyDiagramCalculation object

```
class tc_python.step_or_map_diagrams.AxisType
    Bases: tc_python.step_or_map_diagrams.AbstractAxisType
```

Factory class providing objects for configuring a logarithmic or linear axis by using AxisType.linear() or AxisType.logarithmic().

classmethod linear()

Creates an object for configuring a linear calculation axis.

Default: A minimum number of 40 steps.

Note: The returned object can be configured regarding the maximum step size *or* the minimum number of steps on the axis.

Returns A new *Linear* object

classmethod logarithmic()

Creates an object for configuring a logarithmic calculation axis.

Default: A scale factor of 1.1

Note: The returned object can be configured regarding the scale factor.

Returns A new *Logarithmic* object

class tc_python.step_or_map_diagrams.CalculationAxis(quantity:

Union[tc_python.quantity_factory.ThermodynamicQuanstr])

Bases: object

A calculation axis used for property and phase diagram calculations.

Note: A calculation axis is defining the varied condition and the range of variation. It is the same concept as in Thermo-Calc *Graphical Mode* or *Console Mode*.

Default: A *Linear* axis with a *minimum number of 40 steps*

```
set_max(max: float)
```

Sets the maximum quantity value of the calculation axis.

There is no default value set, it always needs to be defined.

Parameters max – The maximum quantity value of the axis [unit according to the axis quantity]

Returns This CalculationAxis object

set_min (min: float)

Sets the minimum quantity value of the calculation axis.

There is no default value set, it always needs to be defined.

Parameters min – The minimum quantity value of the axis [unit according to the axis quantity]

Returns This CalculationAxis object

set_start_at (at: float)

Sets the starting point of the calculation on the axis.

Default: The default starting point is the center between the minimum and maximum quantity value

Parameters at – The starting point on the axis [unit according to the axis quantity]

Returns This CalculationAxis object

with_axis_type (axis_type: tc_python.step_or_map_diagrams.AxisType)

Sets the type of the axis.

Default: A Linear axis with a minimum number of 40 steps

Parameters axis_type – The axis type (linear or logarithmic)

Returns This CalculationAxis object

class tc_python.step_or_map_diagrams.Direction(value)

Bases: enum.Enum

An enumeration.

DECREASE_FIRST_AXIS = 3

 $DECREASE_SECOND_AXIS = 4$

 $INCREASE_FIRST_AXIS = 0$

```
INCREASE_SECOND_AXIS = 1
```

```
class tc_python.step_or_map_diagrams.InitialEquilibrium(first_axis: float, sec-
ond_axis: float)
```

Bases: object

add_equilibria_at_all_phase_changes()

This generates one start point for each set of phase change in the chosen direction of the specified axis This ensures finding all possible phase boundary lines (not just the first one) along such an axis direction.

Default behavior is to only generate one start point at the first phase change.

Returns This *InitialEquilibrium* object

add_equilibria_at_first_phase_change()

This generates one start point at the first phase change.

This is the default behavior.

Returns This InitialEquilibrium object

set_direction (*direction_enum:* tc_python.step_or_map_diagrams.Direction) Specifies along which axes the initial equilibria should be added.

The default direction is INCREASE_FIRST_AXIS.

Parameters direction_enum-

Returns This InitialEquilibrium object

class tc_python.step_or_map_diagrams.Linear Bases: tc_python.step_or_map_diagrams.AxisType

Represents a linear axis.

get_type() \rightarrow str

Convenience method for getting axis type.

Returns The type

set_max_step_size (max_step_size: float)
Sets the axis to use the maximum step size configuration.

Default: This is not the default which is minimum number of steps

Note: Either maximum step size or minimum number of steps can be used but not both at the same time.

Parameters max_step_size – The maximum step size [unit according to the axis quantity]

Returns This Linear object

set_min_nr_of_steps (min_nr_of_steps: float = 40)

Sets the axis to use the minimum number of steps configuration.

Default: This is the default option (with a minimum number of steps of 40)

Note: Either maximum step size or minimum number of steps can be used but not both at the same time.

Parameters min_nr_of_steps - The minimum number of steps

Returns This Linear object

```
class tc_python.step_or_map_diagrams.Logarithmic(scale_factor: float = 1.1)
Bases: tc_python.step_or_map_diagrams.AxisType
```

Represents a logarithmic axis.

Note: A logarithmic axis is useful for low fractions like in a gas phase where 1E-7 to 1E-2 might be an interesting range. For the pressure a logarithmic axis is often also useful.

 $get_type() \rightarrow str$ Convenience method for getting axis type.

Returns The type

set_scale_factor (scale_factor: float = 1.1)
Sets the scale factor.

Default: 1.1

Parameters scale_factor – The scale factor setting the maximum factor between two calculated values, must be larger than 1.0

Returns This Logarithmic object

class tc_python.step_or_map_diagrams.**PhaseDiagramCalculation** (*calculator*) **Bases:** tc_python.step_or_map_diagrams.AbstractPhaseDiagramCalculation

Configuration for a phase diagram calculation.

Note: Specify the conditions, the calculation is performed with *calculate()*.

add_initial_equilibrium (*initial_equilibrium*: tc_python.step_or_map_diagrams.InitialEquilibrium) Add initial equilibrium start points from which a phase diagram is calculated.

Scans along the axis variables and generates start points when the scan procedure crosses a phase boundary.

It may take a little longer to execute than using the minimum number of start points, as some lines may be calculated more than once. But the core remembers all node points and subsequently stops calculations along a line when it finds a known node point.

It is also possible to create a sequence of start points from one initial equilibria.

Parameters initial_equilibrium – The initial equilibrium

Returns This *PhaseDiagramCalculation* object

calculate (*keep_previous_results: bool = False, timeout_in_minutes: float = 0.0*) \rightarrow *tc_python.step_or_map_diagrams.PhaseDiagramResult* Performs the phase diagram calculation.

Warning: If you use *keep_previous_results=True*, you must not use another calculator or even get results in between the calculations using *calculate()*. Then the previous results will actually be lost.

Parameters

- **keep_previous_results** If True, results from any previous call to this method are appended. This can be used to combine calculations with multiple start points if the mapping fails at a certain condition.
- timeout_in_minutes Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

Returns A new *PhaseDiagramResult* object which later can be used to get specific values from the calculated result.

disable_global_minimization()

Disables global minimization.

Default: Enabled

Returns This PhaseDiagramCalculation object

dont_keep_default_equilibria()

Do not keep the initial equilibria added by default.

This is only relevant in combination with add_initial_equilibrium().

This is the default behavior.

Returns This PhaseDiagramCalculation object

enable_global_minimization()

Enables global minimization.

Default: Enabled

Returns This PhaseDiagramCalculation object

get_components() \rightarrow List[str]

Returns the names of the components in the system (including all components auto-selected by the database(s)).

Returns The component names

get_gibbs_energy_addition_for (*phase: str*) \rightarrow float

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters phase – Specify the name of the (stoichiometric or solution) phase with the addition

Returns Gibbs energy addition to G per mole formula unit.

$\texttt{get_system_data()} \rightarrow \textit{tc_python.abstract_base.SystemData}$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

keep_default_equilibria()

Keep the initial equilibria added by default. This is only relevant in combination with *add_initial_equilibrium()*.

Default behavior is to not keep default equilibria.

Returns This PhaseDiagramCalculation object

remove_all_conditions()

Removes all set conditions.

Returns This PhaseDiagramCalculation object

remove_all_initial_equilibria()

Removes all previously added initial equilibria.

Returns This PhaseDiagramCalculation object

remove_condition (*quantity: Union*[tc_python.quantity_factory.ThermodynamicQuantity, *str*]) Removes the specified condition.

Parameters quantity – The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example X(Cr))

Returns This ThermodynamicCalculation object

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

Parameters command - The Thermo-Calc Console Mode command

Returns This PhaseDiagramCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

set_condition(quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str], value:

float) Sets the specified condition.

Parameters

- quantity The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example X(Cr))
- **value** The value of the condition

Returns This PhaseDiagramCalculation object

set_gibbs_energy_addition_for (phase: str, gibbs_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- **gibbs_energy** Addition to G per mole formula unit

Returns This PhaseDiagramCalculation object

set_phase_to_dormant (phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This *PhaseDiagramCalculation* object

set_phase_to_entered (phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

Parameters

- phase The phase name or ALL_PHASES for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

Returns This *PhaseDiagramCalculation* object

set_phase_to_fixed (phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

Parameters

• phase – The phase name

• **amount** – The fixed phase fraction (between 0.0 and 1.0)

Returns This PhaseDiagramCalculation object

set_phase_to_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This PhaseDiagramCalculation object

with_first_axis (axis: tc_python.step_or_map_diagrams.CalculationAxis)
 Sets the first calculation axis.

Parameters axis – The axis

Returns This PhaseDiagramCalculation object

with_options (options: tc_python.step_or_map_diagrams.PhaseDiagramOptions)
 Sets the simulation options.

Parameters options – The simulation options

Returns This PhaseDiagramCalculation object

with_reference_state (component: str, phase: str = 'SER', temperature: float = - 1.0, pressure:

float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

- component The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

Returns This PhaseDiagramCalculation object

with_second_axis (*axis*: tc_python.step_or_map_diagrams.CalculationAxis) Sets the second calculation axis.

Parameters axis – The axis

Returns This *PhaseDiagramCalculation* object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)
 Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. user) databases loaded as a *.tdb-file.

Parameters system_modifications - The system modification to be performed

Returns This PhaseDiagramCalculation object

class tc_python.step_or_map_diagrams.PhaseDiagramOptions

Bases: object

Simulation options for phase diagram calculations.

```
disable_approximate_driving_force_for_metastable_phases()
```

Disables the approximation of the driving force for metastable phases.

Default: Enabled

Note: When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces use are correct. disable_approximate_driving_force_for_metastable_phases() to force the calculation to converge for the metastable phases.

Returns This PhaseDiagramOptions object

disable_control_step_size_during_minimization()

Disables stepsize control during minimization (non-global).

Default: Enabled

Returns This PhaseDiagramOptions object

disable_force_positive_definite_phase_hessian()

Disables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

Returns This PhaseDiagramOptions object

dont_use_auto_start_points()

Switches the usage of automatic starting points for the mapping off.

Default: Switched on

Returns This *PhaseDiagramOptions* object

dont_use_inside_mesh_points()

Switches the usage of inside meshing points for the mapping off.

Default: Switched off

Returns This PhaseDiagramOptions object

enable_approximate_driving_force_for_metastable_phases()

Enables the approximation of the driving force for metastable phases.

Default: Enabled

Note: When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

driving forces correct, If it is important that these are use disable_approximate_driving_force_for_metastable_phases() to force the calculation to converge for the metastable phases.

Returns This PhaseDiagramOptions object

enable_control_step_size_during_minimization()

Enables stepsize control during normal minimization (non-global).

Default: Enabled

Returns This *PhaseDiagramOptions* object

enable_force_positive_definite_phase_hessian()

Enables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

Returns This PhaseDiagramOptions object

set_global_minimization_max_grid_points (max_grid_points: int = 2000)

Sets the maximum number of grid points in global minimization. ** Only applicable if global minimization is actually used**.

Default: 2000 points

Parameters max_grid_points – The maximum number of grid points

Returns This PhaseDiagramOptions object

set_global_minimization_test_interval (*global_test_interval: int = 0*) Sets the interval for the global test.

Default: 0

Parameters global_test_interval – The global test interval

Returns This PhaseDiagramOptions object

set_max_no_of_iterations (max_no_of_iterations: int = 500)
Set the maximum number of iterations.

Default: max. 500 iterations

Note: As some models give computation times of more than 1 CPU second/iteration, this number is also used to check the CPU time and the calculation stops if 500 CPU seconds/iterations are used.

Parameters max_no_of_iterations - The max. number of iterations

Returns This PhaseDiagramOptions object

set_no_of_mesh_along_axis (no_of_mesh_along_axis: int = 3)
Sets the number of meshes along an axis for the mapping.

Default: 3

Parameters no_of_mesh_along_axis - The number of meshes

Returns This PhaseDiagramOptions object

set_required_accuracy(accuracy: float = 1e-06)

Sets the required relative accuracy.

Default: 1.0E-6

Note: This is a relative accuracy, and the program requires that the relative difference in each variable must be lower than this value before it has converged. A larger value normally means fewer iterations but less accurate solutions. The value should be at least one order of magnitude larger than the machine precision.

Parameters accuracy – The required relative accuracy

Returns This PhaseDiagramOptions object

```
set_smallest_fraction (smallest_fraction: float = 1e-12)
```

Sets the smallest fraction for constituents that are unstable.

It is normally only in the gas phase that you can find such low fractions.

The **default value** for the smallest site-fractions is 1E-12 for all phases except for IDEAL phase with one sublattice site (such as the GAS mixture phase in many databases) for which the default value is always as 1E-30.

Parameters smallest_fraction – The smallest fraction for constituents that are unstable

Returns This PhaseDiagramOptions object

```
use_auto_start_points()
```

Switches the usage of automatic starting points for the mapping on.

Default: Switched on

Returns This PhaseDiagramOptions object

use_inside_mesh_points()

Switches the usage of inside meshing points for the mapping off.

Default: Switched off

Returns This PhaseDiagramOptions object

```
class tc_python.step_or_map_diagrams.PhaseDiagramResult (result)
    Bases: tc python.abstract base.AbstractResult
```

Result of a phase diagram calculation, it can be evaluated using quantities or Console Mode syntax.

add_coordinate_for_phase_label (x: float, y: float)

Sets a coordinate in the result plot for which the stable phases will be evaluated and provided in the result data object. This can be used to plot the phases of a region into the phase diagram or just to programmatically evaluate the phases in certain regions.

Warning: This method takes coordinates of the plot axes and not of the calculation axis.

Parameters

- x The coordinate of the first plot axis ("x-axis") [unit of the plot axis]
- y The coordinate of the second plot axis ("y-axis") [unit of the plot axis]

Returns This PhaseDiagramResult object

get_values_grouped_by_quantity_of (x_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str], y_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str]) \rightarrow tc_python.step_or_map_diagrams.PhaseDiagramResultValues

Returns x-y-line data grouped by the multiple datasets of the specified quantities (for example in dependency of components). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity. Usually the result data represents the phase diagram.

Note: The different datasets will contain NaN-values between different subsections and are not sorted (because they are unsortable due to their nature).

Note: Its possible to use functions as axis variables, either by using ThermodynamicQuan*tity.user_defined_function*, or by using an expression that contains '='.

Example get_values_grouped_by_quantity_of('T', ThermodynamicQuantity.user_defined_function('HM.T'))

Example get_values_grouped_by_quantity_of('T', 'CP=HM.T')

Parameters

- **x_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example 'T'), or even a function (for example 'f=T*1.01')
- y quantity The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example 'NV'), or even a function (for example 'CP=HM.T')

Returns The phase diagram data

get_values_grouped_by_stable_phases_of (x_quantity: Union/tc_python.quantity_factory.ThermodynamicQuantit str], *y_quantity:* Union/tc_python.quantity_factory.ThermodynamicQuantity,

 $str]) \rightarrow tc_python.step_or_map_diagrams.PhaseDiagramResultValues$ Returns x-y-line data grouped by the sets of "stable phases" (for example "LIQUID" or "LIQUID" + FCC_A1"). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity. Usually the result data represents the phase diagram.

Note: The different datasets will contain *NaN*-values between different subsections and are not sorted (because they are unsortable due to their nature).

Note: Its possible to use functions as axis variables, either by using *ThermodynamicQuantity.user_defined_function*, or by using an expression that contains '='.

Example get_values_grouped_by_quantity_of('T', ThermodynamicQuantity.user_defined_function('HM.T'))

Example get_values_grouped_by_quantity_of('T', 'CP=HM.T')

Parameters

- **x_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example '*T*'), or even a function (for example '*f*=*T**1.01')
- **y_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), or even a function (for example '*CP=HM.T*')

Returns The phase diagram data

remove_phase_labels()

Erases all added coordinates for phase labels.

Returns This *PhaseDiagramResult* object

save_to_disk (path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load_result_from_disk()

Parameters path – the path to the folder you want the result to be saved in. It can be relative or absolute.

Returns this *PhaseDiagramResult* object

set_phase_name_style (phase_name_style_enum: tc_python.step_or_map_diagrams.PhaseNameStyle = <PhaseNameStyle.NONE: 0>)

Sets the style of the phase name labels that will be used in the result data object (constitution description, ordering description, \dots).

Default: PhaseNameStyle.NONE

Parameters phase_name_style_enum - The phase name style

Returns This *PhaseDiagramResult* object

class tc_python.step_or_map_diagrams.PhaseDiagramResultValues (phase_diagram_values_java)
Bases: object

Represents the data of a phase diagram.

$get_invariants() \rightarrow tc_python.utils.ResultValueGroup$

Returns the x- and y-datasets of all invariants in the phase diagram.

Note: The datasets will normally contain different sections separated by NaN-values.

Returns The invariants dataset object

get_lines () \rightarrow Dict[str, *tc_python.utils.ResultValueGroup*]

Returns the x- and y-datasets of all phase boundaries in the phase diagram.

Note: The datasets will normally contain different sections separated by NaN-values.

Returns Containing the phase boundary datasets with the *quantities* or *stable phases* as keys (depending on the used method to get the values)

get_phase_labels() → List[tc_python.step_or_map_diagrams.PhaseLabel]
Returns the phase labels added for certain coordinates using PhaseDiagramResult.
add_coordinate_for_phase_label().

Returns The list with the phase label data (that contains plot coordinates and stable phases)

get_tie_lines () \rightarrow tc_python.utils.ResultValueGroup Returns the x- and y-datasets of all tie-lines in the phase diagram.

Note: The datasets will normally contain different sections separated by NaN-values.

Returns The tie-line dataset object

```
class tc_python.step_or_map_diagrams.PhaseLabel (phase_label_java)
    Bases: object
```

Represents a *phase label at a plot coordinate*, i.e. the stable phases that are present at that *plot* coordinate.

```
\begin{array}{l} \texttt{get\_text} () \rightarrow \texttt{str} \\ Accessor \ \texttt{for the phase label :return: the phase label} \end{array}
```

 $get_x() \rightarrow List[float]$ Accessor for the x-value :return: the x value

```
get_y() \rightarrow List[float]
Accessor for the y-value :return: the y value
```

class tc_python.step_or_map_diagrams.PhaseNameStyle(value)

Bases: enum.Enum

The style of the phase names used in the labels.

ALL = 1

Adding ordering and constitution description.

```
CONSTITUTION_DESCRIPTION = 3
```

Adding only constitution description.

NONE = 0

Only the phase names.

$ORDERING_DESCRIPTION = 4$

Adding only ordering description.

class tc_python.step_or_map_diagrams.**PropertyDiagramCalculation** (*calculator*) Bases: tc_python.step_or_map_diagrams.AbstractPropertyDiagramCalculation

calculate (*keep_previous_results: bool* = *False*, *timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.step_or_map_diagrams.PropertyDiagramResult* Performs the property diagram calculation.

Warning: If you use *keep_previous_results=True*, you must not use another calculator or even get results in between the calculations using *calculate()*. Then the previous results will actually be lost.

Parameters

- **keep_previous_results** If *True*, results from any previous call to this method are appended. This can be used to combine calculations with multiple start points if the stepping fails at a certain condition.
- **timeout_in_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

Returns A new *PropertyDiagramResult* object which later can be used to get specific values from the calculated result

disable_global_minimization()

Disables global minimization.

Default: Enabled

Returns This PropertyDiagramCalculation object

disable_step_separate_phases()

Disables step separate phases. This is the default setting.

Returns This PropertyDiagramCalculation object

enable_global_minimization()

Enables global minimization.

Default: Enabled

Returns This PropertyDiagramCalculation object

enable_step_separate_phases()

Enables step separate phases.

Default: By default separate phase stepping is disabled

Note: This is an advanced option, it is used mostly to calculate how the Gibbs energy for a number of phases varies for different compositions. This is particularly useful to calculate Gibbs energies for complex phases with miscibility gaps and for an ordered phase that is never disordered (e.g. SIGMA-phase, G-phase, MU-phase, etc.).

Returns This PropertyDiagramCalculation object

get_components() \rightarrow List[str]

Returns the names of the components in the system (including all components auto-selected by the database(s)).

Returns The component names

$get_gibbs_energy_addition_for(phase: str) \rightarrow float$

Used to get the additional energy term (always being a constant) of a given phase. The value given is added

to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters phase – Specify the name of the (stoichiometric or solution) phase with the addition

Returns Gibbs energy addition to G per mole formula unit.

get_system_data() \rightarrow tc_python.abstract_base.SystemData

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with_system_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

remove_all_conditions()

Removes all set conditions.

Returns This PropertyDiagramCalculation object

remove_condition (*quantity: Union*[tc_python.quantity_factory.ThermodynamicQuantity, *str*]) Removes the specified condition.

Parameters quantity – The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example X(Cr))

Returns This PropertyDiagramCalculation object

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

Parameters command - The Thermo-Calc Console Mode command

Returns This PropertyDiagramCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

Sets the specified condition.

Parameters

- **quantity** The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example *X*(*Cr*))
- value The value of the condition

Returns This PropertyDiagramCalculation object

set_gibbs_energy_addition_for (phase: str, gibbs_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters

- **phase** Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs_energy Addition to G per mole formula unit

Returns This PropertyDiagramCalculation object

set_phase_to_dormant (phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters phase – The phase name or ALL_PHASES for all phases

Returns This *PropertyDiagramCalculation* object

set_phase_to_entered (phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

Parameters

- **phase** The phase name or *ALL_PHASES* for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

Returns This PropertyDiagramCalculation object

set_phase_to_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

Parameters

- **phase** The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

Returns This PropertyDiagramCalculation object

set_phase_to_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters phase – The phase name or ALL_PHASES for all phases

Returns This PropertyDiagramCalculation object

with_axis (axis: tc_python.step_or_map_diagrams.CalculationAxis)
 Sets the calculation axis.

tets the ealeulation axis.

Parameters axis - The axis

Returns This PropertyDiagramCalculation object

with_options (options: tc_python.step_or_map_diagrams.PropertyDiagramOptions)
 Sets the simulation options.

Parameters options – The simulation options

Returns This PropertyDiagramCalculation object

with_reference_state (component: str, phase: str = 'SER', temperature: float = - 1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

- **component** The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

Returns This *PropertyDiagramCalculation* object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)
 Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. user) databases loaded as a *.tdb-file.

Parameters system_modifications - The system modification to be performed

Returns This *PropertyDiagramCalculation* object

class tc_python.step_or_map_diagrams.PropertyDiagramOptions
 Bases: object

Simulation options for the property diagram calculations.

disable_approximate_driving_force_for_metastable_phases() Disables the approximation of the driving force for metastable phases.

Default: Enabled

Note: When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use disable_approximate_driving_force_for_metastable_phases() force the to calculation to converge for the metastable phases.

Returns This PropertyDiagramOptions object

disable_control_step_size_during_minimization()

Disables stepsize control during minimization (non-global).

Default: Enabled

Returns This PropertyDiagramOptions object

disable_force_positive_definite_phase_hessian()

Disables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

Returns This PropertyDiagramOptions object

enable_approximate_driving_force_for_metastable_phases()

Enables the approximation of the driving force for metastable phases.

Default: Enabled

Note: When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces correct, use are disable_approximate_driving_force_for_metastable_phases() force the to calculation to converge for the metastable phases.

Returns This *PropertyDiagramOptions* object

enable_control_step_size_during_minimization()

Enables stepsize control during normal minimization (non-global).

Default: Enabled

Returns This PropertyDiagramOptions object

enable_force_positive_definite_phase_hessian()

Enables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

Returns This *PropertyDiagramOptions* object

set_global_minimization_max_grid_points (max_grid_points: int = 2000)

Sets the maximum number of grid points in global minimization. Only applicable if global minimization is actually used.

Default: 2000 points

Parameters max_grid_points - The maximum number of grid points

Returns This PropertyDiagramOptions object

set_global_minimization_test_interval (global_test_interval: int = 0)
Sets the interval for the global test.

Default: 0

Parameters global_test_interval – The global test interval

Returns This PropertyDiagramOptions object

set_max_no_of_iterations (max_no_of_iterations: int = 500)
Set the maximum number of iterations.

Default: max. 500 iterations

Note: As some models give computation times of more than 1 CPU second/iteration, this number is also used to check the CPU time and the calculation stops if 500 CPU seconds/iterations are used.

Parameters max_no_of_iterations - The max. number of iterations

Returns This PropertyDiagramOptions object

set_required_accuracy (accuracy: float = 1e-06)

Sets the required relative accuracy.

Default: 1.0E-6

Note: This is a relative accuracy, and the program requires that the relative difference in each variable must be lower than this value before it has converged. A larger value normally means fewer iterations but less accurate solutions. The value should be at least one order of magnitude larger than the machine precision.

Parameters accuracy – The required relative accuracy

Returns This PropertyDiagramOptions object

```
set_smallest_fraction (smallest_fraction: float = 1e-12)
```

Sets the smallest fraction for constituents that are unstable.

It is normally only in the gas phase that you can find such low fractions.

The **default value** for the smallest site-fractions is 1E-12 for all phases except for IDEAL phase with one sublattice site (such as the GAS mixture phase in many databases) for which the default value is always as 1E-30.

Parameters smallest_fraction – The smallest fraction for constituents that are unstable

Returns This PropertyDiagramOptions object

class tc_python.step_or_map_diagrams.PropertyDiagramResult(result)
 Bases: tc_python.abstract_base.AbstractResult

Result of a property diagram. This can be used to query for specific values.

get_values_grouped_by_quantity_of (x_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity,

str], y_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity,

str], sort_and_merge: bool = True) \rightarrow Dict[str,

tc python.utils.ResultValueGroup

Returns x-y-line data grouped by the multiple datasets of the specified quantities (typically the phases). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity.

Note: The different datasets might contain *NaN*-values between different subsections and might not be sorted **even if the flag `sort_and_merge` has been set** (because they might be unsortable due to their nature).

Note: Its possible to use functions as axis variables, either by using *ThermodynamicQuantity.user_defined_function*, or by using an expression that contains '='.

Example get_values_grouped_by_quantity_of('T', ThermodynamicQuantity.user_defined_function('HM.T'))

Example get_values_grouped_by_quantity_of('T', 'CP=HM.T')

Parameters

- **x_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example '*T*'), or even a function (for example '*f*=*T**1.01')
- **y_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), or even a function (for example '*CP=HM.T*')
- **sort_and_merge** If *True*, the data is sorted and merged into as few subsections as possible (divided by *NaN*)

Returns Containing the datasets with the quantities as their keys

Returns x-y-line data grouped by the sets of "stable phases" (for example "LIQUID" or "LIQUID + FCC_A1"). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity.

Note: The different datasets might contain *NaN*-values between different subsections and different lines of an ambiguous dataset. They might not be sorted **even if the flag** `sort_and_merge` has been set (because they might be unsortable due to their nature).

Note: Its possible to use functions as axis variables, either by using *ThermodynamicQuantity.user_defined_function*, or by using an expression that contains '='.

Exampleget_values_grouped_by_quantity_of('T',ThermodynamicQuantity_user_defined_function('HM.T'))

Example get_values_grouped_by_quantity_of(`T', `CP=HM.T')

Parameters

- **x_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example 'T'), or even a function (for example 'f=T*1.01')
- **y_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), or even a function (for example '*CP=HM.T*')
- **sort_and_merge** If *True*, the data will be sorted and merged into as few subsections as possible (divided by *NaN*)

Returns Containing the datasets with the quantities as their keys

get_values_of (x_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str], y_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str]) \rightarrow [typing.List[float], typing.List[float]]

Returns sorted x-y-line data without any separation. Use get_values_grouped_by_quantity_of() or get_values_grouped_by_stable_phases_of() instead if you need such a separation. The available quantities can be found in the documentation of the factory class ThermodynamicQuantity.

Note: This method will always return sorted data without any *NaN*-values. If it is unsortable that might give data that is hard to interpret. In such a case you need to choose the quantity in another way or use one of the other methods. One example of this is to use quantities with *All*-markers, for example *MassFractionOfAComponent("All")*.

Note: Its possible to use functions as axis variables, either by using *ThermodynamicQuantity.user_defined_function*, or by using an expression that contains '='.

Example get_values_grouped_by_quantity_of('T', ThermodynamicQuantity.user_defined_function('HM.T'))

Example get_values_grouped_by_quantity_of('T', 'CP=HM.T')

Parameters

- **x_quantity** The first Thermodynamic quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example 'T') or even a function (for example 'f=T*1.01')
- **y_quantity** The second Thermodynamic quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), or even a function (for example '*CP=HM.T*')

Returns A tuple containing the x- and y-data in lists

save_to_disk (path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load_result_from_disk()

Parameters path – the path to the folder you want the result to be saved in. It can be relative or absolute.

Returns this PropertyDiagramResult object

set_phase_name_style (phase_name_style_enum: tc_python.step_or_map_diagrams.PhaseNameStyle = <PhaseNameStyle.NONE: 0>)

Sets the style of the phase name labels that will be used in the result data object (constitution description, ordering description, \dots).

Default: PhaseNameStyle.NONE

Parameters phase_name_style_enum - The phase name style

Returns This PropertyDiagramResult object

5.1.6 Module "diffusion"

class tc_python.diffusion.AbstractBoundaryCondition
 Bases: object

The abstract base class for all boundary conditions.

class tc_python.diffusion.AbstractCalculatedGrid
 Bases: tc_python.diffusion.AbstractGrid

The abstract base class for calculated grids.

class tc_python.diffusion.AbstractElementProfile
 Bases: object

The abstract base class for all initial composition profile types.

class tc_python.diffusion.AbstractGrid
 Bases: object

The abstract base class for all grids.

class tc_python.diffusion.AbstractSolver
 Bases: object

Abstract base class for the solvers (Classic, Homogenization and Automatic).

class tc_python.diffusion.**ActivityFluxFunction Bases**: tc_python.diffusion.BoundaryCondition

$\texttt{get_type}() \rightarrow str$

The type of the boundary condition.

Returns The type

 $set_flux_function$ (element_name: str, f: str = '0', g: str = '1', n: float = 1.0, to_time: float = 1.7976931348623157e+308)

The flux for the independent components must be given in the format:

 $J = f(T,P,TIME) * (ACTIVITY^N - g(T,P,TIME))$

where f and g may be functions of time (TIME), temperature (T), and pressure (P), and N is an integer.

f and g must be expressed in DICTRA Console Mode syntax.

Parameters

- element_name The name of the element
- **f** the function f in the formula above
- g the function g in the formula above
- **n** the constant N in the formula above

• to_time – The max-time for which the flux function is used.

class tc_python.diffusion.AutomaticSolver

Bases: tc_python.diffusion.Solver

Solver using the *homogenization model* if any region has more than one phase, otherwise using the *classic model*.

Note: This is the default solver and recommended for most applications.

```
\begin{array}{l} \texttt{get\_type}() \rightarrow str\\ \text{The type of the solver.} \end{array}
```

Returns The type

set_flux_balance_equation_accuracy (accuracy: float = 1e-16)

Only valid if the :class: ClassicSolver` is actually used (i.e. not more than one phase in each region).

Sets the required accuracy during the solution of the flux balance equations. Default: 1.0e-16

Parameters accuracy – The required accuracy

Returns A new AutomaticSolver object

set_tieline_search_variable_to_activity()

Only valid if the :class: ClassicSolver` is actually used (i.e. not more than one phase in each region).

Configures the solver to use the *activity of a component* to find the correct tie-line at the phase interface. Either activity or chemical potential are applied to reduce the degrees of freedom at the local equilibrium. **Default**: This is the default setting

Returns A new AutomaticSolver object

set_tieline_search_variable_to_potential() Ole is is in the second second

Only valid if the :class: ClassicSolver` is actually used (i.e. not more than one phase in each region).

Configures the solver to use the *chemical potential of a component* to find the correct tie-line at the phase interface. Either activity or chemical potential are applied to reduce the degrees of freedom at the local equilibrium. **Default**: To use the activity

Returns A new AutomaticSolver object

class tc_python.diffusion.BoundaryCondition

Bases: tc_python.diffusion.AbstractBoundaryCondition

Contains factory methods for the the different boundary conditions available.

classmethod activity_flux_function()

Factory method that creates a **new** activity-flux-function boundary condition.

This type of boundary condition is used to take into account the finite rate of a surface reaction.

The flux for the independent components must be given in the format:

$J = f(T,P,TIME) * (ACTIVITY^N - g(T,P,TIME))$

where f and g may be functions of time (TIME), temperature (T), and pressure (P), and N is an integer.

f and g must be expressed in DICTRA Console Mode syntax.

Note: The activities are those with user-defined reference states. The function mass transfer coefficient

is the mass transfer coefficient, activity of the corresponding species in the gas is the activity of the corresponding species in the gas and N is a stoichiometric coefficient.

Note: For more details see L. Sproge and J. Ågren, "Experimental and theoretical studies of gas consumption in the gas carburizing process" J. Heat Treat. 6, 9–19 (1988).

Returns A new ActivityFluxFunction object

classmethod closed_system()

Factory method that creates a **new** closed-system boundary condition.

Returns A new *ClosedSystem* object

classmethod fix_flux_value()

Factory method that creates a new fix-flux-value boundary condition.

This type of boundary condition makes it possible to enter functions that yield the flux times the molar volume for the independent components. May be a function of time, temperature and pressure: J(T,P,TIME).

Returns A new FixFluxValue object

classmethod fixed_compositions (unit_enum: tc_python.diffusion.Unit = <Unit.MASS_PERCENT: 3>) Factory method that creates a **new** fixed-composition boundary condition.

Parameters unit_enum – The composition unit

Returns A new *FixedCompositions* object

classmethod mixed_zero_flux_and_activity()

Factory method that creates a new mixed zero-flux and activity boundary condition

Returns A new *MixedZeroFluxAndActivity* object

class tc_python.diffusion.CalculatedGrid

Bases: tc_python.diffusion.AbstractCalculatedGrid

Factory class for grids generated by a mathematical series (linear, geometric, ...). Use *tc_python*. *diffusion*.*PointByPointGrid* instead if you want to use an existing grid from experimental data or a previous calculation.

Note: A region must contain a number of grid points. The composition is only known at these grid points and the software assumes that the composition varies linearly between them. The amount and composition of all the phases present at a single grid point in a certain region are those given by thermodynamic equilibrium keeping the over-all composition at the grid point fixed.

```
classmethod double_geometric (no_of_points: int = 50, lower_geometrical_factor: float = 1.1,
upper_geometrical_factor: float = 0.9)
Factory method that creates a new double geometric grid.
```

Note: Double geometric grids have a high number of grid points in the middle or at both ends of a region. One geometrical factor for the lower (left) and upper (right) half of the region need to specified. In both cases a geometrical factor of larger than one yields a higher density of grid points at the lower end of the half and vice versa for a factor smaller than one.

Parameters

- no_of_points The number of points
- lower_geometrical_factor The geometrical factor for the left half
- upper_geometrical_factor The geometrical factor for the right half

Returns A new DoubleGeometricGrid object

```
classmethod geometric (no_of_points: int = 50, geometrical_factor: float = 1.1)
Factory method that creates a new geometric grid.
```

Note: A grid that yields a varying density of grid points in the region. A geometrical factor larger than one yields a higher density of grid points at the lower end of the region and a factor smaller than one yields a higher density of grid points at the upper end of the region.

Parameters

- no_of_points The number of points
- geometrical_factor The geometrical factor

Returns A new GeometricGrid object

classmethod linear (*no_of_points: int = 50*) Factory method that creates a **new** equally spaced grid.

Parameters no_of_points – The number of points

Returns A new *LinearGrid* object

class tc_python.diffusion.ClassicSolver

Bases: tc_python.diffusion.Solver

Solver using the Classic model.

Note: This solver **never switches** to the homogenization model even if it fails to converge. Use the *tc_python.diffusion.AutomaticSolver* if necessary instead.

get_type () \rightarrow str

Convenience method for getting the type of the solver.

Returns The type of the solver

set_flux_balance_equation_accuracy (accuracy: float = 1e-16)

Sets the required accuracy during the solution of the flux balance equations. Default: 1.0e-16

Parameters accuracy – The required accuracy

Returns A new ClassicSolver object

set_tieline_search_variable_to_activity()

Configures the solver to use the *activity of a component* to find the correct tie-line at the phase interface. Either activity or chemical potential are applied to reduce the degrees of freedom at the local equilibrium. **Default**: This is the default setting

set_tieline_search_variable_to_potential()

Configures the solver to use the chemical potential of a component to find the correct tie-line at the phase

<Unit.MASS PERCENT: 3>)

interface. Either activity or chemical potential are applied to reduce the degrees of freedom at the local equilibrium. **Default**: To use the activity

Returns A new ClassicSolver object

class tc_python.diffusion.ClosedSystem

Bases: tc_python.diffusion.BoundaryCondition

Represents a boundary for a closed system.

 $\texttt{get_type()} \rightarrow str$

Convenience method for getting the type of the boundary condition.

Returns The type of the boundary condition

class tc_python.diffusion.**CompositionProfile**(*unit_enum: tc_python.diffusion.Unit* =

Bases: object

Contains initial concentration profiles for the elements.

add (*element_name: str, profile:* tc_python.diffusion.ElementProfile) Adds a concentration profile for the specified element.

Parameters

• element_name - The name of the element

• profile – The initial concentration profile

Returns A CompositionProfile object

class tc_python.diffusion.**ConstantProfile**(*value: float*) Bases: tc_python.diffusion.ElementProfile

Represents a constant initial concentration profile.

get_type () \rightarrow str The type of the element profile.

Returns The type

class tc_python.diffusion.**ContinuedDiffusionCalculation**(*calculation*) Bases: tc_python.abstract_base.AbstractCalculation

Configuration for a diffusion calculation that is a continuation of a previous isothermal or non-isothermal diffusion calculation. It contains a subset of the settings possible in the original calculation.

Use set_simulation_time() to set a simulation time that is higher than the original calculation.

calculate (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.diffusion.DiffusionCalculationResult* Runs the diffusion calculation.

Parameters timeout_in_minutes – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

Returns A *DiffusionCalculationResult* which later can be used to get specific values from the calculated result

set_simulation_time (simulation_time: float)

Sets the simulation time.

Parameters simulation_time – The simulation time [s]

Returns This DiffusionIsoThermalCalculation object

Defines the boundary condition on the left edge of the system.

Default: A closed-system boundary condition.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Note: You can specify time-dependent boundary conditions by calling with_left_boundary_condition() many times, with different values of the "to" parameter.

Examples:

- with_left_boundary_condition(BoundaryCondition.closed_system(), to=100)
- with_left_boundary_condition(BoundaryCondition.mixed_zero_flux_and_activity().set_activity_for_element("C", surface_activity), to=500)
- with_left_boundary_condition(BoundaryCondition.closed_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

Parameters

- boundary_condition The boundary condition
- to The upper time-limit for boundary_condition.

Returns This DiffusionIsoThermalCalculation object

with_options (*options*: tc_python.diffusion.Options, *to*: *float* = 1.7976931348623157e+308) Sets the general simulation conditions.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Parameters

- **options** The general simulation conditions
- to The upper time-limit for options.

Returns This DiffusionIsoThermalCalculation object

with_right_boundary_condition (boundary_condition: tc_python.diffusion.BoundaryCondition,

to: float = 1.7976931348623157e+308)

Defines the boundary condition on the right edge of the system.

Default: A closed-system boundary condition

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Note: You can specify time-dependent boundary conditions by calling *with_right_boundary_condition()* many times, with different values of the "to" parameter.

Examples:

- with_right_boundary_condition(BoundaryCondition.closed_system(), to=100)
- with_right_boundary_condition(BoundaryCondition.mixed_zero_flux_and_activity().set_activity_for_element("C", surface_activity), to=500)
- with_right_boundary_condition(BoundaryCondition.closed_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

Parameters

- **boundary_condition** The boundary condition
- to The upper time-limit for boundary_condition.

Returns This DiffusionIsoThermalCalculation object

with_solver (solver: tc_python.diffusion.Solver, to: float = 1.7976931348623157e+308)
Sets the solver to use (Classic, Homogenization or Automatic). Default is Automatic.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Parameters

- **solver** The solver to use
- to The upper time-limit for solver.

Returns This DiffusionIsoThermalCalculation object

Sets the timestep control options.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Parameters

- timestep_control The new timestep control options
- to The upper time-limit for timestep_control.

Returns This DiffusionIsoThermalCalculation object

class tc_python.diffusion.DiffusionCalculationResult(result)

Bases: tc_python.abstract_base.AbstractResult

Result of a diffusion calculation. This can be used to query for specific values. For details of the axis variables, search the Thermo-Calc help.

 $get_mass_fraction_at_lower_interface(region: str, component: str) \rightarrow [typ-ing.List[float], typing.List[float]]$

Returns the mass fraction of the specified component at the lower boundary of the specified region, in dependency of time.

Parameters

- **region** The name of the region
- **component** The name of the component

Returns A tuple of two lists of floats (time [s], mass fraction of the specified component)

get_mass_fraction_at_upper_interface(region: str, component: str) \rightarrow [typing.List[float], typing.List[float]]

Returns the mass fraction of the specified component at the upper boundary of the specified region, in dependency of time.

Parameters

- **region** The name of the region
- component The name of the component

Returns A tuple of two lists of floats (time [s], mass fraction of the specified component)

<pre>get_mass_fraction_of_component_at_time (component:</pre>	str,	time:	
Union[tc_python.d	Union[tc_python.diffusion.SimulationTime,		
$float]) \rightarrow$	[typing.List[float],	typ-	
ing.List[float]]			
Paturns the mass fraction of the specified component at the specified tim	10		

Returns the mass fraction of the specified component at the specified time.

Note: Use the enum *tc_python.diffusion.SimulationTime* to choose the first or the last timepoint of the simulation. A timepoint close to the last one should never be specified manually because the actual end of the simulation can slightly deviate.

Parameters

- component The name of the component
- time The time [s]

Returns A tuple of two lists of floats (distance [m], mass fraction of component at the specified time)

<pre>get_mass_fraction_of_phase_at_time</pre>	(phase:	str,	time:
	Union[tc_python.diffus	ion.SimulationTime,	float])
Deturns the mass fraction of the encoding the	\rightarrow [typing.List[float], ty	ping.List[float]]	

Returns the mass fraction of the specified phase.

Note: Use the enum tc python.diffusion.SimulationTime to choose the first or the last timepoint of the simulation. A timepoint close to the last one should never be specified manually because the actual end of the simulation can slightly deviate.

Parameters

- **phase** The name of the phase
- time The time [s]

Returns A tuple of two lists of floats (distance [m], mass fraction of hte phase at the specified time)

```
get_mole_fraction_at_lower_interface(region:
                                                             str, component:
                                                                                  str)
                                                                                        \rightarrow [typ-
                                                   ing.List[float], typing.List[float]]
```

Returns the mole fraction of the specified component at the lower boundary of the specified region, in dependency of time.

Parameters

- region The name of the region
- component The name of the component

Returns A tuple of two lists of floats (time [s], mole fraction of the specified component)

get_mole_fraction_at_upper_interface (region: str, component: str) \rightarrow [typ-

ing.List[float], typing.List[float]] Returns the mole fraction of the specified component at the upper boundary of the specified region, in

dependency of time.

Parameters

- **region** The name of the region
- component The name of the component

Returns A tuple of two lists of floats (time [s], mole fraction of the specified component)

<pre>get_mole_fraction_of_component_at_time(component_at_time)</pre>	componen	<i>t</i> :	str,	time:
U	Union[tc_python.diffusion.SimulationTime,			me,
fl	loat])	\rightarrow	[typing.List[float],	typ-
	ng.List[fl			
Returns the mole fraction of the specified component a	at the spe	cified ti	me.	

Note: Use the enum *tc_python.diffusion.SimulationTime* to choose the first or the last timepoint of the simulation. A timepoint close to the last one should never be specified manually because the actual end of the simulation can slightly deviate.

Parameters

- component The name of the component
- time The time [s]
- **Returns** A tuple of two lists of floats (distance [m], mole fraction of component at the specified time)

<pre>get_mole_fraction_of_phase_at_time</pre>	(phase:	str,	time:
	Union[tc_python.dif	fusion.SimulationTime,	float])
Potums the mole fraction of the encodified phase	\rightarrow [typing.List[float]	, typing.List[float]]	

Returns the mole fraction of the specified phase.

Note: Use the enum *tc_python.diffusion.SimulationTime* to choose the first or the last timepoint of the simulation. A timepoint close to the last one should never be specified manually because the actual end of the simulation can slightly deviate.

Parameters

- **phase** The name of the phase
- time The time [s]
- **Returns** A tuple of two lists of floats (distance [m], mole fraction of the phase at the specified time)
- $\texttt{get_position_of_lower_boundary_of_region} (\textit{region: str}) \rightarrow [\texttt{typing.List[float]}, \texttt{typ-ing.List[float]}]$

Returns the position of the lower boundary of the specified region in dependency of time.

Parameters region – The name of the region

Returns A tuple of two lists of floats (time [s], position of lower boundary of region [m])

ing.List[float]]

 $\texttt{get_position_of_upper_boundary_of_region} (\textit{region: str}) \rightarrow [\texttt{typing.List[float]}, \texttt{typ-structure}) \rightarrow [\texttt{typing.List[float]}, \texttt{typing.List[float]}, \texttt{typing.List[float]}, \texttt{typing.List[float$

Returns the position of the upper boundary of the specified region in dependency of time.

Parameters region – The name of the region

Returns A tuple of two lists of floats (time [s], position of upper boundary of region [m])

$get_regions() \rightarrow List[str]$

Returns the regions of the diffusion simulation.

Note: Automatically generated regions $(R_{\#})$ are included in the list.

Returns The region names

```
get\_time\_steps() \rightarrow List[float]
```

Returns the timesteps of the diffusion simulation.

Returns The timesteps [s]

 $get_total_mass_fraction_of_component(component: str) \rightarrow [typing.List[float], typing.List[float]]$

Returns the total mass fraction of the specified component in dependency of time.

Parameters component – The name of the component

Returns A tuple of two lists of floats (time [s], total mass fraction of the component)

get_total_mass_fraction_of_component_in_phase (component: str, phase: str) → [typing.List[float], typing.List[float]]

Returns the total mass fraction of the specified component in the specified phase in dependency of time.

Parameters

- component The name of the component
- **phase** The name of the phase

Returns A tuple of two lists of floats (time [s], total mass fraction of the component in the phase)

get_total_mass_fraction_of_phase (*phase: str*) \rightarrow [typing.List[float], typing.List[float]] Returns the total mass fraction of the specified phase in dependency of the time.

Parameters phase – The name of the phase

Returns A tuple of two lists of floats (time [s], total mass fraction of the phase)

 $\texttt{get_total_mole_fraction_of_component}(\textit{component}: str) \rightarrow [\texttt{typing.List[float]}, \texttt{typ-index}) \rightarrow [\texttt{typing.List[float]}, \texttt{typ-index}) \rightarrow \texttt{for the started} \rightarrow \texttt{for the sta$

ing.List[float]]

Returns the total mole fraction of the specified component in dependency of time.

Parameters component – The name of the component

Returns A tuple of two lists of floats (time [s], total mole fraction of the component)

get_total_mole_fraction_of_component_in_phase (component: str, phase: str) → [typing.List[float], typing.List[float]]

Returns the total mole fraction of the specified component in the specified phase in dependency of time.

Parameters

- component The name of the component
- **phase** The name of the phase

Returns A tuple of two lists of floats (time [s], total mole fraction of the component in the phase)

get_total_mole_fraction_of_phase (*phase*: *str*) \rightarrow [typing.List[float], typing.List[float]] Returns the total mole fraction of the specified phase in dependency of time.

Parameters phase – The name of the phase

Returns A tuple of two lists of floats (time [s], total mole fraction of the phase)

get_total_volume_fraction_of_phase (*phase: str*) \rightarrow [typing.List[float], typing.List[float]] Returns the total volume fraction of the specified phase in dependency of the time.

Parameters phase – The name of the phase

Returns A tuple of two lists of floats (time [s], total volume fraction of the phase)

Returns the specified result from the simulation, allows all possible settings.

Note: As an alternative, DICTRA Console Mode syntax can be used as well for each quantity and condition.

Warning: This is an advanced mode that is equivalent to the possibilities in the DICTRA Console Mode. Not every combination of settings will return a result.

Parameters

- x_axis The first result quantity
- y_axis The second result quantity
- plot_condition The plot conditions
- independent_variable The independent variable

Returns A tuple of two lists of floats (the x_axis quantity result, the y_axis quantity result) [units according to the quantities]

 $get_velocity_of_lower_boundary_of_region (region: str) \rightarrow [typing.List[float], typing.List[float]]$

Returns the velocity of the lower boundary of the specified region in dependency of time.

Parameters region – The name of the region

Returns A tuple of two lists of floats (time [s], velocity of lower boundary of region [m/s])

 $\texttt{get_velocity_of_upper_boundary_of_region} (\textit{region: str}) \rightarrow [\texttt{typing.List[float]}, \texttt{typing.List[float]}]$

Returns the velocity of the upper boundary of the specified region in dependency of time.

Parameters region – The name of the region

Returns A tuple of two lists of floats (time [s], velocity of upper boundary of region [m/s])

get_width_of_region (*region: str*) \rightarrow [typing.List[float], typing.List[float]] Returns the width of region, in dependency of time.

Parameters region – The name of the region

Returns A tuple of two lists of floats (time [s], width of the specified region [m])

save_to_disk (path: str)

Saves the result to disk. The result can later be loaded using tc_python.server.SetUp. load_result_from_disk().

Note: The result data is represented by a whole folder containing multiple files.

Parameters path – The path to the result folder, can be relative or absolute.

Returns This DiffusionCalculationResult object

with_continued_calculation()

Returns a *ContinuedDiffusionCalculation* that is used for continuing a diffusion calculation with altered settings.

Returns A ContinuedDiffusionCalculation

class tc_python.diffusion.**DiffusionIsoThermalCalculation**(*calculation*) Bases: tc_python.abstract_base.AbstractCalculation

Configuration for an isothermal diffusion calculation.

add_console_command(console_command: str)

Registers a DICTRA Console Mode command for execution. These commands are executed after all other configuration directly before the calculation starts to run. All commands are stored and used until explicitly deleted using tc_python.diffusion.DiffusionIsoThermoCalculation.remove_all_console_commands.

Parameters console_command - The DICTRA Console Mode command

Returns This DiffusionIsoThermalCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw DICTRA-commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten parenthesis, ...).

add_region (region: tc_python.diffusion.Region)

Adds a region to the calculation. Regions are always added in the simulation domain from left to right.

If you want to replace an already added region, call *remove_all_regions()*, and add the regions that you want to keep.

Warning: Regions must have unique names.

Parameters region – The region to be added

Returns This DiffusionIsoThermalCalculation object

- **calculate** (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.diffusion.DiffusionCalculationResult* Runs the diffusion calculation.
 - **Parameters timeout_in_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.
 - **Returns** A *DiffusionCalculationResult* which later can be used to get specific values from the calculated result

get_system_data() \rightarrow tc_python.abstract_base.SystemData

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. *user*) databases loaded as *.*tdb*-file.

Returns The system data

remove_all_console_commands()

Removes all previously added Console Mode commands.

Returns This DiffusionIsoThermalCalculation object

remove_all_regions()

Removes all previously added regions.

:return This DiffusionIsoThermalCalculation object

set_simulation_time (simulation_time: float)

Sets the simulation time.

Parameters simulation_time – The simulation time [s]

Returns This DiffusionIsoThermalCalculation object

set_temperature(temperature: float)

Sets the temperature for the isothermal simulation.

Parameters temperature – The temperature [K]

Returns This DiffusionIsoThermalCalculation object

with_cylindrical_geometry (first_interface_position: float = 0.0)
Sets geometry to cylindrical, corresponds to an infinitely long cylinder of a certain radius.

Default: A planar geometry

Note: With a cylindrical or spherical geometry, the system's zero coordinate (left boundary) is at the centre of the cylinder or sphere by default. By specifying the *first_interface_position*, a different leftmost coordinate can be defined. This allows to model a tube or a hollow sphere geometry. The highest coordinate (right boundary) is defined by the cylinder or sphere radius (i.e. by the width of all regions).

Parameters first_interface_position – The position of the left-most coordinate along the axis, only necessary for modeling a tube geometry [m]

Returns This DiffusionIsoThermalCalculation object

Default: A closed-system boundary condition.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Note: You can specify time-dependent boundary conditions by calling with_left_boundary_condition() many times, with different values of the "to" parameter.

Examples:

- *with_left_boundary_condition(BoundaryCondition.closed_system(), to=100)*
- with_left_boundary_condition(BoundaryCondition.mixed_zero_flux_and_activity().set_activity_for_element("C", surface_activity), to=500)
- with_left_boundary_condition(BoundaryCondition.closed_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

Parameters

- boundary_condition The boundary condition
- to The upper time-limit for boundary_condition.

Returns This DiffusionIsoThermalCalculation object

with_options (*options*: tc_python.diffusion.Options, *to: float* = 1.7976931348623157e+308) Sets the general simulation conditions.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Parameters

- options The general simulation conditions
- to The upper time-limit for options.

Returns This DiffusionIsoThermalCalculation object

with_planar_geometry()

Sets geometry to *planar*.

This is default.

Returns This DiffusionIsoThermalCalculation object

with_reference_state (element: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

- element The name of the element
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- pressure The pressure (in Pa) for the reference state

Returns This DiffusionIsoThermalCalculation object

Defines the boundary condition on the right edge of the system.

Default: A closed-system boundary condition

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Note: You can specify time-dependent boundary conditions by calling with_right_boundary_condition() many times, with different values of the "to" parameter.

Examples:

- *with_right_boundary_condition(BoundaryCondition.closed_system(), to=100)*
- with_right_boundary_condition(BoundaryCondition.mixed_zero_flux_and_activity().set_activity_for_element("C", surface_activity), to=500)
- with_right_boundary_condition(BoundaryCondition.closed_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

Parameters

• **boundary_condition** – The boundary condition

• to – The upper time-limit for boundary_condition.

Returns This DiffusionIsoThermalCalculation object

with_solver (solver: tc_python.diffusion.Solver, to: float = 1.7976931348623157e+308)
Sets the solver to use (Classic, Homogenization or Automatic). Default is Automatic.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Parameters

- **solver** The solver to use
- to The upper time-limit for solver.

Returns This DiffusionIsoThermalCalculation object

with_spherical_geometry (first_interface_position: float = 0.0)

Sets geometry to *spherical*, corresponds to a sphere with a certain radius.

Default: A spherical geometry

Note: With a cylindrical or spherical geometry, the system's zero coordinate (left boundary) is at the centre of the cylinder or sphere by default. By specifying the *first_interface_position*, a different leftmost coordinate can be defined. This allows to model a tube or a hollow sphere geometry. The highest coordinate (right boundary) is defined by the cylinder or sphere radius (i.e. by the width of all regions).

Parameters first_interface_position – The position of the left-most coordinate along the axis, only necessary for modeling a hollow sphere geometry [m]

Returns This DiffusionIsoThermalCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)
 Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a * .tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This DiffusionIsoThermalCalculation object

Sets the timestep control options.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Parameters

- timestep_control The new timestep control options
- to The upper time-limit for timestep_control.

Returns This DiffusionIsoThermalCalculation object

class tc_python.diffusion.**DiffusionNonIsoThermalCalculation**(*calculation*) Bases: tc_python.abstract_base.AbstractCalculation

Configuration for a non-isothermal diffusion calculation.

add_console_command(console_command: str)

Registers a DICTRA Console Mode command for execution. These commands are executed after all other configuration directly before the calculation starts to run. All commands are stored and used until explicitly deleted using *tc_python.diffusion.DiffusionNonIsoThermalCalculation*. *remove all console commands*.

Parameters console_command - The DICTRA Console Mode command

Returns This DiffusionNonIsoThermalCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw DICTRA-commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten parenthesis, ...).

add_region (region: tc_python.diffusion.Region)

Adds a region to the calculation. Regions are always added in the simulation domain from left to right.

If you want to replace an already added region, call *remove_all_regions()*, and add the regions that you want to keep.

Warning: Regions must have unique names.

Parameters region – The region to be added

Returns This DiffusionNonIsoThermalCalculation object

- **calculate** (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.diffusion.DiffusionCalculationResult* Runs the diffusion calculation.
 - **Parameters timeout_in_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.
 - **Returns** A *DiffusionCalculationResult* which later can be used to get specific values from the calculated result

 $get_system_data() \rightarrow tc_python.abstract_base.SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

remove_all_console_commands()

Removes all previously added Console Mode commands.

Returns This DiffusionNonIsoThermalCalculation object

remove_all_regions()

Removes all previously added regions.

Returns This DiffusionNonIsoThermalCalculation object

set_simulation_time (simulation_time: float)
Sets the simulation time.

Parameters simulation_time – The simulation time [s]

Returns This DiffusionNonIsoThermalCalculation object

with_cylindrical_geometry (first_interface_position: float = 0.0)

Sets geometry to cylindrical, corresponds to an infinitely long cylinder of a certain radius.

Default: A planar geometry

Note: With a cylindrical or spherical geometry, the system's zero coordinate (left boundary) is at the centre of the cylinder or sphere by default. By specifying the *first_interface_position*, a different leftmost coordinate can be defined. This allows to model a tube or a hollow sphere geometry. The highest coordinate (right boundary) is defined by the cylinder or sphere radius (i.e. by the width of all regions).

Parameters first_interface_position – The position of the left-most coordinate along the axis, only necessary for modeling a tube geometry [m]

Returns This DiffusionNonIsoThermalCalculation object

Defines the boundary condition on the left edge of the system.

Default: A closed-system boundary condition.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Note: You can specify time-dependent boundary conditions by calling with_left_boundary_condition() many times, with different values of the "to" parameter.

Examples:

- with_left_boundary_condition(BoundaryCondition.closed_system(), to=100)
- with_left_boundary_condition(BoundaryCondition.mixed_zero_flux_and_activity().set_activity_for_element("C", surface_activity), to=500)
- with_left_boundary_condition(BoundaryCondition.closed_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

Parameters

- boundary_condition The boundary condition
- to The upper time-limit for boundary_condition.

Returns This DiffusionNonIsoThermalCalculation object

with_options (*options*: tc_python.diffusion.Options, to: float = 1.7976931348623157e+308) Sets the general simulation conditions.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Parameters

- options The general simulation conditions
- to The upper time-limit for options.

Returns This DiffusionNonIsoThermalCalculation object

with_planar_geometry()

Sets geometry to planar.

This is default.

Returns This DiffusionNonIsoThermalCalculation object

with_reference_state (element: str, phase: str = 'SER', temperature: float = - 1.0, pressure: float

```
= 100000.0)
```

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

- element The name of the element
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- pressure The pressure (in Pa) for the reference state

Returns This DiffusionNonIsoThermalCalculation object

Defines the boundary condition on the right edge of the system.

Default: A closed-system boundary condition

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Note: You can specify time-dependent boundary conditions by calling with_right_boundary_condition() many times, with different values of the "to" parameter.

Examples:

- *with_right_boundary_condition(BoundaryCondition.closed_system(), to=100)*
- with_right_boundary_condition(BoundaryCondition.mixed_zero_flux_and_activity().set_activity_for_element("C", surface_activity), to=500)
- with_right_boundary_condition(BoundaryCondition.closed_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

Parameters

- boundary_condition The boundary condition
- to The upper time-limit for boundary_condition.

Returns This DiffusionNonIsoThermalCalculation object

with_solver (solver: tc_python.diffusion.Solver, to: float = 1.7976931348623157e+308)
Sets the solver to use (Classic, Homogenization or Automatic). Default is Automatic.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Parameters

- **solver** The solver to use
- to The upper time-limit for solver.

Returns This DiffusionNonIsoThermalCalculation object

with_spherical_geometry (first_interface_position: float = 0.0)

Sets geometry to *spherical*, corresponds to a sphere with a certain radius.

Default: A spherical geometry

Note: With a cylindrical or spherical geometry, the system's zero coordinate (left boundary) is at the centre of the cylinder or sphere by default. By specifying the *first_interface_position*, a different leftmost coordinate can be defined. This allows to model a tube or a hollow sphere geometry. The highest coordinate (right boundary) is defined by the cylinder or sphere radius (i.e. by the width of all regions).

Parameters first_interface_position – The position of the left-most coordinate along the axis, only necessary for modeling a hollow sphere geometry [m]

Returns This DiffusionNonIsoThermalCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)
 Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. user) databases loaded as a *.tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This DiffusionNonIsoThermalCalculation object

- with_temperature_profile (temperature_profile: tc_python.utils.TemperatureProfile)
 Sets the temperature profile to use with this calculation.
 - **Parameters temperature_profile** The temperature profile object (specifying time / temperature points)

Returns This DiffusionNonIsoThermalCalculation object

Sets the timestep control options.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

Parameters

- timestep_control The new timestep control options
- to The upper time-limit for timestep_control.

Returns This DiffusionNonIsoThermalCalculation object

class	<pre>tc_python.diffusion.DoubleGeometricGrid(no_of_points:</pre>	int	=	50,
	lower_geometrical_	factor:	float	=
	1.1, upper_geomet	rical_factor	r: floa	t =
_	0.9)			

Bases: *tc_python.diffusion.CalculatedGrid*

Represents a double geometric grid.

get_lower_geometrical_factor() \rightarrow float Returns the lower geometrical factor (for the left half).

Returns The lower geometrical factor

```
\begin{array}{l} \texttt{get\_no\_of\_points}() \rightarrow \text{int} \\ \text{Returns number of grid points.} \end{array}
```

Returns The number of grid points

 $\begin{array}{l} \texttt{get_type}() \rightarrow str\\ Type of the grid. \end{array}$

Returns The type of the grid

```
get_upper_geometrical_factor()
```

Returns the upper geometrical factor (for the right half).

Returns The upper geometrical factor

set_lower_geometrical_factor (geometrical_factor: float = 1.1)

Sets the lower (left half) geometrical factor.

Note: A geometrical factor of larger than one yields a higher density of grid points at the lower end of the half and vice versa for a factor smaller than one.

Parameters geometrical_factor - The geometrical factor for the left half

Returns This DoubleGeometricGrid object

set_no_of_points (no_of_points: int = 50)
Sets the number of grid points.

Parameters no_of_points - The number of points

Returns This DoubleGeometricGrid object

set_upper_geometrical_factor (*geometrical_factor: float = 0.9*) Sets the upper (right half) geometrical factor.

Note: A geometrical factor of larger than one yields a higher density of grid points at the lower end of the half and vice versa for a factor smaller than one.

Parameters geometrical_factor - The geometrical factor for the right half

Returns This DoubleGeometricGrid object

class tc_python.diffusion.ElementProfile

Bases: tc_python.diffusion.AbstractElementProfile

Factory class providing objects for configuring a step, function or linear initial concentration profile.

classmethod constant (value: float)

Factory method that creates a **new** constant initial concentration profile.

Parameters value – The constant composition in the region. [unit as defined in *CompositionProfile*].

Returns A new ConstantProfile object

classmethod funct (dictra_console_mode_function: str)

Factory method that creates a **new** initial concentration profile defined by a function in DICTRA Console Mode syntax.

Parameters dictra_console_mode_function – The function, expressed in DICTRA Console Mode syntax.

Returns A new FunctionProfile object

Note: This is an advanced feature, preferably a complex concentration profile should be generated using third party libraries and added to the simulation using *tc_python.diffusion*. *PointByPointGrid*.

classmethod linear(start_value: float, end_value: float)

Factory method that creates a **new** linear initial concentration profile.

Parameters

- **start_value** Composition at the left side of the region [unit as defined in *CompositionProfile*].
- **end_value** Composition at the right side of the region [unit as defined in *CompositionProfile*].

Returns A new LinearProfile object

classmethod step (*lower_boundary: float, upper_boundary: float, step_at: float*)

Factory method that creates a **new** initial concentration profile with a step at the specified distance, otherwise the composition is constant at the specified values.

Parameters

- **lower_boundary** Composition before the step [unit as defined in *CompositionProfile*].
- **upper_boundary** Composition after the step [unit as defined in *CompositionProfile*].
- **step_at** The distance where the step should be [m].

Returns A new *StepProfile* object

class tc_python.diffusion.FixFluxValue
 Bases: tc_python.diffusion.BoundaryCondition

get_type() \rightarrow str

The type of the boundary condition.

Returns The type

set_flux (element_name: str, J: str = '0', to_time: float = 1.7976931348623157e+308)
Enter functions that yield the flux times the molar volume for the specified element. May be a function of
time, temperature and pressure: J(T,P,TIME).

Parameters

- element_name The name of the element
- **J** the function *J*(*T*,*P*,*TIME*)
- to_time The max-time for which the flux function is used.

Bases: tc_python.diffusion.BoundaryCondition

Represents a boundary having fixed composition values.

get_type () \rightarrow str The type of the boundary condition.

Returns The type

set_composition (*element_name: str, value: float*) Sets the composition for the specified element.

Note: The boundary composition needs to be specified for each element.

Parameters

- element_name The name of the element
- **value** The composition value [unit according to the constructor parameter]

```
class tc_python.diffusion.FunctionProfile (dictra_console_mode_function: str)
Bases: tc python.diffusion.ElementProfile
```

Creates an initial concentration profile defined by a function in DICTRA Console Mode syntax.

Note: This is an advanced feature, preferably a complex concentration profile should be generated using third party libraries and added to the simulation using *tc_python.diffusion.PointByPointGrid*.

get_type() \rightarrow str

The type of the element profile.

Returns The type

class tc_python.diffusion.GeneralLowerHashinShtrikman

Bases: tc_python.diffusion.HomogenizationFunctions

General lower Hashin-Shtrikman bounds: the outermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

class tc_python.diffusion.GeneralLowerHashinShtrikmanExcludedPhase(excluded_phases:

List[str] =

[])

Bases: tc_python.diffusion.HomogenizationFunctions

General lower Hashin-Shtrikman bounds: the outermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

class tc_python.diffusion.GeneralUpperHashinShtrikman

 $Bases: \verb+tc_python.diffusion.HomogenizationFunctions$

General upper Hashin-Shtrikman bounds: the innermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

Bases: tc_python.diffusion.HomogenizationFunctions

General upper Hashin-Shtrikman bounds: the innermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

[])

class tc_python.diffusion.**GeometricGrid** (*no_of_points: int = 50, geometrical_factor: float = 1.1*)

Bases: tc_python.diffusion.CalculatedGrid

Represents a geometric grid.

get_geometrical_factor() \rightarrow float Returns the geometrical factor.

Returns The geometrical factor

get_no_of_points () \rightarrow int Returns the number of grid points.

Returns The number of grid points

 $\begin{array}{l} \texttt{get_type} () \rightarrow \textit{str} \\ \text{Returns the type of grid.} \end{array}$

Returns The type

set_geometrical_factor (*geometrical_factor: float* = 1.1) Sets the geometrical factor.

Note: A geometrical factor larger than one yields a higher density of grid points at the lower end of the region and a factor smaller than one yields a higher density of grid points at the upper end of the region.

Parameters geometrical_factor - The geometrical factor

Returns This GeometricGrid object

set_no_of_points (no_of_points: int = 50)

Sets the number of grid points.

Parameters no_of_points – The number of points

Returns This GeometricGrid object

class tc_python.diffusion.GridPoint(distance: float)
 Bases: object

Represents a grid point, this is used in combination with grids of the type *tc_python.diffusion*. *PointByPointGrid*.

add_composition (element: str, value: float)

Adds a composition for the specified element to the grid point.

Parameters

• element – The element

• value – The composition value [unit as defined for the grid]

Returns This GridPoint object

class tc_python.diffusion.HashinShtrikmanBoundMajority

Bases: tc_python.diffusion.HomogenizationFunctions

Hashin-Shtrikman bounds with majority phase as matrix phase: the outermost shell consists of the phase with the highest local volume fraction.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

Bases: tc_python.diffusion.HomogenizationFunctions

Hashin-Shtrikman bounds with majority phase as matrix phase: the outermost shell consists of the phase with the highest local volume fraction.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

class tc_python.diffusion.**HashinShtrikmanBoundPrescribed**(*matrix_phase: str*) Bases: tc_python.diffusion.HomogenizationFunctions

Hashin-Shtrikman bounds with prescribed phase as matrix phase: the outermost shell consists of the prescribed phase.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

class tc_python.diffusion.HashinShtrikmanBoundPrescribedExcludedPhase (matrix_phase:

```
str,
ex-
cluded_phases:
List[str]
= [])
```

Bases: tc_python.diffusion.HomogenizationFunctions

class tc_python.diffusion.HomogenizationFunction(value)

Bases: enum.Enum

Homogenization function used for the *homogenization solver*. Many homogenization functions are based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase. **Default**: *RULE_OF_MIXTURES* (i.e. upper Wiener bounds)

GENERAL_LOWER_HASHIN_SHTRIKMAN = 0

General lower Hashin-Shtrikman bounds: the outermost shell consists of the phase with the most sluggish kinetics.

```
GENERAL_UPPER_HASHIN_SHTRIKMAN = 1
```

General upper Hashin-Shtrikman bounds: the innermost shell consists of the phase with the most sluggish kinetics.

HASHIN_SHTRIKMAN_BOUND_MAJORITY = 2

Hashin-Shtrikman bounds with majority phase as matrix phase: the outermost shell consists of the phase with the highest local volume fraction.

INVERSE_RULE_OF_MIXTURES = 4

Lower Wiener bounds: the geometrical interpretation are continuous layers of each phase orthogonal to the direction of diffusion

$RULE_OF_MIXTURES = 3$

Upper Wiener bounds: the geometrical interpretation are continuous layers of each phase parallel with the direction of diffusion

class tc_python.diffusion.HomogenizationFunctions

Bases: object

classmethod general_lower_hashin_shtrikman()

Factory method that creates a **new** homogenization function of the type

GeneralLowerHashinShtrikman.

General lower Hashin-Shtrikman bounds: the outermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

Returns A new GeneralLowerHashinShtrikman object

 $\verb+classmethod general_lower_hashin_shtrikman_excluded_phase (excluded_phases:$

List[str] = []) Factory method that creates a **new** homogenization function of the type *GeneralLowerHashinShtrikmanExcludedPhase*.

General lower Hashin-Shtrikman bounds: the outermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase. The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

Parameters excluded_phases - The excluded phases

Returns A new GeneralLowerHashinShtrikmanExcludedPhase object

classmethod general_upper_hashin_shtrikman()

Factory method that creates a **new** homogenization function of the type *GeneralUpperHashinShtrikman*.

General upper Hashin-Shtrikman bounds: the innermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

Returns A new General Upper Hashin Shtrikman object

classmethod general_upper_hashin_shtrikman_excluded_phase (excluded_phases:

List[str] = []) Factory method that creates a **new** homogenization function of the type *GeneralUpperHashinShtrikmanExcludedPhase*.

General upper Hashin-Shtrikman bounds: the innermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase. The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

Parameters excluded_phases – The excluded phases

Returns A new General UpperHashinShtrikmanExcludedPhase object

classmethod hashin_shtrikman_bound_majority()

Factory method that creates a **new** homogenization function of the type *HashinShtrikmanBoundMajority*.

Hashin-Shtrikman bounds with majority phase as matrix phase: the outermost shell consists of the phase with the highest local volume fraction.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

Returns A new *HashinShtrikmanBoundMajority* object

classmethod hashin_shtrikman_bound_majority_excluded_phase (excluded_phases:

List[str] = [])

Factory method that creates a **new** homogenization function of the type HashinShtrikmanBoundMajorityExcludedPhase.

Hashin-Shtrikman bounds with majority phase as matrix phase: the outermost shell consists of the phase with the highest local volume fraction. Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase. The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

Parameters excluded_phases – The excluded phases

Returns A new HashinShtrikmanBoundMajorityExcludedPhase object

classmethod hashin_shtrikman_bound_prescribed(matrix_phase: str)

Factory method that creates a **new** homogenization function of the type *HashinShtrikmanBoundPrescribed*.

Hashin-Shtrikman bounds with prescribed phase as matrix phase: the outermost shell consists of the prescribed phase.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

Parameters matrix_phase – The matrix phase

Returns A new HashinShtrikmanBoundPrescribed object

classmethod hashin_shtrikman_bound_prescribed_excluded_phase(matrix_phase:

str. excluded phases: List[str] = []) Factory homogenization function method that creates а new of the type HashinShtrikmanBoundPrescribedExcludedPhase.

Hashin-Shtrikman bounds with prescribed phase as matrix phase: the outermost shell consists of the prescribed phase.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase. The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

Parameters

- matrix_phase The matrix phase
- excluded_phases The excluded phases

Returns A new HashinShtrikmanBoundPrescribedExcludedPhase object

classmethod inverse_rule_of_mixtures()

Factory method that creates a **new** homogenization function of the type InverseRuleOfMixtures.

Lower Wiener bounds: the geometrical interpretation are continuous layers of each phase orthogonal to the direction of diffusion.

Returns A new *InverseRuleOfMixtures* object

classmethod inverse_rule_of_mixtures_excluded_phase (excluded_phases: List[str]

Factory method that creates a **new** homogenization function of the type InverseRuleOfMixturesExcludedPhase.

Lower Wiener bounds: the geometrical interpretation are continuous layers of each phase orthogonal to the direction of diffusion. Excluded phases are not considered in the diffusion calculations.

Parameters excluded_phases – The excluded phases

Returns A new *InverseRuleOfMixturesExcludedPhase* object

classmethod labyrinth_factor_f(matrix_phase: str)

Factory method that creates a **new** homogenization function of the type LabyrinthFactorF.

The labyrinth factor functions implies that all diffusion takes place in a single continuous matrix phase. The impeding effect on diffusion by phases dispersed in the matrix phase is taken into account by multiplying the flux with the volume fraction of the matrix phase.

Parameters matrix_phase - The matrix phase

Returns A new LabyrinthFactorF object

classmethod labyrinth_factor_f2 (matrix_phase: str)

Factory method that creates a new homogenization function of the type LabyrinthFactorF2.

The labyrinth factor functions implies that all diffusion takes place in a single continuous matrix phase. The impeding effect on diffusion by phases dispersed in the matrix phase is taken into account by multiplying the flux with the volume fraction of the matrix phase squared.

Parameters matrix_phase - The matrix phase

Returns A new LabyrinthFactorF2 object

classmethod rule_of_mixtures()

Factory method that creates a **new** homogenization function of the type *RuleOfMixtures*.

Upper Wiener bounds: the geometrical interpretation are continuous layers of each phase parallel with the direction of diffusion.

Returns A new *RuleOfMixtures* object

classmethod rule_of_mixtures_excluded_phase(excluded_phases: List[str] = [])

Factory method that creates a **new** homogenization function of the type *RuleOfMixturesExcludedPhase*.

Upper Wiener bounds: the geometrical interpretation are continuous layers of each phase parallel with the direction of diffusion. Excluded phases are not considered in the diffusion calculations.

Parameters excluded_phases – The excluded phases

Returns A new *RuleOfMixturesExcludedPhase* object

class tc_python.diffusion.HomogenizationSolver

Bases: tc_python.diffusion.Solver

Solver using the Homogenization model.

Note: This solver always uses the homogenization model, even if all regions have only one phase. The solver is significantly slower than the Classic model. Use the *tc_python.diffusion.AutomaticSolver* instead if you do not need that behavior.

disable_global_minimization()

Disables global minimization to be used in equilibrium calculations. Default: Disabled

Note: In general, using global minimization **significantly increases the simulation time**, but there is also a significantly reduced risk for non-converged equilibrium calculations.

Returns A new HomogenizationSolver object

disable_interpolation_scheme()

Configures the simulation not use *any interpolation scheme*. **Default**: To use the *logarithmic interpolation scheme* with 10000 discretization steps

Note: The homogenization scheme can be switched on by using *with_linear_interpolation_scheme* or *with_logarithmic_interpolation_scheme*.

enable_global_minimization()

Enables global minimization to be used in equilibrium calculations. Default: Disabled

Note: In general, using global minimization **significantly increases the simulation time**, but there is also a significantly reduced risk for non-converged equilibrium calculations.

Returns A new *HomogenizationSolver* object

get_type() \rightarrow str

The type of solver.

Returns The type

set_fraction_of_free_memory_to_use (fraction: float)

Sets the maximum fraction of free physical memory to be used by the interpolation scheme. **Default**: 1 / 10 of the free physical memory

Parameters fraction – The maximum free physical memory fraction to be used

Returns A new HomogenizationSolver object

set_memory_to_use (memory_in_megabytes: float)

Sets the maximum physical memory in megabytes to be used by the interpolation scheme. **Default**: 1000 MBytes of the free physical memory

Parameters memory_in_megabytes - The maximum physical memory to be used

Returns A new HomogenizationSolver object

with_function (*homogenization_function:* tc_python.diffusion.HomogenizationFunctions) Sets the *homogenization function* used by the *homogenization model*.

> **Parameters homogenization_function** – The homogenization function used by the homogenization model

Returns A new HomogenizationSolver object

with_linear_interpolation_scheme (steps: int = 10000)

Configures the simulation to use the *linear interpolation scheme*. **Default**: To use the *logarithmic interpolation scheme* with 10000 discretization steps

Parameters steps - The number of discretization steps in each dimension

Returns A new HomogenizationSolver object

List[str] = [])

with_logarithmic_interpolation_scheme (steps: int = 10000)

Configures the simulation to use the *linear interpolation scheme*. **Default**: To use the *logarithmic interpolation scheme* with 10000 discretization steps

Parameters steps – The number of discretization steps in each dimension

Returns A new HomogenizationSolver object

class tc_python.diffusion.InverseRuleOfMixtures

Bases: tc_python.diffusion.HomogenizationFunctions

Lower Wiener bounds: the geometrical interpretation are continuous layers of each phase orthogonal to the direction of diffusion.

class tc_python.diffusion.InverseRuleOfMixturesExcludedPhase (excluded_phases:

Bases: tc_python.diffusion.HomogenizationFunctions

Lower Wiener bounds: the geometrical interpretation are continuous layers of each phase orthogonal to the direction of diffusion.

Excluded phases are not considered in the diffusion calculations.

class tc_python.diffusion.**LabyrinthFactorF** (*matrix_phase: str*) Bases: tc_python.diffusion.HomogenizationFunctions

The labyrinth factor functions implies that all diffusion takes place in a single continuous matrix phase. The impeding effect on diffusion by phases dispersed in the matrix phase is taken into account by multiplying the flux with the volume fraction of the matrix phase.

class tc_python.diffusion.**LabyrinthFactorF2** (*matrix_phase: str*) Bases: tc_python.diffusion.HomogenizationFunctions

The labyrinth factor functions implies that all diffusion takes place in a single continuous matrix phase. The impeding effect on diffusion by phases dispersed in the matrix phase is taken into account by multiplying the flux with the volume fraction of the matrix phase squared.

class tc_python.diffusion.**LinearGrid** (*no_of_points: int = 50*) Bases: tc_python.diffusion.CalculatedGrid

Represents an equally spaced grid.

 $\begin{array}{l} \texttt{get_no_of_points} \ (\) \ \rightarrow \ int \\ Returns \ the \ number \ of \ grid \ points. \end{array}$

Returns The number of grid points

 $\begin{array}{l} \texttt{get_type}() \rightarrow \text{str} \\ \text{Type of the grid.} \end{array}$

Returns The type

set_no_of_points (no_of_points: int = 50)
Sets the number of grid points.

Parameters no_of_points – The number of points

Returns This LinearGrid object

class tc_python.diffusion.**LinearProfile** (*start_value: float, end_value: float*) Bases: tc_python.diffusion.ElementProfile

Represents a linear initial concentration profile.

 $\begin{array}{l} \texttt{get_type} \ (\) \ \rightarrow \ \texttt{str} \\ \\ \text{The type of the element profile.} \end{array}$

Returns The type

class tc_python.diffusion.MixedZeroFluxAndActivity

Bases: tc_python.diffusion.BoundaryCondition

Represents a boundary having zero-flux as well as fixed-activity conditions.

Default: On that boundary for every element without an explicitly defined condition, a zero-flux boundary condition is used.

 $\texttt{get_type}() \rightarrow \text{str}$

The type of the boundary condition.

Returns The type

Sets an activity expression for an element at the boundary. Enter a formula that the software evaluates during the calculation.

The formula can be:

- a function of the variable TIME
- a constant

The formula must be written with these rules:

- a number must begin with a number (not a .)
- a number must have a dot or an exponent (*E*)

The operators +, -, *, /, ** (exponentiation) can be used and with any level of parenthesis. As shown, the following operators must be followed by open and closed parentheses ()

- *SQRT*(*X*) is the square root
- *EXP(X)* is the exponential
- *LOG(X)* is the natural logarithm
- *LOG10(X)* is the base 10 logarithm
- SIN(X), COS(X), TAN(X), ASIN(X), ACOS(X), ATAN(X)
- SINH(X), COSH(X), TANH(X), ASINH(X), ACOSH(X), ATANH(X)
- SIGN(X)
- *ERF*(*X*) is the error function

Default: the expression entered is used for the entire simulation.

Parameters

- **element_name** The name of the element
- **activity** The activity
- to_time The max-time for which the activity is used.

set_zero_flux_for_element (element_name: str)

Sets a zero-flux condition for an element at the boundary. **Default for all elements at the boundary** without an explicitly defined condition

Parameters element_name - The name of the element

class tc_python.diffusion.Options

```
Bases: object
```

General simulation conditions for the diffusion calculations.

disable_forced_starting_values_in_equilibrium_calculations()

Disables forced starting values for the equilibrium calculations. The default is 'enable_automatic_forced_starting_values_in_equilibrium_calculations'.

Returns This Options object

disable_save_results_to_file()

Disables the saving of results to file during the simulation. Default: Saving of the results at every timestep

Returns This Options object

enable_automatic_forced_starting_values_in_eq_calculations()

Lets calculation engine decide if forced start values for the equilibrium calculations should be used. **This is the default setting**.

Returns This Options object

enable_forced_starting_values_in_equilibrium_calculations()

Enables forced start values for the equilibrium calculations. The default is 'enable_automatic_forced_starting_values_in_equilibrium_calculations'.

Returns This Options object

enable_save_results_to_file (every_nth_step: int = - 1)

Enables and configures saving of results to file during the simulation. They can be saved for every n-th or optionally for every timestep (-1). **Default**: Saving of the results at every timestep

Parameters every_nth_step - -1 or a value ranging from 0 to 99

Returns This Options object

enable_time_integration_method_automatic()

Enables automatic selection of integration method. This is the default method.

Returns This Options object

enable_time_integration_method_euler_backwards()

Enables *Euler backwards* integration. **The default method is en**able_time_integration_method_automatic.

Note: This method is more stable but less accurate and may be necessary if large fluctuations occur in the profiles.

Returns This Options object

enable_time_integration_method_trapezoidal()

Enables *trapezoidal* integration.

Note: If large fluctuations occur in the profiles, it may be necessary to use the more stable but less accurate *Euler backwards method*.

Returns This Options object

```
set default driving force for phases allowed to form at interf (driving force:
                                                                       float = 1e-
                                                                       05)
```

Sets the default required driving force for phases allowed to form at the interfaces. Default: 1.0e-5

Note: The required driving force (evaluated as DGM(ph)) is used for determining whether an inactive phase is stable, i.e. actually formed. DGM represents the driving force normalized by RT and is dimensionless.

Parameters driving_force – The driving force (*DGM*(*ph*)) [-]

Returns This Options object

```
class tc_python.diffusion.PointByPointGrid(unit_enum:
                                                              tc_python.diffusion.Unit
                                                                                    =
                                                 <Unit.MASS PERCENT: 3>)
```

Bases: tc python.diffusion.AbstractGrid

Represents a point-by-point grid. This is setting the grid and the compositions at once, it is typically used to enter a measured composition profile or the result from a previous calculation.

Note: If a point-by-point grid is used, it is not necessary to specify the grid and composition profile separately.

add_point (grid_point: tc_python.diffusion.GridPoint) Adds a grid point to the grid.

Parameters grid_point – The grid point

Returns This PointByPointGrid object

get_type() \rightarrow str Type of the grid.

Returns The type

class tc_python.diffusion.Region(name: str) Bases: object

Represents a region of the simulation domain that can contain more that one phase.

Note: The first added phase represents the matrix phase, while all later added phases are *spheriod phases*, i.e. precipitate phases.

add_phase (*phase_name: str, is_matrix_phase: bool = False*)

Adds a phase to the region, each region must contain at least one phase.

Note: Normally the matrix phase and the precipitate phases are automatically chosen based on the presence of all profile elements in the phase and if it has diffusion data. If multiple phases have equal properties, the phase that was added first is chosen. The matrix phase can be explicitly set by using is matrix phase=True.

Note: If multiple phases are added to a region, the *homogenization model* is applied. That means that average properties of the local phase mixture are used.

Parameters

- **phase_name** The phase name
- **is_matrix_phase** If set to *True* this phase is explicitly set as matrix phase for the region, if no phase is set to *True*, the matrix phase is chosen automatically

Returns This Region object

add_phase_allowed_to_form_at_left_interface (phase_name: str, driving_force: float =

1e-05)

Adds a phase allowed to form at the left boundary of the region (an *inactive phase*). The phase will only appear at the interface as a new automatic region if the driving force to form it is sufficiently high.

Parameters

- phase_name The phase name
- **driving_force** The driving force for the phase to form (*DGM*(*ph*))

Returns This *Region* object

```
add_phase_allowed_to_form_at_right_interface (phase_name: str, driving_force: float
= 1e-05)
```

Adds a phase allowed to form at the right boundary of the region (an *inactive phase*). The phase will only appear at the interface as a new automatic region if the driving force to form it is sufficiently high.

Parameters

- phase_name The phase name
- **driving_force** The driving force for the phase to form (*DGM*(*ph*))

Returns This Region object

remove_all_phases()

Removes all previously added phases from the region.

Returns This *Region* object

set_width(width: float)

Defined the width of the region.

Note: This method needs only to be used if a calculated grid has been defined (using with_grid()).

Parameters width – The width [m]

Returns This Region object

with_composition_profile (initial_compositions: tc_python.diffusion.CompositionProfile)
 Defines the initial composition profiles for all elements in the region.

Note: This method needs only to be used if a calculated grid has been defined (using with_grid()).

Parameters initial_compositions – The initial composition profiles for all elements **Returns** This *Region* object

with_grid (grid: tc_python.diffusion.CalculatedGrid)

Defines a calculated grid in the region. If measured composition profiles or the result from a previous calculation should be used, instead with_point_by_point_grid_containing_compositions() needs to be applied.

Note: The composition profiles need to be defined separately using with_composition_profile(), additionally the region width needs to be specified using set_width().

Parameters grid – The grid

Returns This Region object

with_point_by_point_grid_containing_compositions (grid:

tc_python.diffusion.PointByPointGrid) Defines a point-by-point grid in the region. This is setting the grid and the compositions at once, it is typically used to enter a measured composition profile or the result from a previous calculation. If the composition profile should be calculated (linear, geometric, ...) with_grid() should be used instead.

Note: If a point-by-point grid is used, with_grid(), with_composition_profile() and set_width() are unnecessary and must not be used.

Parameters grid – The point-by-point grid

Returns This Region object

class tc_python.diffusion.RuleOfMixtures

Bases: tc_python.diffusion.HomogenizationFunctions

Upper Wiener bounds: the geometrical interpretation are continuous layers of each phase parallel with the direction of diffusion.

```
class tc_python.diffusion.RuleOfMixturesExcludedPhase(excluded_phases: List[str] =
```

Bases: tc_python.diffusion.HomogenizationFunctions

Upper Wiener bounds: the geometrical interpretation are continuous layers of each phase parallel with the direction of diffusion.

[1)

Excluded phases are not considered in the diffusion calculations.

class tc_python.diffusion.SimulationTime(value)
 Bases: enum.Enum

Specifying special time steps for the evaluation of diffusion results.

Note: These placeholders should be used because especially the actual last timestep will slightly differ from the specified end time of the simulation.

FIRST = 0

Represents the first timestep of the simulation

LAST = 1

Represents the last timestep of the simulation

class tc_python.diffusion.Solver

Bases: $tc_python.diffusion.AbstractSolver$

Factory class providing objects representing a solver.

classmethod automatic()

Factory method that creates a new *automatic solver*. This is the default solver and recommended for most applications.

Note: This solver uses the homogenization model if any region has more than one phase, otherwise it uses the classic model.

Returns A new AutomaticSolver object

classmethod classic()

Factory method that creates a **new** classic solver.

Note: This solver never switches to the homogenization model even if the solver fails to converge. Use the *tc_python.diffusion.AutomaticSolver* if necessary instead.

Returns A new *ClassicSolver* object

classmethod homogenization()

Factory method that creates a **new** homogenization solver.

Note: This solver always uses the homogenization model, even if all regions have only one phase. The solver is **significantly slower than the Classic model**. Use the *tc_python.diffusion*. *AutomaticSolver* instead if you do not need that behavior.

Returns A new HomogenizationSolver object

```
class tc_python.diffusion.StepProfile(lower_boundary: float, upper_boundary: float,
```

step_at: float)
Bases: tc python.diffusion.ElementProfile

Represents an initial constant concentration profile with a step at the specified position.

 $\texttt{get_type}() \rightarrow str$ The type of the element profile.

Returns The type

```
class tc_python.diffusion.TimestepControl
```

Bases: object

Settings that control the time steps in the simulation.

disable_check_interface_position()

Disables checking of the interface position, i.e. the timesteps are not controlled by the phase interface displacement during the simulation. The default setting is :func:`enable_automatic_check_interface_position`.

Returns This *TimestepControl* object

enable_automatic_check_interface_position()

Lets calculation engine decide if checking of the interface position should be used. **This is the default setting**.

Returns This TimestepControl object

enable_check_interface_position()

Enables checking of the interface position, i.e. the timesteps are controlled by the phase interface displacement during the simulation. **The default setting is :func:`enable_automatic_check_interface_position`**.

Returns This *TimestepControl* object

set_initial_time_step (initial_time_step: float = 1e-07)
Sets the initial timestep. Default: 1.0e-7 s

Parameters initial_time_step – The initial timestep [s]

Returns This TimestepControl object

set_max_absolute_error (absolute_error: float = 1e-05)
Sets the maximum absolute error. Default: 1.0e-5

Parameters absolute_error - The maximum absolute error

Returns This *TimestepControl* object

set_max_relative_error (*relative_error*: *float* = 0.05) Sets the maximum relative error. **Default**: 0.05

Parameters relative_error - The maximum relative error

Returns This TimestepControl object

```
set_max_timestep_allowed_as_percent_of_simulation_time (max_timestep_allowed_as_percent_of_simulat
```

float = 10.0)

The maximum timestep allowed during the simulation, specified in percent of the simulation time. **Default**: 10.0%

Parameters max_timestep_allowed_as_percent_of_simulation_time - The maximum timestep allowed [%]

Returns This *TimestepControl* object

set_max_timestep_increase_factor (max_timestep_increase_factor: float = 2.0)
Sets the maximum timestep increase factor. Default: 2

Note: For example, if 2 is entered the maximum time step is twice as long as the previous time step taken.

Parameters max_timestep_increase_factor – The maximum timestep increase factor

Returns This *TimestepControl* object

set_smallest_time_step_allowed(smallest_time_step_allowed: float = 1e-07)

Sets the smallest time step allowed during the simulation. This is required when using the automatic procedure to determine the time step. **Default**: 1.0e-7 s

Parameters smallest_time_step_allowed – The smalles timestep allowed [s]

Returns This *TimestepControl* object

class tc_python.diffusion.**Unit**(*value*) Bases: enum.Enum Represents a composition unit.

MASS_FRACTION = 2 Mass fraction.

MASS_PERCENT = 3 Mass percent.

MOLE_FRACTION = 0 Mole fraction.

MOLE_PERCENT = 1 Mole percent.

U_FRACTION = 4 U fraction

5.1.7 Module "propertymodel"

class tc_python.propertymodel.**PropertyModelCalculation** (*calculator*) Bases: tc_python.abstract_base.AbstractCalculation

Configuration for a Property Model calculation.

Note: Specify the settings, the calculation is performed with calculate().

```
add_poly_command (poly_command: str)
```

Registers a POLY Console Mode command for execution. These commands are executed after all other configuration directly before the calculation starts to run. All commands are stored and used until explicitly deleted using *remove_all_poly_commands*.

Parameters poly_command - The POLY Console Mode command

Returns This *PropertyModelCalculation* object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw POLY-commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten parenthesis, ...).

calculate (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.propertymodel.PropertyModelResult* Runs the Property Model calculation.

- **Parameters timeout_in_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a CalculationEngineException will be thrown.
- **Returns** A *PropertyModelResult* which later can be used to get specific values from the simulation.

get_argument_default (*argument_id: str*) → object

Returns the default value for the specified argument. The argument id can be obtained with get_arguments().

Parameters argument_id - The argument id

Returns The default value (the type depends on the argument)

 $\texttt{get_argument_description} (\textit{argument_id: str}) \rightarrow \texttt{str}$

Returns the detailed description of the argument. The id can be obtained with get_arguments().

Parameters argument_id – The argument id

Returns The detailed description

 $\texttt{get}_\texttt{arguments}() \rightarrow \texttt{Set}[\texttt{str}]$

Returns a list of the arguments of the Property Model.

Note: The arguments are the 'UI-panel components' defined in the Property Model interface method provide_ui_panel_components(). They have the same id as specified in the Property Model. The naming is different because there is no UI present.

Returns The ids of the available arguments

```
get_dynamic_arguments() \rightarrow Set[str]
```

Returns a list of the dynamic arguments of the Property Model.

Note: Dynamic arguments are "extra" arguments created by pressing the "plus" button that can occur next to the UI-panel for some models, when running the Property Model from within Thermo-Calc. You can use them also from the API by *invoke_dynamic_argument()*.

Returns The ids of the available dynamic arguments

```
\texttt{get\_model\_description()} \rightarrow str
```

Returns the description text of the current model.

Returns the description

 $\texttt{get_model_parameter_value} (\textit{model_parameter_id: str}) \rightarrow \texttt{float}$

Returns the current value of an optimizable model parameter. The id can be obtained with get_model_parameters().

Parameters model_parameter_id - The model parameter id

Returns The current value [unit according to the parameter meaning]

```
get_model_parameters() \rightarrow Set[str]
```

Returns a list of the optimizable model parameters.

Note: The model parameters are an optional set of variables that can be used within the Property Model. Typically they are used to provide the possibility to inject parameter values during an optimization into the model. This allows the dynamic development of Property Models that need to be fitted to experimental data. The model parameters are controlled with the Property Model interface methods *provide_model_parameters* and *set_model_parameter*.

Returns The ids of the optimizable model parameters

$get_system_data() \rightarrow tc_python.abstract_base.SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

invoke_dynamic_argument (argument_id: str)

Increases the number of instances of this dynamic argument by one, the argument will have an id such as *argument_1*, *argument_2*, ... if the dynamic argument is called *argument*.

Note: You can obtain all available dynamic arguments by using get_dynamic_arguments().

Parameters argument_id - argument_id: The argument id

Returns This *PropertyModelCalculation* object

remove_all_conditions()

Removes all set classic POLY conditions.

Note: This does not affect the compositions set by set_composition().

Returns This PropertyModelCalculation object

remove_all_poly_commands()

Removes all previously added POLY Console Mode commands.

Returns This *PropertyModelCalculation* object

remove_dependent_element()

Removes a manually set dependent element. This method does not affect the automatic choice of the dependent element if set_composition() is used.

Returns This PropertyModelCalculation object

set_argument (argument: str, value: str)

Sets the specified model argument to the specified value. The id can be obtained with get_arguments().

Parameters

- **argument** The argument id
- **value** The value [unit according to the argument meaning]

Returns This PropertyModelCalculation object

set_composition (element_name: str, value: float)

Sets the composition of a element. The unit for the composition can be changed using set_composition_unit().

Default: Mole percent (CompositionUnit.MOLE_PERCENT)

Parameters

- element_name The element
- **value** The composition value [composition unit defined for the calculation]

Returns This PropertyModelCalculation object

```
set_composition_unit (unit_enum: tc_python.utils.CompositionUnit = <Compositio-
nUnit.MOLE PERCENT: 2>)
```

Sets the composition unit.

Default: Mole percent (CompositionUnit.MOLE_PERCENT).

Parameters unit_enum – The new composition unit

Returns This PropertyModelCalculation object

set_condition (classic_condition: str, value: float)

Adds a classic POLY condition. If that method is used, all conditions need to be specified in such a way. If this method is used, it is necessary to set the dependent element manually using set_dependent_element().

Default if not specified: pressure P = 1e5 Pa, system size N = 1, Temperature T = 1000 K

Warning: It is not possible to mix POLY-commands and compositions using set_composition().

Note: It should not be necessary for most users to use this method, try to use *set_composition()* instead.

Warning: As this method runs raw POLY-commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten parenthesis, ...).

Parameters

- **classic_condition** The classic POLY condition (for example: *X*(*CR*))
- **value** The value of the condition

Returns This PropertyModelCalculation object

set_dependent_element (dependent_element_name: str)
 Sets the dependent element manually.

Note: It should not be necessary for most users to use this method. Setting the dependent element manually is only necessary and allowed if *set_condition()* is used.

Parameters dependent_element_name – The name of the dependent element

Returns This PropertyModelCalculation object

set_model_parameter (model_parameter_id: str, value)

Resets an optimizable model parameter. The id can be obtained with get_model_parameters ().

Parameters

- model_parameter_id The model parameter id
- **value** The new value of the parameter

Returns This PropertyModelCalculation object

set_temperature (temperature: float = 1000)

Sets the temperature. **Default**: 1000 K

Parameters temperature – The temperature [K]

Returns This PropertyModelCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)
 Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. user) databases loaded as a * .tdb-file.

Parameters system_modifications - The system modification to be performed

Returns This *PropertyModelCalculation* object

```
class tc_python.propertymodel.PropertyModelResult(result)
```

Bases: tc_python.abstract_base.AbstractResult

The result of a Property Model calculation.

get_result_quantities() \rightarrow Set[str]

Returns a list of the available result quantities defined in the Property Model.

Returns The ids of the defined result quantities

get_result_quantity_description (*result_quantity_id*) \rightarrow str

Returns the detailed description of the result quantity. The id can be obtained by get_result_quantities().

Parameters result_quantity_id - The result quantity id

Returns The detailed description

 $get_single_equilibrium_result (result_quantity_id: str = ") \rightarrow tc_python.single_equilibrium.SingleEquilibriumResult$

Returns a result quantity value. The available result quantities can be obtained by get_result_quantities().

Parameters result_quantity_id – The id of the result quantity.

Returns The requested value [unit depending on the quantity], if the result is a SingleEquilibriumResult, is returned.

 $\texttt{get_value_of} (\textit{result_quantity_id: str}) \rightarrow \texttt{Union[float, Dict[str, float]]}$

Returns a result quantity value. The available result quantities can be obtained by get_result_quantities().

Parameters result_quantity_id – The id of the result quantity

Returns The requested value [unit depending on the quantity]. If the result is parameterized, parameter-value pairs are returned.

save_to_disk (path: str)

Saves the result to disk. The result can later be loaded using tc_python.server.SetUp. load_result_from_disk().

Note: The *result data* is represented by a whole folder possibly containing multiple files.

Parameters path – The path to the result folder, can be relative or absolute.

Returns This *PropertyModelResult* object

5.1.8 Module "material_to_material"

class tc_python.material_to_material.AbstractConstantCondition
 Bases: object

The abstract base class for all constant conditions.

class tc_python.material_to_material.AbstractMaterialToMaterialCalculationAxis
 Bases: object

The abstract base class of all calculation axis.

class tc_python.material_to_material.ConstantCondition
 Bases: tc_python.material_to_material.AbstractConstantCondition

A constant condition.

classmethod fraction_of_material_b (*fraction_of_material_b: float = 0.5*) Creates a constant fraction of material B condition object.

Note: The unit depends on the composition unit setting in the calculator object.

Parameters fraction_of_material_b – The fraction of material B [weight-fraction or mole-fraction]

Returns The condition object

```
classmethod temperature (temperature: float = 1000)
Creates a constant temperature condition object.
```

Parameters temperature – The temperature [K]

Returns The condition object

class tc_python.material_to_material.FractionOfMaterialBAxis (from_fraction:

float = 0.0, to_fraction: float = 1.0, start_fraction: float = 0.5) Bases: tc_python.material_to_material.MaterialToMaterialCalculationAxis

A fraction of material B axis.

Bases: tc_python.material_to_material.ConstantCondition

A constant fraction of material B condition.

class tc_python.material_to_material.MaterialToMaterialCalculationAxis

Bases: tc_python.material_to_material.AbstractMaterialToMaterialCalculationAxis

A calculation axis.

```
classmethod fraction_of_material_b (from_fraction: float = 0.0, to_fraction: float = 1.0,
```

Start_fraction: float = 0.5) Creates a fraction of material B axis object.

Note: The unit depends on the composition unit setting in the calculator.

Parameters

- from_fraction The left axis limit [weight-fraction or mole-fraction]
- to_fraction The right axis limit [weight-fraction or mole-fraction]
- **start_fraction** The start fraction of the calculation [weight-fraction or mole-fraction]

Returns A new FractionOfMaterialBAxis axis object

```
classmethod temperature (from_temperature: float = 1000, to_temperature: float = 3000,
start_temperature: float = 2000)
```

Creates a temperature calculation axis object.

Parameters

- **from_temperature** The left axis limit [K]
- to_temperature The right axis limit [K]
- **start_temperature** The start temperature of the calculation [K]

Returns A new *TemperatureAxis* condition object

class tc_python.material_to_material.MaterialToMaterialCalculationContainer(instance)
 Bases: object

Provides access to the calculation objects for all Material to Material calculations.

These are specialised calculations for mixtures of two materials A and B. Otherwise they behave identical to the corresponding regular single equilibrium, property diagram and phase diagram calculations.

with_phase_diagram_calculation (default_conditions: bool = True, components: List[str] = []) \rightarrow $tc_python.material_to_material.MaterialToMaterialPhaseDiagramCalculation$

Creates a Material to Material phase diagram (map) calculation.

Parameters

- **default_conditions** If *True*, automatically sets the conditions N=1 and P=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

Returns A new MaterialToMaterialPhaseDiagramCalculation object

with_property_diagram_calculation (default_conditions: bool = True, components: List[str] = []) \rightarrow

```
tc python.material to material.MaterialToMaterialPropertyDiagramCalcula
```

Creates a Material to Material property diagram (step) calculation.

Parameters

- default_conditions If *True*, automatically sets the conditions N=1 and P=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

Returns A new MaterialToMaterialPropertyDiagramCalculation object

with_single_equilibrium_calculation(default_conditions: bool = True,

components: $List[str] = []) \rightarrow$

tc_python.material_to_material.MaterialToMaterialSingleEquilibriumCalc

Creates a Material to Material single equilibrium calculation.

Parameters

- **default_conditions** If *True*, automatically sets the conditions N=1 and P=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

Returns A new MaterialToMaterialSingleEquilibriumCalculation object

class tc_python.material_to_material.**MaterialToMaterialPhaseDiagramCalculation**(*calculator*) Bases: tc_python.step_or_map_diagrams.AbstractPhaseDiagramCalculation

Configuration for a Material to Material phase diagram calculation.

Note: Specify the conditions, the calculation is performed with *calculate()*.

add_initial_equilibrium (*initial_equilibrium*: tc_python.step_or_map_diagrams.InitialEquilibrium) Add initial equilibrium start points from which a phase diagram is calculated.

Scans along the axis variables and generates start points when the scan procedure crosses a phase boundary.

It may take a little longer to execute than using the minimum number of start points, as some lines may be calculated more than once. But the core remembers all node points and subsequently stops calculations along a line when it finds a known node point.

It is also possible to create a sequence of start points from one initial equilibria.

Parameters initial_equilibrium – The initial equilibrium

Returns This MaterialToMaterialPhaseDiagramCalculation object

calculate (keep_previous_results: bool = False, timeout_in_minutes: float = 0.0) \rightarrow tc_python.material_to_material.MaterialToMaterialPhaseDiagramResult Performs the phase diagram calculation.

Warning: If you use *keep_previous_results=True*, you must not use another calculator or even get results in between the calculations using *calculate()*. Then the previous results will actually be lost.

Parameters

- **keep_previous_results** If *True*, results from any previous call to this method are appended. This can be used to combine calculations with multiple start points if the mapping fails at a certain condition.
- timeout_in_minutes Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.
- **Returns** A new *MaterialToMaterialPhaseDiagramResult* object which later can be used to get specific values from the calculated result.

disable_global_minimization()

Disables global minimization.

Default: Enabled

Returns This MaterialToMaterialPhaseDiagramCalculation object

dont_keep_default_equilibria()

Do not keep the initial equilibria added by default.

This is only relevant in combination with add_initial_equilibrium().

This is the default behavior.

Returns This Material To Material Phase Diagram Calculation object

enable_global_minimization()

Enables global minimization.

Default: Enabled

Returns This Material To Material Phase Diagram Calculation object

get_components() \rightarrow List[str]

Returns the names of the components in the system (including all components auto-selected by the database(s)).

Returns The component names

get_gibbs_energy_addition_for (*phase: str*) \rightarrow float

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters phase – Specify the name of the (stoichiometric or solution) phase with the addition

Returns Gibbs energy addition to G per mole formula unit.

$get_system_data() \rightarrow tc_python.abstract_base.SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

keep_default_equilibria()

Keep the initial equilibria added by default. This is only relevant in combination with add_initial_equilibrium().

Default behavior is to not keep default equilibria.

Returns This MaterialToMaterialPhaseDiagramCalculation object

remove_all_initial_equilibria()

Removes all previously added initial equilibria.

Returns This MaterialToMaterialPhaseDiagramCalculation object

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

Parameters command - The Thermo-Calc Console Mode command

Returns This Material To Material Phase Diagram Calculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

set_activities (activities: Dict[str, float])

Sets the constant activity conditions.

Note: The activity conditions are identical for both materials.

Parameters activities – The constant activities

Returns This MaterialToMaterialPhaseDiagramCalculation object

set_composition_unit (unit: tc_python.utils.CompositionUnit = <CompositionUnit.MASS_PERCENT: 0>) Sets the composition unit of both materials A and B.

Default: Weight percent

Parameters unit – The composition unit of both materials A and B

Returns This Material To Material Phase Diagram Calculation object

set_gibbs_energy_addition_for (phase: str, gibbs_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs_energy Addition to G per mole formula unit

Returns This Material To Material Phase Diagram Calculation object

set_material_a (composition: Dict[str, float], dependent_component: str = None)
Sets the composition of the material A.

The unit is set with set_composition_unit().

Tip: The material can also have constant activity conditions, they are set in set_activities().

Parameters

- composition The composition of the material A
- dependent_component The dependent component of the material A

Returns This MaterialToMaterialPhaseDiagramCalculation object

set_material_b (*composition: Dict[str, float]*, *dependent_component: str = None*) Sets the composition of the material B.

The unit is set with set_composition_unit().

Tip: The material can also have constant activity conditions, they are set in set_activities().

Parameters

- composition The composition of the material B
- dependent_component The dependent component of the material B

Returns This MaterialToMaterialPhaseDiagramCalculation object

set_phase_to_dormant (phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This MaterialToMaterialPhaseDiagramCalculation object

set_phase_to_entered (phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

Parameters

- phase The phase name or ALL_PHASES for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

Returns This MaterialToMaterialPhaseDiagramCalculation object

set_phase_to_fixed (phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

Parameters

- **phase** The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

Returns This MaterialToMaterialPhaseDiagramCalculation object

set_phase_to_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters phase – The phase name or ALL_PHASES for all phases

Returns This MaterialToMaterialPhaseDiagramCalculation object

set_pressure (pressure: float)

Sets the pressure (i.e. the condition *P*).

Note: If the flag *default_conditions=True* has been set during the creation of the calculator, the pressure is set to 1000 hPa by default.

Parameters pressure – The pressure [Pa]

Returns This MaterialToMaterialPhaseDiagramCalculation object

set_system_size (system_size: float)

Sets the system size (i.e. the condition 'N', the number of moles).

Note: If the flag *default_conditions=True* has been set during the creation of the calculator, the system size is set to 1.0 moles by default.

Parameters system_size – The system size [mole]

Returns This MaterialToMaterialPhaseDiagramCalculation object

with_first_axis (axis: tc_python.material_to_material.MaterialToMaterialCalculationAxis)
Sets the first axis (either temperature of fraction of material B). This calculation type requires that both
temperature and fraction of material B axis are set.

Parameters axis – The axis

Returns This MaterialToMaterialPhaseDiagramCalculation object

with_options (*options*: tc_python.step_or_map_diagrams.PhaseDiagramOptions) Sets the simulation options.

Parameters options – The simulation options

Returns This PhaseDiagramCalculation object

with_reference_state (component: str, phase: str = 'SER', temperature: float = - 1.0, pressure:

float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set

as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

- **component** The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

Returns This MaterialToMaterialPhaseDiagramCalculation object

with_second_axis (axis: tc_python.material_to_material.MaterialToMaterialCalculationAxis)

Sets the second axis (either temperature of fraction of material B). This calculation type requires that both temperature and fraction of material B axis are set.

Parameters axis – The axis

Returns This MaterialToMaterialPhaseDiagramCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)
 Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a * .tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This MaterialToMaterialPhaseDiagramCalculation object

class tc_python.material_to_material.MaterialToMaterialPhaseDiagramResult(result)
 Bases: tc_python.step_or_map_diagrams.PhaseDiagramResult

Result of a Material to Material phase diagram calculation, it can be evaluated using quantities or Console Mode syntax.

add_coordinate_for_phase_label (x: float, y: float)

Sets a coordinate in the result plot for which the stable phases will be evaluated and provided in the result data object. This can be used to plot the phases of a region into the phase diagram or just to programmatically evaluate the phases in certain regions.

Warning: This method takes coordinates of the **plot** axes and not of the calculation axis.

Parameters

- **x** The coordinate of the first **plot** axis ("x-axis") [unit of the **plot** axis]
- y The coordinate of the second plot axis ("y-axis") [unit of the plot axis]

Returns This MaterialToMaterialPhaseDiagramResult object

 $\texttt{get_values_grouped_by_quantity_of} (x_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str], y_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str]) \rightarrow tc_python.step_or_map_diagrams.PhaseDiagramResultValues$

Returns x-y-line data grouped by the multiple datasets of the specified quantities (for example in dependency of components). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity. Usually the result data represents the phase diagram.

Note: The different datasets will contain *NaN*-values between different subsections and are not sorted (because they are unsortable due to their nature).

Note: Its possible to use functions as axis variables, either by using ThermodynamicQuantity. user_defined_function, or by using an expression that contains '='.

Example get_values_grouped_by_quantity_of('T', ThermodynamicQuantity.user_defined_function('HM.T'))

Example get_values_grouped_by_quantity_of('T', 'CP=HM.T')

Parameters

- **x_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example '*T*'), *MATERIAL_B_FRACTION*, or even a function (for example '*f*=*T**1.01')
- **y_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), *MATERIAL_B_FRACTION*, or even a function (for example '*CP=HM.T*')

Returns The phase diagram data

 $\texttt{get_values_grouped_by_stable_phases_of} (x_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity str], y_quantity: \\ Union[tc_python.quantity_factory.ThermodynamicQuantity, str]) \rightarrow tc_python.step_or_map_diagrams_PhaseDiagramResultValues str]$

 $str]) \rightarrow tc_python.step_or_map_diagrams.PhaseDiagramResultValues$ Returns x-y-line data grouped by the sets of "stable phases" (for example "LIQUID" or "LIQUID + FCC_A1"). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity. Usually the result data represents the phase diagram.

Note: The different datasets will contain *NaN*-values between different subsections and are not sorted (because they are unsortable due to their nature).

Note: Its possible to use functions as axis variables, either by using ThermodynamicQuantity. user_defined_function, or by using an expression that contains '='.

Example get_values_grouped_by_quantity_of('T', ThermodynamicQuantity.user_defined_function('HM.T')) **Example** get_values_grouped_by_quantity_of('T', 'CP=HM.T')

Parameters

- **x_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example '*T*'), *MATERIAL_B_FRACTION*, or even a function (for example '*f*=*T**1.01')
- **y_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), *MATERIAL_B_FRACTION*, or even a function (for example '*CP=HM.T*')

Returns The phase diagram data

remove_phase_labels()

Erases all added coordinates for phase labels.

Returns This MaterialToMaterialPhaseDiagramResult object

save_to_disk (path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load_result_from_disk()

Parameters path – the path to the folder you want the result to be saved in. It can be relative or absolute.

Returns this MaterialToMaterialPhaseDiagramResult object

set_phase_name_style (phase_name_style_enum: tc_python.step_or_map_diagrams.PhaseNameStyle

= <*PhaseNameStyle.NONE:* 0>)

Sets the style of the phase name labels that will be used in the result data object (constitution description, ordering description, \dots).

Default: PhaseNameStyle.NONE

Parameters phase_name_style_enum – The phase name style

Returns This MaterialToMaterialPhaseDiagramResult object

class tc_python.material_to_material.**MaterialToMaterialPropertyDiagramCalculation**(*calculator*) Bases: tc_python.step_or_map_diagrams.AbstractPropertyDiagramCalculation

Configuration for a Material to Material property diagram calculation.

Note: Specify the conditions and possibly other settings, the calculation is performed with *calculate()*.

calculate (keep_previous_results: bool = False, timeout_in_minutes: float = 0.0) \rightarrow tc_python.material_to_material.MaterialToMaterialPropertyDiagramResult Performs the Material to Material property diagram calculation.

Warning: If you use *keep_previous_results=True*, you must not use another calculator or even get results in between the calculations using *calculate()*. Then the previous results will actually be lost.

Parameters

- **keep_previous_results** If *True*, results from any previous call to this method are appended. This can be used to combine calculations with multiple start points if the stepping fails at a certain condition.
- **timeout_in_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.
- **Returns** A new *MaterialToMaterialPropertyDiagramResult* object which later can be used to get specific values from the calculated result

disable_global_minimization()

Disables global minimization.

Default: Enabled

Returns This MaterialToMaterialPropertyDiagramCalculation object

disable_step_separate_phases()

Disables *step separate phases*. This is the **default** setting.

Returns This MaterialToMaterialPropertyDiagramCalculation object

enable_global_minimization()

Enables global minimization.

Default: Enabled

Returns This MaterialToMaterialPropertyDiagramCalculation object

enable_step_separate_phases()

Enables step separate phases.

Default: By default separate phase stepping is *disabled*

Note: This is an advanced option, it is used mostly to calculate how the Gibbs energy for a number of phases varies for different compositions. This is particularly useful to calculate Gibbs energies for complex phases with miscibility gaps and for an ordered phase that is never disordered (e.g. SIGMA-phase, G-phase, MU-phase, etc.).

Returns This MaterialToMaterialPropertyDiagramCalculation object

get_components() \rightarrow List[str]

Returns the names of the components in the system (including all components auto-selected by the database(s)).

Returns The component names

$\texttt{get_gibbs_energy_addition_for}\ (\textit{phase: str}) \ \rightarrow \textit{float}$

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters phase – Specify the name of the (stoichiometric or solution) phase with the addition

Returns Gibbs energy addition to G per mole formula unit.

$get_system_data() \rightarrow tc_python.abstract_base.SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

Parameters command - The Thermo-Calc Console Mode command

Returns This MaterialToMaterialPropertyDiagramCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

set_activities: Dict[str, float])

Sets the constant activity conditions.

Note: The activity conditions are identical for both materials.

Parameters activities – The constant activities

Returns This MaterialToMaterialPropertyDiagramCalculation object

set_composition_unit (unit: tc_python.utils.CompositionUnit = <CompositionUnit.MASS_PERCENT: 0>) Sets the composition unit of both materials A and B.

Default: Weight percent

Parameters unit - The composition unit of both materials A and B

Returns This MaterialToMaterialPropertyDiagramCalculation object

set_gibbs_energy_addition_for (phase: str, gibbs_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters

• phase - Specify the name of the (stoichiometric or solution) phase with the addition

• gibbs_energy – Addition to G per mole formula unit

Returns This MaterialToMaterialPropertyDiagramCalculation object

set_material_a (composition: Dict[str, float], dependent_component: str = None)
Sets the composition of the material A.

The unit is set with set_composition_unit().

Tip: The material can also have constant activity conditions, they are set in set_activities().

Parameters

- composition The composition of the material A
- dependent_component The dependent component of the material A

Returns This MaterialToMaterialPropertyDiagramCalculation object

set_material_b (composition: Dict[str, float], dependent_component: str = None)
Sets the composition of the material B.

The unit is set with set_composition_unit().

Tip: The material can also have constant activity conditions, they are set in set_activities ().

Parameters

- composition The composition of the material B
- dependent_component The dependent component of the material B

Returns This MaterialToMaterialPropertyDiagramCalculation object

set_phase_to_dormant (phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters phase – The phase name or ALL_PHASES for all phases

Returns This MaterialToMaterialPropertyDiagramCalculation object

set_phase_to_entered (phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

Parameters

- **phase** The phase name or ALL_PHASES for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

Returns This MaterialToMaterialPropertyDiagramCalculation object

set_phase_to_fixed (phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

Parameters

- phase The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

Returns This MaterialToMaterialPropertyDiagramCalculation object

set_phase_to_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This MaterialToMaterialPropertyDiagramCalculation object

set_pressure (pressure: float)

Sets the pressure (i.e. the condition *P*).

Note: If the flag *default_conditions=True* has been set during the creation of the calculator, the pressure is set to 1000 hPa by default.

Parameters pressure – The pressure [Pa]

Returns This MaterialToMaterialPropertyDiagramCalculation object

set_system_size (system_size: float)

Sets the system size (i.e. the condition 'N', the number of moles).

Note: If the flag *default_conditions=True* has been set during the creation of the calculator, the system size is set to 1.0 moles by default.

Parameters system_size – The system size [mole]

Returns This MaterialToMaterialPropertyDiagramCalculation object

with_axis (axis: tc_python.material_to_material.MaterialToMaterialCalculationAxis)

Sets the axis (either temperature of fraction of material B). This calculation type requires that either temperature or fraction of material B is set as a constant condition - the other one is set as an axis.

Parameters axis – The axis

Returns This MaterialToMaterialPropertyDiagramCalculation object

with_constant_condition (condition: tc_python.material_to_material.ConstantCondition)

Sets the constant condition (either temperature of fraction of material B). This calculation type requires that either temperature or fraction of material B is set as a constant condition - the other one is set as an axis.

Parameters condition - The condition

Returns This MaterialToMaterialPropertyDiagramCalculation object

with_options (options: tc_python.step_or_map_diagrams.PropertyDiagramOptions)
 Sets the simulation options.

Parameters options – The simulation options

Returns This MaterialToMaterialPropertyDiagramCalculation object

with_reference_state (component: str, phase: str = 'SER', temperature: float = - 1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

- **component** The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

Returns This MaterialToMaterialPropertyDiagramCalculation object

with_system_modifications (*system_modifications:* tc_python.abstract_base.SystemModifications) Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a * .tdb-file.

Parameters system_modifications - The system modification to be performed

Returns This MaterialToMaterialPropertyDiagramCalculation object

class tc_python.material_to_material.MaterialToMaterialPropertyDiagramResult(result)
 Bases: tc_python.step_or_map_diagrams.PropertyDiagramResult

Result of a Material to Material property diagram. It can be used to query for specific values.

get_values_grouped_by_quantity_of (x_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity,

str], *y_quantity:* Union[tc_python.quantity_factory.ThermodynamicQuantity, *str*], *sort_and_merge:* bool = True) \rightarrow Dict[str,

tc_python.utils.ResultValueGroup]

Returns x-y-line data grouped by the multiple datasets of the specified quantities (typically the phases). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity.

Note: The different datasets might contain *NaN*-values between different subsections and might not be sorted **even if the flag `sort_and_merge` has been set** (because they might be unsortable due to their nature).

Note: Its possible to use functions as axis variables, either by using ThermodynamicQuantity. user_defined_function, or by using an expression that contains '='.

Example get_values_grouped_by_quantity_of('T', tity.user_defined_function('HM.T'))

ThermodynamicQuan-

Example get_values_grouped_by_quantity_of('T', 'CP=HM.T')

Parameters

- **x_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example '*T*'), *MATERIAL_B_FRACTION*, or even a function (for example '*f*=*T**1.01')
- **y_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), *MATERIAL_B_FRACTION*, or even a function (for example '*CP=HM.T*')
- sort_and_merge If *True*, the data is sorted and merged into as few subsections as possible (divided by *NaN*)

Returns Containing the datasets with the quantities as their keys

Note: The different datasets might contain *NaN*-values between different subsections and different lines of an ambiguous dataset. They might not be sorted **even if the flag `sort_and_merge` has been set** (because they might be unsortable due to their nature).

Note: Its possible to use functions as axis variables, either by using ThermodynamicQuantity. user_defined_function, or by using an expression that contains '='.

Exampleget_values_grouped_by_quantity_of('T',ThermodynamicQuantity_user_defined_function('HM.T'))

Example get_values_grouped_by_quantity_of('T', 'CP=HM.T')

Parameters

ThermodynamicQuantity.

- **x_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example '*T*'), *MATERIAL_B_FRACTION*, or even a function (for example '*f*=*T**1.01')
- **y_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), *MATERIAL_B_FRACTION*, or even a function (for example '*CP=HM.T*')

 sort_and_merge – If True, the data will be sorted and merged into as few subsections as possible (divided by NaN)

Returns Containing the datasets with the quantities as their keys

get_values_of (x_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str], y_quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str]) \rightarrow [typing.List[float], typing.List[float]]

Returns sorted x-y-line data without any separation. Use get_values_grouped_by_quantity_of() or get_values_grouped_by_stable_phases_of() instead if you need such a separation. The available quantities can be found in the documentation of the factory class ThermodynamicQuantity.

Note: This method will always return sorted data without any *NaN*-values. If it is unsortable that might give data that is hard to interpret. In such a case you need to choose the quantity in another way or use one of the other methods. One example of this is to use quantities with *All*-markers, for example *MassFractionOfAComponent("All")*.

Note: Its possible to use functions as axis variables, either by using ThermodynamicQuantity. user_defined_function(), or by using an expression that contains '='.

Example get_values_grouped_by_quantity_of('T', ThermodynamicQuantity.user_defined_function('HM.T'))

Example get_values_grouped_by_quantity_of('T', 'CP=HM.T')

Parameters

- **x_quantity** The first thermodynamic quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example 'T', MATERIAL_B_FRACTION, or even a function (for example 'f=T*1.01').
- **y_quantity** The second thermodynamic quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example 'NV'), *MATERIAL_B_FRACTION*, or even a function (for example '*CP=HM.T*')

Returns A tuple containing the x- and y-data in lists

save_to_disk (path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load_result_from_disk()

Parameters path – the path to the folder you want the result to be saved in. It can be relative or absolute.

Returns this MaterialToMaterialPropertyDiagramResult object

set_phase_name_style (phase_name_style_enum: tc_python.step_or_map_diagrams.PhaseNameStyle

= <*PhaseNameStyle.NONE:* 0>)

Sets the style of the phase name labels that will be used in the result data object (constitution description, ordering description, \ldots).

Default: PhaseNameStyle.NONE

Parameters phase_name_style_enum – The phase name style

Returns This MaterialToMaterialPropertyDiagramResult object

class tc_python.material_to_material.MaterialToMaterialSingleEquilibriumCalculation(calculato
 Bases: tc_python.single_equilibrium.AbstractSingleEquilibriumCalculation

Configuration for a Material to Material single fraction of B calculation.

Note: Specify the conditions and possibly other settings, the calculation is performed with *calculate()*.

calculate (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.material_to_material.MaterialToMaterialSingleEquilibriumResult* Performs the material calculation.

Note: The calculation result is **no** temporary result object.

- **Parameters timeout_in_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout_in_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.
- **Returns** A new *MaterialToMaterialSingleEquilibriumResult* object which can be used to get specific values from the calculated result. It is undefined behavior to use that object after the state of the calculation has been changed.

disable_global_minimization()

Turns the global minimization completely off.

Returns This MaterialToMaterialSingleEquilibriumCalculation object

enable_global_minimization()

Turns the global minimization on (using the default settings).

Returns This MaterialToMaterialSingleEquilibriumCalculation object

get_components() \rightarrow List[str]

Returns a list of components in the system (including all components auto-selected by the database(s)).

Returns The components

$get_gibbs_energy_addition_for(phase: str) \rightarrow float$

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters phase – Specify the name of the (stoichiometric or solution) phase with the addition

Returns Gibbs energy addition to G per mole formula unit.

get_system_data() \rightarrow tc_python.abstract_base.SystemData

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

```
run_poly_command(command: str)
```

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

Parameters command - The Thermo-Calc Console Mode command

Returns This MaterialToMaterialSingleEquilibriumCalculation object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

set_activities (activities: Dict[str, float])

Sets the constant activity conditions.

Note: The activity conditions are identical for both materials.

Parameters activities – The constant activities

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_component_to_entered(component: str)

Sets the specified component to the status ENTERED, that is the default state.

Parameters component – The component name or ALL_COMPONENTS

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_component_to_suspended (component: str, reset_conditions: bool = False)
Sets the specified component to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters

- **reset_conditions** if 'True' also remove composition conditions for the component if they are defined
- component The component name or ALL_COMPONENTS

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_composition_unit (unit: tc_python.utils.CompositionUnit = <CompositionUnit.MASS_PERCENT: 0>)
Sets the server side A and B

Sets the composition unit of **both materials A and B**.

Default: Weight percent

Parameters unit – The composition unit of both materials A and B

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_gibbs_energy_addition_for (phase: str, gibbs_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs_energy Addition to G per mole formula unit

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_material_a (composition: Dict[str, float], dependent_component: str = None)
Sets the composition of the material A.

The unit is set with set_composition_unit().

Tip: The material can also have constant activity conditions, they are set in set_activities().

Parameters

- composition The composition of the material A
- dependent_component The dependent component of the material A

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_material_b (composition: Dict[str, float], dependent_component: str = None)
Sets the composition of the material B.

The unit is set with set_composition_unit().

Tip: The material can also have constant activity conditions, they are set in *set_activities()*.

Parameters

- composition The composition of the material B
- dependent_component The dependent component of the material B

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_phase_to_dormant (phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_phase_to_entered (*phase: str, amount: float = 1.0*) Sets the phase to the status ENTERED, that is the default state.

Parameters

- **phase** The phase name or ALL_PHASES for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_phase_to_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

Parameters

- phase The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_phase_to_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

Parameters phase – The phase name or *ALL_PHASES* for all phases

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_pressure (pressure: float)

Sets the pressure (i.e. the condition *P*).

Note: If the flag *default_conditions=True* has been set during the creation of the calculator, the pressure is set to 1000 hPa by default.

Parameters pressure – The pressure [Pa]

Returns This MaterialToMaterialSingleEquilibriumCalculation object

set_system_size (system_size: float)

Sets the system size (i.e. the condition 'N', the number of moles).

Note: If the flag *default_conditions=True* has been set during the creation of the calculator, the system size is set to 1.0 moles by default.

Parameters system_size – The system size [mole]

Returns This MaterialToMaterialSingleEquilibriumCalculation object

with_first_constant_condition (condition: tc_python.material_to_material.ConstantCondition)
 Sets the first constant condition (either temperature of fraction of material B).

Parameters condition – The condition

Returns This MaterialToMaterialSingleEquilibriumCalculation object

with_options (options: tc_python.single_equilibrium.SingleEquilibriumOptions)
 Sets the simulation options.

Parameters options – The simulation options

Returns This SingleEquilibriumCalculation object

with_reference_state (component: str, phase: str = 'SER', temperature: float = - 1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

Parameters

- component The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

Returns This MaterialToMaterialSingleEquilibriumCalculation object

with_second_constant_condition (condition: tc_python.material_to_material.ConstantCondition)
 Sets the second constant condition (either temperature of fraction of material B).

Parameters condition - The condition

Returns This MaterialToMaterialSingleEquilibriumCalculation object

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications) Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. user) databases loaded as a *.tdb-file.

Parameters system_modifications – The system modification to be performed

Returns This MaterialToMaterialSingleEquilibriumCalculation object

class tc_python.material_to_material.MaterialToMaterialSingleEquilibriumResult(result)
 Bases: tc_python.single_equilibrium.SingleEquilibriumResult

Result of a Material To Material calculation for a single fraction of material B, it can be evaluated using a quantity or Console Mode syntax.

change_pressure (pressure: float)

Change the pressure and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with higher performance. The properties are calculated at the new pressure using the phase amount, temperature and composition of phases from the initial equilibrium. Use $get_value_of()$ to obtain them.

Parameters pressure – The pressure [Pa]

Returns This MaterialToMaterialSingleEquilibriumResult object

change_temperature (temperature: float)

Change the temperature and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with high performance. The properties are calculated at the new temperature using the phase amount, pressure and composition of phases from the initial equilibrium. Use $get_value_of()$ to obtain them.

Note: This is typically used when calculating room temperature properties (e.g. density) for a material when it is assumed that the equilibrium phase amount and composition freeze-in at a higher temperature during cooling.

Parameters temperature – The temperature [K]

Returns This MaterialToMaterialSingleEquilibriumResult object

$\texttt{get_components()} \rightarrow List[str]$

Returns the names of the components selected in the system (including any components auto-selected by the database(s)).

Returns The names of the selected components

```
\texttt{get\_conditions()} \rightarrow List[str]
```

Returns the conditions.

Returns The selected conditions

get_phases() \rightarrow List[str]

Returns the phases present in the system due to its configuration. It also contains all phases that have been automatically added during the calculation, this is the difference to the method System. get_phases_in_system().

Returns The names of the phases in the system including automatically added phases

get_stable_phases() \rightarrow List[str]

Returns the stable phases (i.e. the phases present in the current equilibrium).

Returns The names of the stable phases

- get_value_of (quantity: Union[tc_python.quantity_factory.ThermodynamicQuantity, str]) \rightarrow float Returns a value from a single equilibrium calculation.
 - **Parameters quantity** The thermodynamic quantity to get the value of; a Console Mode syntax strings can be used as an alternative (for example "NPM(FCC_A1)")

Returns The requested value

run_poly_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine. This affects only the state of the result object.

Parameters command - The Thermo-Calc Console Mode command

Returns This MaterialToMaterialSingleEquilibriumResult object

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

```
save_to_disk (path: str)
```

Saves the result to disk. Note that the result is a folder, containing potentially many files. The result can later be loaded with load_result_from_disk()

Parameters path – the path to the folder you want the result to be saved in. It can be relative or absolute.

Returns this MaterialToMaterialSingleEquilibriumResult object

A constant temperature condition.

5.1.9 Package "process_metallurgy"

5.1.9.1 Module "base"

```
class tc_python.process_metallurgy.base.AbstractAddition
```

Bases: object

The base class for representing an addition to an equilibrium calculation or process simulation.

```
get_composition() \rightarrow Dict[str, float]
```

Returns the composition of the addition - without containing a dependent component.

Returns The composition [in the unit provided by getCompositionUnit()]

```
abstract get_composition_unit()
```

Returns the composition unit used in this addition.

Returns The composition unit

```
\texttt{get\_dependent\_component()} \rightarrow str
```

Returns the dependent component.

Returns The dependent component or an empty string if no dependent component is defined

```
get_elements () \rightarrow Set[str]
Returns all elements of the addition.
```

Returns The elements

$\texttt{get_id}() \rightarrow str$

Returns the unique ID of the addition.

Returns The unique ID of the addition

```
get_temperature() \rightarrow float
```

Returns the temperature of the addition. This refers to the temperature before it is added to the process.

Returns The temperature [K]

```
\texttt{is\_do\_scale()} \rightarrow bool
```

Returns if the composition of the addition is being scaled to 100% / 1 or not.

Returns If the composition is scaled

```
is\_empty() \rightarrow bool
```

Returns if the addition is "empty", i.e., has zero amount.

Returns If the addition is empty

```
class tc_python.process_metallurgy.base.ActivityReference(value)
```

Bases: enum.Enum

The reference for a slag activity calculation. The actual reference phase depends on the component for which the activity is request and can be obtained by using these methods on the result object: tc_python.process_metallurgy.process.ProcessSimulationResult. get_formula_for_activity_of_slag() or tc_python.process_metallurgy. equilibrium.EquilibriumResult.get_formula_for_activity_of_slag().

LIQUID = 1

The reference is liquid slag.

SOLID = 0

The reference is solid slag.

```
class tc_python.process_metallurgy.base.PhaseGroup(value)
    Bases: enum.Enum
```

The phase group, such a group is collecting all phases that belong to a certain type.

```
ALL_METAL = 2
```

All metal phases.

 $ALL_SLAG = 5$

All slag phases.

GAS = 6 All gas phases.

LIQUID_METAL = 0 All liquid metal phases.

LIQUID SLAG = 4

All liquid slag phases.

SOLID_METAL = 1 All solid metal phases.

 $SOLID_SLAG = 3$

All solid slag phases.

class tc_python.process_metallurgy.base.ProcessDatabase(value)
 Bases: enum.Enum

The database used for a Process Metallurgy calculation.

```
LATEST = 0
```

The latest database available.

OXDEMO = 1

The database OXDEMO.

TCOX10 = 4

The database TCOX10.

TCOX11 = 5

The database TCOX11.

TCOX8 = 2

The database TCOX8.

TCOX9 = 3

The database TCOX9.

class tc_python.process_metallurgy.base.ProcessMetallurgyOptions Bases: tc_python.single_equilibrium.SingleEquilibriumOptions

The options for a process metallurgy calculation.

disable_approximate_driving_force_for_metastable_phases()

Disables the approximation of the driving force for metastable phases.

Default: Enabled

Note: When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct. use disable_approximate_driving_force_for_metastable_phases() to force the calculation to converge for the metastable phases.

Returns This *ProcessMetallurgyOptions* object

disable_control_step_size_during_minimization()

Disables step size control during minimization (non-global).

Default: Enabled

Returns This *ProcessMetallurgyOptions* object

disable_force_positive_definite_phase_hessian()

Disables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

Returns This *ProcessMetallurgyOptions* object

enable_approximate_driving_force_for_metastable_phases()

Enables the approximation of the driving force for metastable phases.

Default: Enabled

Note: When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it important driving forces is that these are correct. use disable_approximate_driving_force_for_metastable_phases() force to the calculation to converge for the metastable phases.

Returns This *ProcessMetallurgyOptions* object

enable_control_step_size_during_minimization()

Enables step size control during normal minimization (non-global).

Default: Enabled

Returns This *ProcessMetallurgyOptions* object

enable_force_positive_definite_phase_hessian()

Enables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

Returns This *ProcessMetallurgyOptions* object

set_global_minimization_max_grid_points (max_grid_points: int = 2000)

Sets the maximum number of grid points in global minimization. Only applicable if global minimization is actually used.

Default: 2000 points

Parameters max_grid_points – The maximum number of grid points

Returns This ProcessMetallurgyOptions object

set_max_no_of_iterations (max_no_of_iterations: int = 2000)
Sets the maximum number of iterations for the CALPHAD minimizer.

Default: max. 2000 iterations

Note: As some models give computation times of more than 1 CPU second/iteration, this number is also used to check the CPU time and the calculation stops if 500 CPU seconds/iterations are used.

Parameters max_no_of_iterations – The max. number of iterations

Returns This ProcessMetallurgyOptions object

set_process_minimization_policy (minimization_policy: tc_python.process_metallurgy.base.ProcessMinimizationPolicy Sets the minimization policy for the process metallurgy calculation. It is possible to choose policies that try different methods if one method fails.

Parameters minimization_policy - The minimization policy to be used

Returns This *ProcessMetallurgyOptions* object

set_required_accuracy (accuracy: float = 1e-06)
Set the required relative second or accuracy.

Sets the required relative accuracy.

Default: 1.0E-6

Note: This is a relative accuracy, and the program requires that the relative difference in each variable must be lower than this value before it has converged. A larger value normally means fewer iterations

but less accurate solutions. The value should be at least one order of magnitude larger than the machine precision.

Parameters accuracy – The required relative accuracy

Returns This *ProcessMetallurgyOptions* object

```
set_smallest_fraction (smallest_fraction: float = 1e-16)
```

Sets the smallest fraction for constituents that are unstable.

It is normally only in the gas phase that you can find such low fractions.

The **default value** for the smallest site-fractions is 1E-16 for all phases except for IDEAL phase with one sublattice site (such as the GAS mixture phase in many databases) for which the default value is always as 1E-30.

Parameters smallest_fraction – The smallest fraction for constituents that are unstable

Returns This ProcessMetallurgyOptions object

```
class tc_python.process_metallurgy.base.ProcessMinimizationPolicy(value)
    Bases: enum.Enum
```

The policy for the CALPHAD minimization routine used in a calculation.

Note: This affects the runtime and stability of a calculation. Global minimization is more stable but more timeconsuming. Local minimization is much faster but can miss new phases coming up. Global test is a compromise between both approaches.

```
GLOBAL = 0
```

Always use global minimization.

```
GLOBAL\_TEST = 1
```

Always use global test.

```
GLOBAL\_TEST\_GLOBAL = 5
```

First try global test, if that fails use global minimization.

```
LOCAL = 2
```

Always use local minimization.

 $LOCAL_GLOBAL_TEST = 3$

First try local minimization, if that fails use global test.

```
LOCAL__GLOBAL_TEST__GLOBAL = 4
```

First try local minimization, if that fails try global test and if that fails use global minimization.

```
class tc_python.process_metallurgy.base.SlagProperty(value)
```

Bases: enum.Enum

The slag property, different definitions are available. The actual definition of a certain slag property for the current system can be obtained using these methods on the result object: tc_python.process_metallurgy.process.ProcessSimulationResult.getFormulaForSlagProperty() or tc_python.process_metallurgy.equilibrium. EquilibriumResult.getFormulaForSlagProperty().

Note: If not all components required by the definition of slag property are available in a given system, the slag property will return *NaN*.

```
B2 = 0
          Basicity B2 (based on: CaO / SiO2).
     B3 = 1
          Basicity B3 (based on: CaO / MgO / SiO2).
     B4 = 3
          Basicity B4 (based on: CaO / MgO / SiO2 / Al2O3).
     BAS2 = 2
          Basicity Bas2 (based on: CaO / MgO / SiO2 / Al2O3 / TiO2).
     BELLS_RATIO = 4
          Basicity Bell's ratio (based on: CaO / MgO / SiO2 / Al2O3).
     LOG 10 SULPHUR CAPACITY = 5
          Logarithmic sulphur capacity of the slag.
     LS = 6
          Sulphur capacity Ls of the slag.
class tc_python.process_metallurgy.base.SlagType(value)
     Bases: enum. Enum
     The type of slag considered for a slag property calculation.
```

ALL = 2

The slag property is calculated for all slag, i.e. for both the liquid and solid slag.

LIQUID = 1

The slag property is calculated for all liquid slag.

SOLID = 0

The slag property is calculated for all solid slag.

5.1.9.2 Module "equilibrium"

class tc_python.process_metallurgy.equilibrium.AbstractEquilibriumAddition
 Bases: tc_python.process_metallurgy.base.AbstractAddition

The base class for representing an addition to an equilibrium calculation.

```
set_amount (amount: float)
```

Change the amount of the addition.

Parameters amount – The new amount [in the amount unit of this addition]

Returns This AbstractEquilibriumAddition object

set_component_composition (*component: str, content: float*) Change the composition of a component of the addition.

Parameters

- component The component to be changed
- content The new content of the component [in the composition unit defined for this addition]

Returns This AbstractEquilibriumAddition object

class tc_python.process_metallurgy.equilibrium.**AdiabaticEquilibriumCalculation** (*calculator*) Bases: tc_python.process_metallurgy.equilibrium.EquilibriumCalculation

An *adiabatic* Process Metallurgy equilibrium calculation. Such calculations can for example be used to determine the global equilibrium state of a process.

add_addition (*addition:* tc_python.process_metallurgy.equilibrium.AbstractEquilibriumAddition) Adds an addition to the calculation.

Parameters addition - A EquilibriumAddition or EquilibriumGasAddition

Returns This AdiabaticEquilibriumCalculation object

add_poly_command(command: str)

Adds a Thermo-Calc Console syntax POLY module command which will be executed when performing the calculate () method.

If multiple commands are added, they will be executed in the order of addition. Each command will only be executed one.

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

Parameters command – The POLY module command in Thermo-Calc console syntax

Returns This AdiabaticEquilibriumCalculation object

- **calculate** (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.process_metallurgy.equilibrium.EquilibriumResult* Runs the Process Metallurgy equilibrium calculation.
 - Parameters timeout_in_minutes The calculation will be aborted after that time, default: no timeout

Returns A new EquilibriumResult object

remove_addition (*addition*: tc_python.process_metallurgy.equilibrium.AbstractEquilibriumAddition) Removes an addition from the calculation.

Parameters addition – The addition to be removed

Returns This AdiabaticEquilibriumCalculation object

remove_all_additions()

Removes all additions from the calculation.

Returns This AdiabaticEquilibriumCalculation object

set_pressure (pressure: float = 100000.0)

Sets the pressure.

Parameters pressure – The pressure [Pa]

Returns This AdiabaticEquilibriumCalculation object

with_options (*options*: tc_python.process_metallurgy.base.ProcessMetallurgyOptions) Sets the options for the calculation.

Parameters options - The options

Returns This AdiabaticEquilibriumCalculation object

class tc python.process metallurgy.equilibrium.EquilibriumAddition (composition: Dict[str, float], amount: float, temperature: float 293.15, composition_unit: tc_python.utils.CompositionUnit = <*Com*positio*nUnit.MASS_PERCENT: 0>*, do_scale: bool = False) Bases: tc_python.process_metallurgy.equilibrium.AbstractEquilibriumAddition

An addition to an equilibrium calculation.

Tip: By setting $do_scale=True$, the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not summing to 100% / 1. An example could be a slag addition which is provided like this: 90 wt-% CaO - 5 wt-% Al2O3 - 4 wt-% SiO2.

Parameters

- composition The composition
- **amount** The amount [kg]
- temperature The initial addition temperature (default: 20 °C) [K]
- **composition_unit** The composition unit
- do_scale If the composition is scaled to 100% / fraction of 1

 $\texttt{get_amount()} \rightarrow \texttt{float}$

Returns the amount of this addition.

Returns The amount [kg]

get_composition_unit () \rightarrow *tc_python.utils.CompositionUnit* Returns the composition unit used in this addition.

Returns The composition unit

class tc_python.process_metallurgy.equilibrium.EquilibriumCalculation(metallurgical_reaction)
 Bases: object

A Process Metallurgy equilibrium calculation. Such calculations can for example be used to determine the global equilibrium state of a process.

abstract add_addition (*addition*: tc_python.process_metallurgy.equilibrium.AbstractEquilibriumAddition) Adds an addition to the calculation.

Parameters addition – The addition

Returns This EquilibriumCalculation object

abstract add_poly_command(command: str)

Adds a Thermo-Calc Console syntax POLY module command which will be executed when performing the calculate () method.

If multiple commands are added, they will be executed in the order of addition. Each command will only be executed one.

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

Parameters command - The POLY module command in Thermo-Calc console syntax

Returns This EquilibriumCalculation object

abstract calculate (timeout_in_minutes: float = 0.0) $\rightarrow tc_python.process_metallurgy.equilibrium.EquilibriumResult$ Runs the Process Metallurgy equilibrium calculation.

Parameters timeout_in_minutes - The calculation will be aborted after that time, default: no timeout

Returns A new EquilibriumResult object

abstract remove_addition (*addition*: tc_python.process_metallurgy.equilibrium.AbstractEquilibriumAddition) Removes an addition from the calculation.

Parameters addition – The addition to be removed

Returns This EquilibriumCalculation object

abstract remove_all_additions()

Removes all additions from the calculation.

Returns This EquilibriumCalculation object

abstract set_pressure (*pressure: float* = 100000.0) Sets the pressure.

Parameters pressure – The pressure [Pa]

Returns This EquilibriumCalculation object

update_addition (addition: tc_python.process_metallurgy.equilibrium.AbstractEquilibriumAddition)

Replaces an already added addition with an updated one. This is usually used to change the composition or amount of an addition while iterating over them. Typically, this is done for stepping or mapping calculations.

Note: The calculation must already contain the addition object to be updated.

Parameters addition – The new addition containing updated values

 $Returns \ This \ \texttt{IsoThermalMetallurgyCalculation} \ object$

abstract with_options (*options*: tc_python.process_metallurgy.base.ProcessMetallurgyOptions) Sets the options for the calculation.

Parameters options – The options

Returns This EquilibriumCalculation object

class tc_python.process_metallurgy.equilibrium.EquilibriumGasAddition(composition:

Dict[str, float], amount: float, temperature: float 293.15, amount_unit: tc_python.utils.GasAmountUn = <GasAmountUnit.NORM_CUBIC_METER: *0>*, composition_unit: tc_python.utils.GasComposition = <Gas-Compositio*nUnit.VOLUME PERCENT:* 4>, do_scale: bool False) Bases: tc_python.process_metallurgy.equilibrium.AbstractEquilibriumAddition

A gas addition to an equilibrium calculation.

Tip: By setting $do_scale=True$, the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not summing to 100% / 1. An example could be a gas addition which is provided like this: 90 vol-% Ar - 10 vol-% O2.

 $\begin{array}{l} \texttt{get_amount} () \to \texttt{float} \\ \text{Returns the amount of this addition.} \end{array}$

Note: The amount unit can be obtained using get_amount_unit().

Returns The amount [in the amount unit]

```
get_amount_unit() \rightarrow tc_python.utils.GasAmountUnit
Returns the amount unit used in this addition.
```

Returns The amount unit

get_composition_unit() \rightarrow *tc_python.utils.GasCompositionUnit* Returns the composition unit used in this addition.

Returns The composition unit

class tc_python.process_metallurgy.equilibrium.EquilibriumResult(result)
 Bases: tc_python.abstract_base.AbstractResult

The result of a Process Metallurgy equilibrium calculation.

```
\begin{array}{l} \texttt{get\_activity\_of\_slag} (\textit{component: str, reference: tc\_python.process\_metallurgy.base.ActivityReference} \\ = <ActivityReference.LIQUID: 1>) \rightarrow \texttt{float} \\ \texttt{Returns the activity of a component in the slag.} \end{array}
```

Parameters

- **component** The component
- reference The reference for the activity, can be liquid or solid slag, default: liquid slag

Returns The activity of the component [-]

```
get_amount() \rightarrow float
```

Returns the total amount.

Returns The total amount [kg]

 $\texttt{get_amount_of_elements()} \rightarrow Dict[str, float]$

Returns the amount of each element.

Returns The amount of the elements [kg]

get_amount_of_phase_groups () \rightarrow Dict[*tc_python.process_metallurgy.base.PhaseGroup*, float] Returns the amount of each phase group (e.g., for example all liquid slag).

Returns The amount of the phase groups [kg]

get_amount_of_phases() \rightarrow Dict[str, float]

Returns the amount of each phase.

Returns The amount of the phases [kg]

get_components() ightarrow Set[str]

Returns all components defined for the elements present in this result.

Returns The components present in this result

Parameters composition_unit - The composition unit, default: mass percent

Returns The composition

Parameters

- **phase** The phase name
- composition_unit The composition unit, default: mass percent
- composition_type Defines if the composition is given by element (e.g., 75 wt-% Fe 25 wt-% Cr) or by component (e.g. 65 wt-% Al2O3 35 wt-% CaO). In case of a metallic phase, the composition is given by element even if *component* is selected. Default: by component.

Returns The composition

Returns the composition of a phase group (e.g., all liquid slag) in the result.

Parameters

- phase_group The phase group
- composition_unit The composition unit, default: mass percent
- composition_type Defines if the composition is given by element (e.g., 75 wt-% Fe 25 wt-% Cr) or by component (e.g. 65 wt-% Al2O3 35 wt-% CaO). In case of a metallic phase, the composition is given by element even if *component* is selected. Default: by component.

Returns The composition

get_elements() \rightarrow Set[str]

Returns all elements defined for the result.

Returns All elements present in this result

 $get_formula_for_activity_of_slag(component: str, reference: tc_python.process_metallurgy.base.ActivityReference = <ActivityReference.LIQUID: 1>) <math>\rightarrow$ str

Returns the Thermo-Calc Console syntax formula used for calculating the activity of a component in the slag (e.g. AC(AL2O3, IONIC_LIQ). The actual activity can be obtained using get_activity_of_slag().

Parameters

- **component** The component
- **reference** The reference for the activity, can be liquid or solid slag, **default: liquid slag**

Returns The formula for calculating the activity

get_formula_for_slag_property (slag_property: tc_python.process_metallurgy.base.SlagProperty, slag_type: tc_python.process_metallurgy.base.SlagType =

 $\langle SlagType.ALL: 2 \rangle) \rightarrow str$

Returns the Thermo-Calc Console syntax formula used for calculating a property of the slag (e.g.

B(CAO)/B(SIO2). The actual slag property can be obtained using get_slag_property().

Parameters

- slag_property The slag property
- **slag_type** The part of the slag for which the property will be calculated. Can be all slag, the liquid or the solid slag. **Default: all slag**

Returns The formula for calculating the slag property

get_fraction_of_phase_groups (unit:		tc_python.utils.PhaseUnit		=	
	<phaseunit.mass_f< th=""><th>RACTION:</th><th>1>)</th><th>\rightarrow</th></phaseunit.mass_f<>	RACTION:	1>)	\rightarrow	
	Dict[tc_python.proce.	ss_metallurgy.bas	se.PhaseGroup,	float]	
Returns the fraction of the phase grou	ips (e.g., all liquid slag	g) in the result.	<u>^</u>		

Parameters unit – The unit of the fraction, default: volume fraction

Returns The phase fractions

```
get_fraction_of_phases (unit: tc_python.utils.PhaseUnit = <PhaseUnit.MASS_FRACTION:
I>) \rightarrow Dict[str, float]
```

Returns the fraction of the stable phases in the result.

Parameters unit – The unit of the fraction, default: volume fraction

Returns The phase fractions

get_gas_components() \rightarrow Set[str]

Returns all components of the gas phase defined for the elements present in this result.

Returns The components of the gas phase present in this result

get_oxygen_partial_pressure() \rightarrow float

Returns the partial pressure of oxygen in the result.

Returns The partial pressure [Pa]

```
get_pressure() \rightarrow float
```

Returns the pressure in the result.

Returns The pressure [Pa]

```
get_slag_property (slag_property: tc_python.process_metallurgy.base.SlagProperty, slag_type:
```

 $tc_python.process_metallurgy.base.SlagType = <SlagType.ALL: 2>) <math>\rightarrow$ float Returns a property of the slag. These properties are mostly used to describe the property of a slag to pick up sulfur.

Parameters

- slag_property The slag property
- **slag_type** The part of the slag for which the property will be calculated. Can be all slag, the liquid or the solid slag. **Default: all slag**

Returns The slag property [unit depending on the property]

```
get_stable_phases() \rightarrow Set[str]
```

Returns the stable phases in the result.

Returns The stable phases

```
get_stable_phases_in_phase_group (phase_group: tc_python.process_metallurgy.base.PhaseGroup)
```

 \rightarrow Set[str]

Returns the stable phases of a phase group (e.g., all liquid slag) in the result.

Parameters phase_group – The phase group

Returns The stable phases

get_temperature() ightarrow float

Returns the temperature in the result.

Returns The temperature [K]

get_value_of (*classic_expression: str*) \rightarrow float Returns a value for a thermodynamic quantity.

Warning: It should normally not be required to use this method, use the appropriate method available in the API instead.

Parameters classic_expression – The thermodynamic quantity to get the value of in Thermo-Calc Console Mode syntax (for example "NPM(FCC_A1)")

Returns The requested value

get_viscosity_dynamic_of_phase (*phase*: *str*) \rightarrow float Returns the dynamic viscosity of a phase in the result.

Parameters phase – The phase name

Returns The dynamic viscosity [Pa*s]

 $\texttt{get_viscosity_kinematic_of_phase}(\textit{phase: str}) \rightarrow \texttt{float}$

Returns the kinematic viscosity of a phase in the result.

Parameters phase – The phase name

Returns The kinematic viscosity [m**2/s]

class tc_python.process_metallurgy.equilibrium.**IsoThermalEquilibriumCalculation**(*calculation*) Bases: tc_python.process_metallurgy.equilibrium.EquilibriumCalculation

An *isothermal* Process Metallurgy equilibrium calculation. Such calculations can for example be used to determine the global equilibrium state of a process.

add_addition (*addition:* tc_python.process_metallurgy.equilibrium.AbstractEquilibriumAddition) Adds an addition to the calculation.

Parameters addition - A EquilibriumAddition or EquilibriumGasAddition

Returns This IsoThermalEquilibriumCalculation object

add_poly_command(command: str)

Adds a Thermo-Calc Console syntax POLY module command which will be executed when performing the calculate () method.

If multiple commands are added, they will be executed in the order of addition. Each command will only be executed one.

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

Parameters command - The POLY module command in Thermo-Calc console syntax

Returns This IsoThermalEquilibriumCalculation object

calculate (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.process_metallurgy.equilibrium.EquilibriumResult* Runs the Process Metallurgy equilibrium calculation.

Parameters timeout_in_minutes - The calculation will be aborted after that time, default: no timeout

Returns A new EquilibriumResult object

remove_addition (*addition*: tc_python.process_metallurgy.equilibrium.AbstractEquilibriumAddition) Removes an addition from the calculation.

Parameters addition – The addition to be removed

Returns This IsoThermalEquilibriumCalculation object

remove_all_additions()

Removes all additions from the calculation.

Returns This IsoThermalEquilibriumCalculation object

set_pressure (*pressure: float = 100000.0*) Sets the pressure.

Parameters pressure – The pressure [Pa]

Returns This IsoThermalEquilibriumCalculation object

set_temperature(temperature: float)

Sets the temperature.

Parameters temperature – The temperature [K]

Returns This IsoThermalEquilibriumCalculation object

update_addition (*addition:* tc_python.process_metallurgy.equilibrium.AbstractEquilibriumAddition) Replaces an already added addition with an updated one.

Tip: This is usually used to change the composition or amount of an addition while iterating over multiple values. Typically, this is done for stepping or mapping calculations.

Note: The calculation must already contain the addition object to be updated.

Parameters addition – A previously added addition object with the updated values

Returns This IsoThermalEquilibriumCalculation object

with_options (*options:* tc_python.process_metallurgy.base.ProcessMetallurgyOptions) Sets the options for the calculation.

Parameters options – The options

Returns This IsoThermalEquilibriumCalculation object

5.1.9.3 Module "process"

class tc_python.process_metallurgy.process.AbstractContinuousAddition
 Bases: tc_python.process_metallurgy.process.AbstractProcessAddition

The base class representing an addition in a process simulation that is added continuously over a period of time.

class tc_python.process_metallurgy.process.AbstractProcessAddition
 Bases: tc_python.process_metallurgy.base.AbstractAddition

The base class for representing an addition in a process simulation.

class tc_python.process_metallurgy.process.AbstractSingleTimeAddition
 Bases: tc_python.process_metallurgy.process.AbstractProcessAddition

The base class representing an addition in a process simulation that is added at a distinct time point.

class tc_python.process_metallurgy.process.**BulkZone** (*density*:

density: float, phase_group_to_transfer: tc_python.process_metallurgy.base.PhaseGroup, name: str)

Bases: tc_python.process_metallurgy.process.Zone

A bulk zone in a process simulation, this is representing a large volume in the process, for example the steel melt or the top slag. A zone is a volume in a process that has identical temperature and composition. It has well-defined boundaries to other zones.

Tip: This is a generic class and seldom used directly. Use instead *MetalBulkZone* or *SlaqBulkZone*.

add_addition: tc_python.process_metallurgy.process.AbstractSingleTimeAddition, time: float = 0.0)

Adds a single-time addition at the specified time point to the zone. The addition will be dissolved immediately.

Parameters

- $\bullet \ \textbf{addition} A \ \textit{SingleTimeAddition} \ \textbf{or} \ \textit{SingleTimeGasAddition} \\$
- time The time point [s]

Returns This BulkZone object

add_continuous_addition (*addition:* tc_python.process_metallurgy.process.AbstractContinuousAddition, from time: float = 0.0, to time: float = nan)

Adds a constant addition continuously during the specified time period to the zone. All added material will be dissolved immediately.

Parameters

- $\bullet \ \textbf{addition} A \ \textit{ContinuousAddition} \ \textbf{or} \ \textit{ContinuousGasAddition} \\$
- **from_time** The start time point [s]
- to_time The end time point [s]

Returns This BulkZone object

add_power (*power*: *float*, *from_time*: *float* = 0.0, *to_time*: *float* = *nan*)

Adds a constant power during a specified time period to the zone (for example heating or cooling).

Parameters

• power – The power [MW]

- **from_time** The start time point [s]
- to_time The end time point [s]

Returns This *BulkZone* object

disable_degassing()

Disables degassing for this zone, i.e. all gas formed at any time step will be staying in this zone.

Returns This BulkZone object

enable_degassing()

Enables degassing for this zone, i.e. any gas formed at any time step will be removed after that time step. This gas will be transferred into the *exhaust gas zone*. **This is the default**.

Returns This BulkZone object

```
get_density() \rightarrow float
```

Returns the density of the zone

Returns The density [kg/m**3]

$get_elements() \rightarrow Set[str]$

Returns the elements present in the zone. The elements are determined by the additions.

Returns The elements

$\texttt{get_id()} \rightarrow str$

Returns the unique id of the zone. :return: The zone id

get_phase_group_to_transfer() \rightarrow tc_python.process_metallurgy.base.PhaseGroup

Returns the phase group that is transferred from the attached reaction zones back to this zone after each time step.

Returns The phase group

is_degassing_enabled () \rightarrow bool Returns if degassing is enabled in the zone.

Returns If degassing is enabled

class tc_python.process_metallurgy.process.ContinuousAddition (composition:

Dict[str, float], rate: float, temperature: float = 293.15, composition_unit: tc_python.utils.CompositionUnit = <CompositionUnit.MASS_PERCENT: 0>, do_scale: bool = False)

Bases: tc_python.process_metallurgy.process.AbstractContinuousAddition

An addition in a process simulation that is added continuously during a period of time.

It is assumed that the material added during that period is dissolved instantaneously.

Tip: By setting $do_scale=True$, the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not scaling to 100% / 1. An example could be a slag addition which is provided like this: 90 wt-% CaO - 5 wt-% Al2O3 - 4 wt-% SiO2.

Parameters

- composition The composition
- **rate** The rate of addition [kg/s]
- temperature The initial addition temperature (default: 20 °C) [K]
- composition_unit The composition unit
- do scale If the composition is scaled to 100% / fraction of 1

```
get_composition_unit() \rightarrow tc_python.utils.CompositionUnit
```

Returns the composition unit used in this addition.

Returns The composition unit

get rate() \rightarrow float

Returns the rate of addition.

Returns The addition rate [kg/s]

class tc_python.process_metallurgy.process.ContinuousGasAddition (composition:

Dict[str, float], rate: float, temperature: float 293.15, rate unit: tc python.utils.GasRateUnit <Gas-= RateUnit.NORM_CUBIC_METER_PER_SE 0>, composition_unit: tc_python.utils.GasCompositionUnit <Gas-= CompositionUnit.VOLUME_PERCENT: 4>, do_scale: bool = False) Bases: tc_python.process_metallurgy.process.AbstractContinuousAddition

A gas addition in a process simulation that is added continuously during a period of time.

It is assumed that the gas added during that period is dissolved instantaneously.

Tip: By setting *do scale=True*, the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not scaling to 100% / 1. An example could be a gas addition which is provided like this: 90 vol-% Ar - 10 vol-% O2.

```
get_composition_unit() \rightarrow tc_python.utils.GasCompositionUnit
     Returns the composition unit used in this addition.
```

Returns The composition unit

get_rate() \rightarrow float Returns the rate of addition. **Note:** The rate unit can be obtained using get_rate_unit().

Returns The addition rate [in the rate unit]

get_rate_unit () \rightarrow *tc_python.utils.GasRateUnit* Returns the rate unit used in this addition.

Returns The rate unit

class tc_python.process_metallurgy.process.ExhaustGasResult(result)

Bases: object

A result representing the exhaust gas zone, here all exhaust gas generated during the process is accumulated.

The data is returned for each time point of the process simulation. These time points can be obtained from this method: *ProcessSimulationResult.get_time_points()*.

```
get_amount() \rightarrow List[float]
```

Returns the amount of exhaust gas present at each time point.

This is the amount of gas accumulated since the beginning of the process.

Returns The accumulated amount of gas at each time point [kg]

```
\texttt{get\_amount\_of\_components()} \rightarrow Dict[str, List[float]]
```

Returns the amount of each exhaust gas component present at each time point.

This is the amount of gas accumulated since the beginning of the process. This is different from the **current composition at each time point** obtained using get_composition().

Returns The accumulated amount of each gas component at each time point [kg]

Returns the current composition of the exhaust gas zone at each time point. This is the composition at each time point. This is different from the accumulated amount obtained using get_amount_of_components().

Parameters

- **composition_type** The type of the composition, can be by gas component or by element, **default: by gas component**
- unit The composition unit, default: mass percent

Returns The current composition of the gas components at each time point

 $get_pressure() \rightarrow List[float]$

Returns the pressure of the exhaust gas zone at each time point.

Returns The pressure [Pa]

```
\texttt{get\_stable\_phases()} \rightarrow \texttt{Set[str]}
```

Returns the stable phases within the exhaust gas zone at each time point.

Returns The stable phases

get_temperature() \rightarrow List[float]

Returns the temperature of the exhaust gas at each time point.

Returns The temperature at each time point [K]

class tc_python.process_metallurgy.process.MassTransferCoefficients
 Bases: object

The mass transfer coefficients between a reaction zone and a bulk zone vs. time.

add (mass_transfer_coefficient: float, time: float = 0.0)

Adds the mass transfer coefficient valid beginning at a time point.

This value is valid until another value is defined for a later time point.

Parameters

- mass_transfer_coefficient The mass transfer coefficient [m/s]
- time The time-point where the mass transfer coefficient begins to be valid [s]

Returns This *MassTransferCoefficients* object

class tc_python.process_metallurgy.process.**MetalBulkZone**(*density: float*) **Bases:** tc_python.process_metallurgy.process.Zone

A metallic bulk zone in a process simulation.

This is representing a large volume in the process, for example the steel melt. A zone is a volume in a process that has identical temperature and composition. It has well-defined boundaries to other zones.

The name of this zone is automatically defined and unique.

```
add_addition (addition: tc_python.process_metallurgy.process.AbstractSingleTimeAddition, time:
```

float = 0.0)

Adds a single-time addition at the specified time point to the zone. The addition will be dissolved immediately.

Parameters

- addition A SingleTimeAddition or SingleTimeGasAddition
- time The time point [s]

Returns This *MetalBulkZone* object

add_continuous_addition (*addition*: tc_python.process_metallurgy.process.AbstractContinuousAddition, from_time: float = 0.0, to_time: float = nan)

Adds a constant addition continuously during the specified time period to the zone. All added material will be dissolved immediately.

Parameters

- $\bullet \ \textbf{addition} A \ \textit{ContinuousAddition} \ \textbf{or} \ \textit{ContinuousGasAddition} \\$
- **from_time** The start time point [s]
- to_time The end time point [s]

Returns This MetalBulkZone object

add_power (*power: float, from_time: float = 0.0, to_time: float = nan*)

Adds a constant power during a specified time period to the zone (for example heating or cooling).

Parameters

- power The power [MW]
- **from_time** The start time point [s]
- to_time The end time point [s]

Returns This MetalBulkZone object

disable_degassing()

Disables degassing for this zone, i.e. all gas formed at any time step will be staying in this zone.

Returns This MetalBulkZone object

enable_degassing()

Enables degassing for this zone, i.e. any gas formed at any time step will be removed after that time step. This gas will be transferred into the *exhaust gas zone*. **This is the default**.

Returns This *MetalBulkZone* object

```
get_density() 
ightarrow float
```

Returns the density of the zone

Returns The density [kg/m**3]

```
get_elements() \rightarrow Set[str]
```

Returns the elements present in the zone. The elements are determined by the additions.

Returns The elements

 $\texttt{get_id()} \rightarrow \texttt{str}$

Returns the unique name / id of the zone.

Returns The zone name / id

 $\texttt{get_phase_group_to_transfer()} \rightarrow \textit{tc_python.process_metallurgy.base.PhaseGroup}$

Returns the phase group that is transferred from the attached reaction zones back to this zone after each time step.

Returns The phase group

is_degassing_enabled() \rightarrow bool

Returns if degassing is enabled in the zone.

Returns If degassing is enabled

class tc_python.process_metallurgy.process.ProcessSimulationCalculation(calculation)
 Bases: object

A Process Metallurgy process simulation. Such calculations represent complete metallurgical processes with several zones and simulate their evolution over time.

- **calculate** (*timeout_in_minutes: float* = 0.0) \rightarrow *tc_python.process_metallurgy.process.ProcessSimulationResult* Runs the Process Metallurgy process simulation.
 - Parameters timeout_in_minutes The calculation will be aborted after that time, default: no timeout

Returns A new ProcessSimulationResult object

set_end_time (*end_time: float*) Sets the end time of a process.

Parameters end_time – The end time point [s]

Returns This *ProcessSimulationCalculation* object

set_initial_time_step (initial_time_step: float = 1.0)
Sets the initial time step used in the process simulation.

Note: All later time steps are automatically determined to limit the expected temperature change during that step, this is controlled by *set_max_allowed_temp_change_per_step()*.

Parameters initial_time_step – The initial time step [s]

Returns This *ProcessSimulationCalculation* object

set_max_allowed_temp_change_per_step (max_allowed_temp_change: float = 10.0)

The maximum allowed temperature change per time step. This is implicitly also limiting the composition change during a time step and required for numerical stability.

Parameters max_allowed_temp_change – The maximum allowed temperature change [K]

Returns This *ProcessSimulationCalculation* object

set_max_time_step (max_time_step: float = 180.0)

The maximum time step chosen by the automatic time step control.

Note: All time steps are automatically determined to limit the expected temperature change during that step, this is controlled by *set_max_allowed_temp_change_per_step()*.

Parameters max_time_step – The maximum time step [s]

Returns This ProcessSimulationCalculation object

set_min_time_step (min_time_step: float = 1.0)

The minimum time step chosen by the automatic time step control.

Note: All time steps are automatically determined to limit the expected temperature change during that step, this is controlled by *set_max_allowed_temp_change_per_step()*.

Parameters min_time_step – The minimum time step [s]

Returns This *ProcessSimulationCalculation* object

set_pressure (pressure: float = 100000.0)

Sets a constant pressure during the complete process.

Parameters pressure – The pressure [Pa]

Returns This ProcessSimulationCalculation object

set_pressure_in_time_period (*pressure_in_pa: float, from_time: float = 0.0, to_time: float =*

nan

Sets a constant pressure during a time period.

Default: 1.0e5 Pa.

Parameters

- pressure_in_pa The pressure [Pa]
- **from_time** The start time [s]
- to_time The end time [s]

Returns This ProcessSimulationCalculation object

with_options (*options*: tc_python.process_metallurgy.base.ProcessMetallurgyOptions) Sets the options for the process simulation.

Parameters options – The options

Returns This ProcessSimulationCalculation object

with_reaction_zone (reaction_zone: tc_python.process_metallurgy.process.ReactionZone)

Sets the reaction zone of the process simulation. The bulk zones attached to this reaction zone are configured in the reaction zone object.

Note: In the present release, only one reaction zone is supported.

Parameters reaction_zone - The reaction zone object

Returns This *ProcessSimulationCalculation* object

class tc_python.process_metallurgy.process.ProcessSimulationResult(result)
 Bases: tc_python.abstract_base.AbstractResult

The result of a Process Metallurgy process simulation.

 $\texttt{get_activity_of_slag} (zone: Union[tc_python.process_metallurgy.process.Zone, str], component: str, reference: tc_python.process_metallurgy.base.ActivityReference = <ActivityReference.LIQUID: 1>) <math>\rightarrow$ List[float]

Returns the activity of a component in the slag in a zone at each time point.

Parameters

- zone The zone object or the zone name
- component The component
- **reference** The reference for the activity, can be liquid or solid slag, **default: liquid slag**

Returns The activity of the component at each time point [-]

get_amount (*zone: Union*[tc_python.process_metallurgy.process.Zone, *str*]) \rightarrow List[float] Returns the amount of a zone at each time point.

Parameters zone – The zone object or the zone name

Returns The amount at each time point [kg]

 $get_amount_of_elements() \rightarrow Dict[str, List[float]]$

Returns the total amount of each element in the simulation at each time point.

Returns The total amount of the elements at each time point [kg]

get_amount_of_phase_groups (zone: Union[tc_python.process_metallurgy.process.Zone,

 $str]) \rightarrow Dict[tc_python.process_metallurgy.base.PhaseGroup, List[float]]$

Returns the amount of each phase group (e.g., for example all liquid slag) in a zone at each time point.

Parameters zone – The zone object or the zone name

Returns The amount of the phase groups at each time point [kg]

get_amount_of_phases (*zone:* Union[tc_python.process_metallurgy.process.Zone, str]) \rightarrow Dict[str, List[float]]

Returns the amount of each phase in a zone at each time point.

Parameters zone – The zone object or the zone name

Returns The amount of the phases at each time point [kg]

get_components () \rightarrow Set[str] Returns all components defined in the simulation.

Returns The components

Dict[str, List[float]]

Returns the composition of a zone per element at each time point.

Parameters

- **zone** The zone object or the zone name
- composition_unit The composition unit, default: mass percent

Returns The composition at each time point

Returns the composition of a phase in a zone at each time point.

Parameters

- **zone** The zone object or the zone name
- **phase** The phase name
- composition_unit The composition unit, default: mass percent
- **composition_type** Defines if the composition is given by element (e.g., 75 wt-% Fe 25 wt-% Cr) or by component (e.g. 65 wt-% Al2O3 35 wt-% CaO). In case of a metallic phase, the composition is given by element even if *component* is selected. **Default: by component**.

Returns The composition at each time point

Returns the composition of a phase group (e.g., all liquid slag) in a zone at each time point.

Parameters

- **zone** The zone object or the zone name
- **phase_group** The phase group
- composition_unit The composition unit, default: mass percent
- composition_type Defines if the composition is given by element (e.g., 75 wt-% Fe 25 wt-% Cr) or by component (e.g. 65 wt-% Al2O3 35 wt-% CaO). In case of a metallic phase, the composition is given by element even if *component* is selected. Default: by component.

Returns The composition at each time point

```
get_elements() \rightarrow Set[str]
```

Returns all elements present in the simulation.

Returns The elements

 $get_enthalpy() \rightarrow List[float]$

Returns the total enthalpy of the process at each time point.

Returns The enthalpy at each time point [J]

get_exhaust_gas() \rightarrow tc_python.process_metallurgy.process.ExhaustGasResult Returns the result for the exhaust gas zone.

This result object can be used to evaluate the exhaust gas zone at each time point.

Returns The exhaust gas zone result object.

get_formula_for_activity_of_slag (zone: Union[tc_python.process_metallurgy.process.Zone, str], component: str, reference:

 $tc_python.process_metallurgy.base.ActivityReference = <ActivityReference.LIQUID: 1>) <math>\rightarrow$ List[str]

Returns the Thermo-Calc Console syntax formula used for calculating the activity of a component in the slag (e.g. *AC(AL2O3, IONIC_LIQ)* in a zone at each time point. The actual activity can be obtained using get_activity_of_slag().

Parameters

- **zone** The zone object or the zone name
- component The component
- **reference** The reference for the activity, can be liquid or solid slag, **default: liquid slag**

Returns The formula for calculating the activity at each time point

Returns the Thermo-Calc Console syntax formula used for calculating a property of the slag (e.g. B(CAO)/B(SIO2) in a zone at each time point. The actual slag property can be obtained using $get_slag_property()$.

Parameters

- **zone** The zone object or the zone name
- slag_property The slag property
- **slag_type** The part of the slag for which the property will be calculated. Can be all slag, the liquid or the solid slag. **Default: all slag**

Returns The formula for calculating the slag property at each time point

et_fraction_of_phase_groups(zone:		Union[tc_python.process_metallurgy.process.Zone,				
	str],	unit: tc_python.utils.PhaseUnit		utils.PhaseUnit	=	
	<i><phaseunit.mass_fraction:< i=""> 1>)</phaseunit.mass_fraction:<></i>				\rightarrow	
	Dict[tc_python.process_metallurgy.base.PhaseGroup,					
Determe the functions of the phase even	List[float]		、 ·			

Returns the fractions of the phase groups (e.g., all liquid slag) in a zone at each time point.

Parameters

- **zone** The zone object or the zone name
- **unit** The unit of the fraction

Returns The phase fractions at each time point

get_fraction_of_phases (zone: Union[tc_python.process_metallurgy.process.Zone, str], unit: tc python.utils.PhaseUnit = $\langle PhaseUnit.MASS \ FRACTION: 1 \rangle \rangle \rightarrow$

Dict[str, List[float]]

Returns the fractions of all stable phases in a zone at each time point.

Parameters

- **zone** The zone object or the zone name
- unit The unit of the fraction

Returns The phase fractions at each time point

get_gas_components() \rightarrow Set[str]

Returns all components of the gas phase defined for the elements present in the simulation.

Returns The components of the gas phase

get_num_of_performed_steps() \rightarrow List[int]

Returns the accumulated number of performed time steps at each time point.

Note: The number of performed time steps can differ from the index of the time step in the result list because time steps might have been repeated with smaller step size during a process simulation.

Returns The accumulated number of performed time steps

```
get_oxygen_partial_pressure(zone: Union[tc_python.process_metallurgy.process.Zone, str])
```

 \rightarrow List[float]

Returns the partial pressure of oxygen in the zone at each time point.

Parameters zone – The zone object or the zone name

Returns The partial pressure [Pa]

get_pressure (*zone: Union*[tc_python.process_metallurgy.process.Zone, *str*]) \rightarrow List[float] Returns the pressure in a zone at each time point.

Parameters zone – The zone object or the zone name

Returns The pressure at each time point [Pa]

get_slag_property (zone:Union[tc_python.process_metallurgy.process.Zone,str],slag_property:tc_python.process_metallurgy.base.SlagProperty,slag_type:tc_python.process_metallurgy.base.SlagType= <SlagType.ALL:</td>2>)

List[float] Returns a property of the slag in a zone at each time point. These properties are mostly used to describe

the property of a slag to pick up sulfur.

Parameters

- **zone** The zone object or the zone name
- slag_property The slag property
- **slag_type** The part of the slag for which the property will be calculated. Can be all slag, the liquid or the solid slag. **Default: all slag**

Returns The slag property at each time point [unit depending on the property]

get_stable_phases (*zone: Union*[tc_python.process_metallurgy.process.Zone, *str*]) \rightarrow Set[str] Returns the stable phases in a zone.

Parameters zone – The zone object or the zone name

Returns The stable phases

get_stable_phases_in_phase_group (zone: Union[tc_python.process_metallurgy.process.Zone,

str], *phase_group*: tc_python.process_metallurgy.base.PhaseGroup) → Set[str]

Returns the stable phases of a phase group (e.g., all solid slag) in a zone.

Parameters

- zone The zone object or the zone name
- phase_group The phase group

Returns The stable phases of the phase group

get_temperature (*zone: Union*[tc_python.process_metallurgy.process.Zone, *str*]) \rightarrow List[float] Returns the temperature of a zone at each time point.

Parameters zone – The zone object or the zone name

Returns The temperature at each time point [K]

get_time_points() \rightarrow List[float]

Returns the time points of the process simulation. All result quantities are returned for exactly these time points.

Returns The time points [s]

get_value_of (*zone:* Union[tc_python.process_metallurgy.process.Zone, str], classic_expression: str) \rightarrow List[float]

Returns a value for a thermodynamic quantity in a zone at each time point.

Warning: It should normally not be required to use this method, use the appropriate method available in the API instead.

Parameters

- **zone** The zone object or the zone name
- **classic_expression** The thermodynamic quantity to get the value of in Thermo-Calc Console Mode syntax (for example "NPM(FCC_A1)")

Returns The requested value at each time point

get_viscosity_dynamic_of_phase (*zone: Union*[tc_python.process_metallurgy.process.Zone, str], phase: str) \rightarrow List[float]

Returns the dynamic viscosity of a phase in a zone at each time point.

Parameters

- **zone** The zone object or the zone name
- **phase** The phase name

Returns The dynamic viscosity at each time point [Pa*s]

get_viscosity_kinematic_of_phase (zone: Union[tc_python.process_metallurgy.process.Zone,

str], phase: str) \rightarrow List[float] Returns the kinematic viscosity of a phase in a zone at each time point.

Parameters

- **zone** The zone object or the zone name
- phase The phase name

Returns The kinematic viscosity at each time point [m**2/s]

class tc_python.process_metallurgy.process.ReactionZone (area: float, left_zone:

tc_python.process_metallurgy.process.Zone, mass_transfer_coefficient_left: Union[float, tc_python.process_metallurgy.process.MassTransfe right_zone: tc_python.process_metallurgy.process.Zone, mass_transfer_coefficient_right: Union[float, tc_python.process_metallurgy.process.MassTransfe

Bases: tc_python.process_metallurgy.process.Zone

A reaction zone in a process simulation, this is representing the interface layer between two bulk zones that are in contact and can react with each other, for example the steel melt and the top slag. The size of the reaction zone is dynamic and determined by the mass transfer coefficient. A zone is a volume in a process that has identical temperature and composition. It has well-defined boundaries to other zones.

Adds a single-time addition at the specified time point to the zone. The addition will be dissolved immediately.

Parameters

- $\bullet \ \textbf{addition} A \ \textit{SingleTimeAddition} \ \textbf{or} \ \textit{SingleTimeGasAddition} \\$
- time The time point [s]

Returns This ReactionZone object

add_continuous_addition (addition: tc_python.process_metallurgy.process.AbstractContinuousAddition,

 $from_time: float = 0.0, to_time: float = nan)$

Adds a constant addition continuously during the specified time period to the zone. All added material will be dissolved immediately.

Parameters

- $\bullet \ \textbf{addition} A \ \textit{ContinuousAddition} \ \textbf{or} \ \textit{ContinuousGasAddition} \\$
- **from_time** The start time point [s]
- to_time The end time point [s]

Returns This *ReactionZone* object

add_heat_transfer(heat_transfer_coefficient: float)

Adds heat transfer through the reaction zone, i.e., between the two attached bulk zones.

Parameters heat_transfer_coefficient – The heat transfer coefficient [W/(K*m**2)]

Returns This ReactionZone object

add_power (*power: float, from_time: float = 0.0, to_time: float = nan*)

Adds a constant power during a specified time period to the zone (for example heating or cooling).

Parameters

• power – The power [MW]

- **from_time** The start time point [s]
- to_time The end time point [s]

Returns This ReactionZone object

add_transfer_of_phase_group (transfer_of_phase_group: tc_python.process_metallurgy.process.TransferOfPhaseGroup
Adds transfer of a certain phase group through the reaction zone during each time step, i.e. from one of
the attached bulk zones to the other. This is for example used to model inclusion flotation from the steel
melt to the slag.

Parameters transfer_of_phase_group – The transfer of phase group configuration, can be time-dependent.

Returns This *ReactionZone* object

disable_degassing()

Disables degassing for this zone, i.e. all gas formed at any time step will be staying in this zone.

Returns This ReactionZone object

enable_degassing()

Enables degassing for this zone, i.e. any gas formed at any time step will be removed after that time step. This gas will be transferred into the *exhaust gas zone*. **This is the default**.

Returns This ReactionZone object

 $\texttt{get_elements()} \rightarrow Set[str]$

Returns the elements present in the zone. The elements are determined by the additions.

Returns The elements

 $\texttt{get_id()} \rightarrow str$

Returns the unique id of the zone. :return: The zone id

$\texttt{is_degassing_enabled()} \rightarrow \texttt{bool}$

Returns if degassing is enabled in the zone.

Returns If degassing is enabled

class tc_python.process_metallurgy.process.SingleTimeAddition(composition:

Dict[str, float], amount: float, temperature: float = 293.15, composition_unit: tc_python.utils.CompositionUnit = <CompositionUnit.MASS_PERCENT: 0>, do_scale: bool = False)

Bases: tc_python.process_metallurgy.process.AbstractSingleTimeAddition

An addition in a process simulation that is added at a distinct time point.

It is assumed that the addition is dissolved instantaneously.

Tip: By setting $do_scale=True$, the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not summing to 100% / 1. An example could be a slag addition which is provided like this: 90 wt-% CaO - 5 wt-% Al2O3 - 4 wt-% SiO2.

Parameters

- composition The composition
- **amount** The amount [kg]
- temperature The initial addition temperature (default: 20 °C) [K]
- composition_unit The composition unit
- do_scale If the composition is scaled to 100% / fraction of 1

get_amount() \rightarrow float

Returns the amount of this addition.

Returns The amount [kg]

get_composition_unit () \rightarrow *tc_python.utils.CompositionUnit* Returns the composition unit used in this addition.

Returns The composition unit

class tc_python.process_metallurgy.process.SingleTimeGasAddition(composition:

Dict[str, float], amount: float, temperature: float = 293.15. amount unit: tc_python.utils.GasAmountUnit <GasAmountUnit.NORM_CUBIC_METER: 0>, composition_unit: tc_python.utils.GasCompositionUnit <Gas-CompositionUnit.VOLUME PERCENT: 4>. do_scale: bool = False) Bases: tc_python.process_metallurgy.process.AbstractSingleTimeAddition

A gas addition in a process simulation that is added at a distinct time point.

It is assumed that the addition is dissolved instantaneously.

Tip: By setting $do_scale=True$, the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not scaling to 100% / 1. An example could be a gas addition which is provided like this: 90 vol-% Ar - 10 vol-% O2.

Parameters

• **composition** – The composition

- **amount** The amount
- temperature The initial addition temperature (default: 20 °C) [K]
- amount_unit The amount unit
- composition_unit The composition unit
- do_scale If the composition is scaled to 100% / fraction of 1

$\texttt{get}_\texttt{amount}() \rightarrow \texttt{float}$

Returns the amount of this addition.

Note: The amount unit can be obtained using get_amount_unit().

Returns The amount [in the amount unit]

get_amount_unit() \rightarrow tc_python.utils.GasAmountUnit Returns the amount unit used in this addition.

Returns The amount unit

get_composition_unit() \rightarrow tc_python.utils.GasCompositionUnit Returns the composition unit used in this addition.

Returns The composition unit

class tc_python.process_metallurgy.process.**SlagBulkZone**(*density: float*) Bases: tc_python.process_metallurgy.process.Zone

A slag bulk zone in a process simulation.

This is representing a large volume in the process, for example the top slag. A zone is a volume in a process that has identical temperature and composition. It has well-defined boundaries to other zones.

The name of this zone is automatically defined and unique.

add_addition (addition: tc_python.process_metallurgy.process.AbstractSingleTimeAddition, time:

float = 0.0)

Adds a single-time addition at the specified time point to the zone. The addition will be dissolved immediately.

Parameters

- addition A SingleTimeAddition or SingleTimeGasAddition
- time The time point [s]

Returns This SlagBulkZone object

add_continuous_addition (addition: tc_python.process_metallurgy.process.AbstractContinuousAddition, from time: float = 0.0, to time: float = nan)

Adds a constant addition continuously during the specified time period to the zone. All added material will be dissolved immediately.

Parameters

- addition A ContinuousAddition or ContinuousGasAddition
- from_time The start time point [s]
- to_time The end time point [s]

Returns This *SlagBulkZone* object

add_power (power: float, from_time: float = 0.0, to_time: float = nan)

Adds a constant power during a specified time period to the zone (for example heating or cooling).

Parameters

- power The power [MW]
- from_time The start time point [s]
- to_time The end time point [s]

Returns This *SlagBulkZone* object

disable_degassing()

Disables degassing for this zone, i.e. all gas formed at any time step will be staying in this zone.

Returns This *SlagBulkZone* object

enable_degassing()

Enables degassing for this zone, i.e. any gas formed at any time step will be removed after that time step. This gas will be transferred into the *exhaust gas zone*. **This is the default**.

Returns This *SlagBulkZone* object

```
get_density() \rightarrow float
```

Returns the density of the zone

Returns The density [kg/m**3]

$get_elements() \rightarrow Set[str]$

Returns the elements present in the zone. The elements are determined by the additions.

Returns The elements

$get_id() \rightarrow str$

Returns the unique id of the zone. :return: The zone id

get_phase_group_to_transfer() \rightarrow tc_python.process_metallurgy.base.PhaseGroup

Returns the phase group that is transferred from the attached reaction zones back to this zone after each time step.

Returns The phase group

is_degassing_enabled() \rightarrow bool

Returns if degassing is enabled in the zone.

Returns If degassing is enabled

class tc_python.process_metallurgy.process.**TransferOfPhaseGroup**(*phase_group_to_transfer:*

tc_python.process_metallurgy.base.Pha
source_zone:
tc python.process metallurgy.process.2

```
Bases: object
```

The transfer of a percentage of a certain phase group (.e.g., solid slag) between zones during each time step. This is for example used to model inclusion flotation from the steel melt to the slag.

add (*transfer_rate: float*, *time: float* = 0)

Adds the transfer rate valid beginning at a time point.

This value is valid until another value is defined for a later time point.

Parameters

- **transfer_rate** The transfer rate [% of phase group amount/s]
- time The time point where the transfer of a phase group begins to be valid [s]

Returns This *TransferOfPhaseGroup* object

get_phase_group_to_transfer() \rightarrow *tc_python.process_metallurgy.base.PhaseGroup* Returns the phase group to be transferred

Returns The phase group

get_transfer_source_zone_id() \rightarrow str

The id of the source zone of the transfer

Returns This source zone id

class tc_python.process_metallurgy.process.Zone

Bases: object

The base class of a zone in a process simulation. A zone is a volume in a process that has identical temperature and composition. It has well-defined boundaries to other zones.

add_addition (addition: tc_python.process_metallurgy.process.AbstractSingleTimeAddition, time:

float = 0.0)

Adds a single-time addition at the specified time point to the zone. The addition will be dissolved immediately.

Parameters

- addition A SingleTimeAddition or SingleTimeGasAddition
- time The time point [s]

Returns This *Zone* object

add_continuous_addition (*addition*: tc_python.process_metallurgy.process.AbstractContinuousAddition, from time: float = 0.0, to time: float = nan)

Adds a constant addition continuously during the specified time period to the zone. All added material will be dissolved immediately.

Parameters

- $\bullet \ \textbf{addition} A \ \textit{ContinuousAddition} \ \textbf{or} \ \textit{ContinuousGasAddition} \\$
- **from_time** The start time point [s]
- to_time The end time point [s]

Returns This Zone object

add_power (*power*: *float*, *from_time*: *float* = 0.0, *to_time*: *float* = *nan*)

Adds a constant power during a specified time period to the zone (for example heating or cooling).

Parameters

- power The power [MW]
- from_time The start time point [s]
- to_time The end time point [s]

Returns This Zone object

disable_degassing()

Disables degassing for this zone, i.e. all gas formed at any time step will be staying in this zone.

Returns This Zone object

enable_degassing()

Enables degassing for this zone, i.e. any gas formed at any time step will be removed after that time step. This gas will be transferred into the *exhaust gas zone*. **This is the default**.

Returns This *Zone* object

 $\texttt{get_elements}() \rightarrow \texttt{Set[str]}$

Returns the elements present in the zone. The elements are determined by the additions.

Returns The elements

 $\texttt{get_id}() \rightarrow str$

Returns the unique id of the zone. :return: The zone id

 $\texttt{is_degassing_enabled()} \rightarrow bool$

Returns if degassing is enabled in the zone.

Returns If degassing is enabled

5.2 Module "system"

class tc_python.system.MultiDatabaseSystemBuilder(multi_database_system_builder)
Bases: object

Used to select databases, elements, phases etc. and create a System object. The difference to the class System-Builder is that the operations are performed on all the previously selected databases. The system is then used to create calculations.

create_and_select_species (stoichiometry: str)

Specify a species from the already entered elements. The stoichiometry of the species is the chemical formula of the species. The created species will also be automatically selected.

Note: The elements in the chemical formula are normally separated by stoichiometric numbers. Neither parenthesis "()" nor an underscore "_" is allowed in the chemical formula, while the special combination "/-" or "/+" can be used. Consult the Thermo-Calc database documentation for details about the syntax.

Parameters stoichiometry – The stoichiometry of the species

Returns This MultiDatabaseSystemBuilder object

deselect_constituent_on_sublattice (phase_name: str, sublattice_no: int, constituent_name_to_deselect: str)

Rejects a constituent on a sublattice in a phase in both the thermodynamic and the kinetic database.

Parameters

- phase_name The name of the phase
- **sublattice_no** The number of the sublattice (starting with 1)
- constituent_name_to_deselect The name of the constituent to deselect

Returns This MultiDatabaseSystemBuilder object

deselect_phase (phase_name_to_deselect: str)

Rejects a phase for both the thermodynamic and the kinetic database.

Parameters phase_name_to_deselect - The phase name

Returns This MultiDatabaseSystemBuilder object

deselect_species (*species_name: str*) Removes the species from the system.

Parameters species_name - The species

Returns This MultiDatabaseSystemBuilder object

$get_system() \rightarrow tc_python.system.System$

Creates a new System object that is the basis for all calculation types. Several calculation types can be defined later from the object; these are independent.

Returns A new System object

select_constituent_on_sublattice (phase_name: str, sublattice_no: int, constituent_name_to_select: str)

Selects a constituent on a sublattice in a phase in both the thermodynamic and the kinetic database.

Note: Previously the third parameter *constituent_name_to_select* had a wrong name, it has been corrected in version 2021b.

Parameters

- phase_name The name of the phase
- **sublattice_no** The number of the sublattice (starting with 1)
- constituent_name_to_select The name of the constituent to select

Returns This MultiDatabaseSystemBuilder object

select_phase (phase_name_to_select: str)

Selects a phase for both the thermodynamic and the kinetic database.

Parameters phase_name_to_select - The phase name

Returns This MultiDatabaseSystemBuilder object

select_species (species_name: str)

Adds the species to the system. Up to 1000 species can be defined in a single system.

Parameters species_name – The species

Returns This MultiDatabaseSystemBuilder object

with_new_composition_set (composition_set: tc_python.entities.CompositionSet)

Used to enter two or more composition sets for a phase. If a phase has a miscibility gap it is necessary to have two composition sets, one for each possible composition that can be stable simultaneously.

The databases often create the typical composition sets for phases automatically when data are retrieved. The equilibrium calculations (using the default settings with global minimization) will usually add new composition sets if needed.

Note: Precipitation and diffusion calculations can require the user to define additional composition sets. E.g. in the case where the new composition set is needed in the configuration of the calculation.

Parameters composition_set - the composition set

Returns This MultiDatabaseSystemBuilder object

without_default_phases()

Rejects all the default phases from both the thermodynamic and the kinetic database, any phase now needs to be selected manually for the databases.

Returns This MultiDatabaseSystemBuilder object

class tc_python.system(system_instance)

Bases: object

A system containing selections for databases, elements, phases etc.

Note: For the defined system, different calculations can be configured and run. **Instances of this class should always be created from a SystemBuilder**.

Note: The system object is **immutable**, i.e. it cannot be changed after is has been created. If you want to change the system, you must instead create a new one.

<pre>convert_composition (input_composition:</pre>	Dict[str;	float],	input_unit:
tc_python.utils.ConversionUnit,			output_unit:
tc_python.utils.ConversionUnit,	dependent	_component:	$str = ") \rightarrow$
Dict[str, float]			1 1 1 6

Provides conversion between composition units for any combination of chemical compounds. It is fast because no thermodynamic equilibrium calculation is involved.

Syntax of the chemical compounds: "Al2O3", "FeO", "CO", "Fe", "C", ...

Note: It is not required that the chemical compounds are components of the database. The only requirement is that all elements are present in the database.

Parameters

- input_composition Composition (for example: {"Al2O3": 25.0, "FeO": 75.0})
- **input_unit** Unit of the input composition
- **output_unit** Requested output unit
- **dependent_component** The dependent component (optional), for example: "*Fe*". If no dependent component is specified the sum of the input composition needs to match 100% / 1

Returns The composition in the requested output unit

get_all_elements_in_databases() \rightarrow List[str]

Returns the names of all elements present in the selected databases, regardless of the actual selection of elements.

Returns A list of element names

get_all_phases_in_databases() \rightarrow List[str]

Returns all phase names present in the selected databases, regardless of selected elements, phases etc.

Returns A list of phase names

$\texttt{get_all_species_in_databases()} \rightarrow List[str]$

Returns all species names present in the selected databases, regardless of the actual selection of elements, phases,

Returns A list of species names

```
get_element_object (element_name: str) → tc_python.entities.Element
```

Returns the Element object of an element. This can be used to obtain detailed information about the element.

Parameters element_name - The element name

Returns A Element: object

```
get_elements_in_system() \rightarrow List[str]
```

Returns the names of all elements present in the selected system.

Note: The list does not contain any elements or components that have been auto-selected by the database(s) in a calculator. Use the get_components() of the calculator object instead to get the complete information.

Returns A list of element names

```
get_phase_object (phase_name: str) \rightarrow tc_python.entities.Phase
```

Returns the Phase object of a phase. This can be used to obtain detailed information about the phase.

Parameters phase_name - The phase name

Returns A Phase: object

get_phases_in_system() \rightarrow List[str]

Returns all phase names present in the system due to its configuration (selected elements, phases, etc.).

Returns A list of phase names

get_references () \rightarrow Dict[str, List[str]]

Provides a dictionary with database references per database in the selected system.

Returns The database references

get_species_in_system() \rightarrow List[str]

Returns the names of all species present in the selected system.

Note: The list does not contain any species or components that have been auto-selected by the database(s) in a calculator. Use the get_components() of the calculator object instead to get the complete information.

Returns The list of species names

get_species_object (species_name: str) \rightarrow tc_python.entities.Species

Returns the Species object of an species. This can be used to obtain detailed information about the species.

Parameters species_name – The species name

Returns A Species: object

$get_system_data() \rightarrow tc_python.abstract_base.SystemData$

Returns the content of the database. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

with batch equilibrium calculation (default conditions: bool = True. List[str] components: = []) \rightarrow tc python.batch equilibrium.BatchEquilibriumCalculation Creates a batch-equilibrium calculation (a vectorized equilibrium calculation).

Note: Use this instead of looping if you want to calculate equilibria for a larger number of compositions and know the conditions in advance. This calculation type has improved performance when calculating a large number of equilibria when each individual calculations is quick. E.g. when evaluating single phase properties for thousands of compositions.

Parameters

- default_conditions If True, automatically sets the conditions N=1 and P = 100000
- components Specify here the components of the system (for example: [AL2O3, ...]), only necessary if they differ from the elements. If this option is used, all elements of the system need to be replaced by a component.

Returns A new BatchEquilibriumCalculation object

with_cct_precipitation_calculation () \rightarrow tc_python.precipitation.PrecipitationCCTCalculation Creates a CCT diagram calculation.

Returns A new PrecipitationCCTCalculation object

with_isothermal_diffusion_calculation () \rightarrow tc_python.diffusion.DiffusionIsoThermalCalculation Creates an isothermal diffusion calculation.

Returns A new DiffusionIsoThermalCalculation object

with isothermal_precipitation_calculation () \rightarrow tc_python.precipitation.PrecipitationIsoThermalCalculation Creates an isothermal precipitation calculation.

Returns A new PrecipitationIsoThermalCalculation object

with_material_to_material() \rightarrow tc_python.material_to_material.MaterialToMaterialCalculationContainer Provides access to all Material to Material calculations. The actual calculation needs to be chosen in the returned object.

Returns A new MaterialToMaterialCalculationContainer object

with_non_isothermal_diffusion_calculation () \rightarrow tc_python.diffusion.DiffusionNonIsoThermalCalculation Creates a non-isothermal precipitation calculation.

. . .

Returns A new PrecipitationNonIsoThermalCalculation object

-

.

with_non_isothermal_precipitation_calculation	\rightarrow () \rightarrow
Creates a non-isothermal precipitation calculation.	$tc_python.precipitation.PrecipitationNonIsoThermalCalcondent for the state of the$
Returns A new PrecipitationNonIsoTherma	Calculation object
with_phase_diagram_calculation(default_conditions	bool = True,
components:	$List[str] = []) \rightarrow$
	map_diagrams.PhaseDiagramCalculation
Creates a phase diagram (map) calculation	

eates a phase diagram (map) calculation.

Parameters

. .

. . .

- **default_conditions** If *True*, automatically sets the conditions N=1 and P=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

Returns A new PhaseDiagramCalculation object

```
with_property_diagram_calculation (default_conditions: bool = True,

components: List[str] = []) \rightarrow

tc_python.step_or_map_diagrams.PropertyDiagramCalculation
```

Creates a property diagram (step) calculation.

Parameters

- **default_conditions** If *True*, automatically sets the conditions N=1 and P=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

Returns A new PropertyDiagramCalculation object

with_property_model_calculation	(model	: str,	path_to_mod	dels:	str	=
	΄',	debug_model:	bool	=	False)	\rightarrow
	$tc_python.propertymodel.PropertyModelCalculation$					

Creates a Property Model calculation.

The parameter *debug_model* is only used when debugging self-developed models.

Parameters

- **model** The Property Model to be calculated.
- **path_to_models** The path where the Property Models are installed. If no value is entered, the Property Models folder used by the normal Thermo-Calc application is used.
- **debug_model** Used when debugging self-developed models.

Returns A new PropertyModelCalculation object

```
with_scheil_calculation() \rightarrow tc_python.scheil.ScheilCalculation
```

Creates a Scheil solidification calculation.

Warning: Scheil calculations do not support the *GAS* phase being selected, this means the `GAS` phase must always be deselected in the system if it is present in the database

Returns A new ScheilCalculation object

```
with_single_equilibrium_calculation (default_conditions: bool = True,

components: List[str] = []) \rightarrow

tc python.single equilibrium.SingleEquilibriumCalculation
```

Creates a single equilibrium calculation.

Parameters

• **default_conditions** – If *True*, automatically sets the conditions N=1 and P=100000

• **components** – Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

Returns A new SingleEquilibriumCalculation object

with_ttt_precipitation_calculation() \rightarrow *tc_python.precipitation.PrecipitationTTTCalculation* Creates a TTT diagram calculation.

Returns A new PrecipitationTTTCalculation object

class tc_python.system.SystemBuilder(system_builder)

Bases: object

Used to select databases, elements, phases etc. and create a System object. The system is then used to create calculations.

create_and_select_species (stoichiometry: str)

Specify a species from the already entered elements. The stoichiometry of the species is the chemical formula of the species. The created species will also be automatically selected.

Note: The elements in the chemical formula are normally separated by stoichiometric numbers. Neither parenthesis "()" nor an underscore "_" is allowed in the chemical formula, while the special combination "/-" or "/+" can be used. Consult the Thermo-Calc database documentation for details about the syntax.

Parameters stoichiometry – The stoichiometry of the species

Returns This SystemBuilder object

deselect_constituent_on_sublattice (phase_name: str, sublattice_no: int, con-

stituent_name_to_deselect: str) Rejects a constituent on a sublattice in a phase in the last specified database only.

Parameters

- phase_name The name of the phase
- **sublattice_no** The number of the sublattice (starting with 1)
- constituent_name_to_deselect The name of the constituent to deselect

Returns This SystemBuilder object

deselect_phase (*phase_name_to_deselect: str*) Rejects a phase in the last specified database only.

Parameters phase_name_to_deselect - The name of the phase

Returns This *SystemBuilder* object

deselect_species (*stoichiometry: str*) Removes the species from the system.

Parameters stoichiometry – The species

Returns This SystemBuilder object

 $get_system() \rightarrow tc_python.system.System$

Creates a new System object that is the basis for all calculation types. Several calculation types can be defined later from the object; these are independent.

Returns A new System object

$get_system_for_scheil_calculations() \rightarrow tc_python.system.System$

Creates a new System object **without gas phases being selected**, that is the basis for all calculation types, but its particularly useful for Scheil solidification calculations, where the model does not allow that a gas phase is selected in the system.

Several calculation types can be defined later from the object; these are independent.

Note: Deprecated in version 2022b: Use get_system() instead. There are no longer any special requirements from the Thermo-Calc program to deselect the gas phase for Scheil. It will be removed in release 2023b.

Returns A new System object

select_constituent_on_sublattice (phase_name: str, sublattice_no: int, constituent_name_to_select: str) Selects a constituent on a sublattice in a phase in the last specified database only.

Note: Previously the third parameter *constituent_name_to_select* had a wrong name, it has been corrected in version 2021b.

Parameters

- phase_name The name of the phase
- **sublattice_no** The number of the sublattice (starting with 1)
- constituent_name_to_select The name of the constituent to select

Returns This SystemBuilder object

select_database_and_elements (*database_name: str, list_of_element_strings: List[str]*)

Selects a thermodynamic or kinetic database and its selected elements (that will be appended). After that, phases can be selected or unselected.

Parameters

- database_name The database name, for example "FEDEMO"
- list_of_element_strings A list of one or more elements as strings, for example ["Fe", "C"]

Returns This SystemBuilder object

select_phase (phase_name_to_select: str)

Selects a phase in the last specified database only.

Parameters phase_name_to_select - The name of the phase

Returns This SystemBuilder object

select_species (stoichiometry: str)

Adds the species to the system. Up to 1000 species can be defined in a single system.

Parameters stoichiometry – The species

Returns This SystemBuilder object

select_user_database_and_elements(path_to_user_database: str, list_of_element_strings:

List[str])

Selects a thermodynamic database which is a user-defined database and select its elements (that will be appended).

Note: By using a r-literal, it is possible to use slashes on all platforms, also on Windows: *select_user_database_and_elements(r"my path/user_db.tdb", ["Fe", "Cr"]])*

Otherwise it is required to use **double** back-slashes on Windows as separator.

Note: On Linux and Mac the path is case-sensitive, also the file ending.

Parameters

- **path_to_user_database** The path to the database file ("database".TDB), defaults to the current working directory. Only the filename is required if the database is located in the same folder as the script.
- list_of_element_strings A list of one or more elements as strings, for example ["Fe", "C"]

Returns This SystemBuilder object

with_new_composition_set (composition_set: tc_python.entities.CompositionSet)

Used to enter composition sets for a phase. If a phase has a miscibility gap it is necessary to have two composition sets, one for each possible composition that can be stable simultaneously.

Parameters composition_set – The composition set

Returns This SystemBuilder object

without_default_phases()

Rejects all default phases in the last specified database only, any phase needs now to be selected manually for that database.

Returns This SystemBuilder object

5.3 Module "entities"

```
class tc_python.entities.CompositionSet (phase_name: str)
```

Bases: object

Used by the method tc_python.system.SystemBuilder.with_new_composition_set() to enter two or more composition sets for a phase.

Parameters phase_name – The name of the phase for which a new composition set is required

set_major_constituents_for_sublattice (sublattice_index: int, major_constituents:

List[str]) Specify the new major constituent(s) for the sublattice.

Default: If not specified, a default is automatically chosen based on the specified composition set.

Note: This is useful in order to make calculations converge faster and more easily (because it may simplify giving start values when calculating the equilibrium as those phases with miscibility gaps should

have different major constituents for each composition set). The databases often set major constituents for several phases automatically when the data is retrieved.

Parameters

- **sublattice_index** Index of the sublattice to set the major constituents for (starting with *I*)
- **major_constituents** Optional list of the major constituents, which must be selected from the phase constitution of the current system.

Returns This CompositionSet object

```
class tc_python.entities.Element (element)
```

```
Bases: object
```

Represents an element, making detailed information about the element accessible.

```
get_enthalpy() \rightarrow float
```

Returns the enthalpy of the element at 298 K, part of the stable element reference state (SER).

Returns The enthalpy [J]

```
get_entropy_diff_0_to_298k() \rightarrow float
```

Returns the entropy difference 0 - 298 K of the element, part of the stable element reference state (SER).

Returns The entropy difference 0 - 298 K [J/K]

get_molar_mass() ightarrow float

Returns the molar mass of the element.

Returns The molar mass [g/mol]

 $\texttt{get_name()} \rightarrow str$

Returns the name of the element.

Returns The element name

get_stable_element_reference() \rightarrow str

Returns the stable element reference (i.e. the stable phase at 298.15 K and 1 bar, reference for all element thermodynamic data).

Returns The name of the stable element reference

$\texttt{is_interstitial()} \rightarrow bool$

Returns if the element is interstitial.

Note: In the diffusion simulations (DICTRA), the assumption that the volume is carried by the substitutional elements only is applied. The interstitial elements are assumed to have zero molar volumes.

Returns If the element is interstitial

```
is_special() \rightarrow bool
```

Returns if the element is special (i.e. vacancies (VA) and electrons (denoted either as /- in gaseous, liquid or solid phases, or ZE in an aqueous solution phase)).

Returns If the element is special

```
\texttt{is\_valid()} \rightarrow bool
```

Returns if the element is valid. Non-valid elements are represented by an empty name.

Returns If the element is valid

```
class tc_python.entities.Phase(phase)
```

Bases: object

Represents a phase, making detailed information about the phase accessible.

 $\begin{array}{l} \texttt{get_name} \ (\) \ \rightarrow \ \texttt{str} \\ \text{Returns the name of the phase.} \end{array}$

Returns The phase name

get_species () \rightarrow Set[*tc_python.entities.Species*] Returns the species of the phase.

Returns A set containing the species

get_species_for_composition_profile() \rightarrow Set[tc_python.entities.Species]

Returns all species that need to be defined in a composition profile of the phase for diffusion simulations - except for one species that needs to be the dependent species.

Note: In a composition profile of a phase for diffusion simulations it is necessary to specify all non-stoichiometric and non-special species. In case of a DILUTE diffusion model, the database enforces the choice of a certain dependent species.

Returns Set with the species

```
get_sublattices() \rightarrow List[tc_python.entities.Sublattice]
```

Returns the sublattices of the phase in a well-defined contiguous order.

Returns A list containing the Sublattice objects

 $get_type() \rightarrow tc_python.entities.PhaseType$ Returns the type of the phase (liquid, ionic liquid, solid, gas).

Returns The type of a phase

has_diffusion_data () \rightarrow bool Returns if diffusion data exists for the phase.

Returns If diffusion data exists for the phase

 $\texttt{has_molar_volume_data()} \rightarrow bool$

Returns if molar volume data exists for the phase.

Returns If molar volume data exists for the phase

```
\texttt{is\_dilute\_diffusion\_model()} \rightarrow bool
```

Returns if diffusion is described using the DILUTE model for the phase. This will always return *False* if no diffusion data is available.

Returns If the DILUTE model is used

 $\texttt{is_gas}() \rightarrow \texttt{bool}$

Returns if the phase is a gas phase.

Returns If the phase is a gas phase

 $\texttt{is_ionic_liquid}() \rightarrow \texttt{bool}$

Returns if the phase is an ionic liquid phase.

Returns If the phase is an ionic liquid phase

 $\texttt{is_liquid()} \rightarrow bool$

Returns if the phase is a liquid or ionic liquid phase.

Returns If the phase is a liquid phase

 $is_solid() \rightarrow bool$

Returns if the phase is a solid phase.

Returns If the phase is a solid phase

```
class tc_python.entities.PhaseType (value)
```

Bases: enum.Enum

The type of a phase.

GAS = 0

Gas phase.

```
IONIC_LIQUID = 2
Ionic liquid phase.
```

LIQUID = 1 Liquid phase.

SOLID = 3 Solid phase.

```
class tc_python.entities.Species(species)
```

Bases: object

Represents a species, making detailed information about the species accessible.

```
get_all_elements () \rightarrow List[Tuple[tc_python.entities.Element, float]]
Returns all the elements that the species is composed of.
```

Returns List of all elements of the species and their stoichiometry

```
\texttt{get\_charge()} \rightarrow \texttt{int}
```

Returns the charge of the species.

Returns The charge of the species

```
get_name() 
ightarrow str
```

Returns the name of the species.

Returns The species name

```
\texttt{is\_element}() \rightarrow bool
```

Returns if the species actually represents an element.

Returns If the species represents an element

```
is_interstitial () \rightarrow bool
Returns if the species is interstitial.
```

Note: In the diffusion simulations (DICTRA), the assumption that the volume is carried by the substitutional elements only is applied. The interstitial elements are assumed to have zero molar volumes.

Returns If the species is interstitial

 $\texttt{is_special()} \rightarrow bool$

Returns if the species is special (i.e. vacancies (VA) and electrons (denoted either as /- in gaseous, liquid or solid phases, or ZE in an aqueous solution phase)).

Returns If the species is special

 $\texttt{is_valid}() \rightarrow \texttt{bool}$

Returns if the species is valid. Non-valid species are represented by an empty name.

Returns If the species is valid

to_element() \rightarrow *tc_python.entities.Element*

Returns the *Element* representation of the species - if the species actually represents an element.

Returns The Element object

class tc_python.entities.Sublattice(sublattice)

Bases: object

Represents a sublattice of a phase.

get_constituents () \rightarrow Set[*tc_python.entities.Species*] Returns the constituents of the sublattice.

Returns A set containing the constituents

 $get_nr_of_sites() \rightarrow float$ Returns the number of sites in the sublattice.

Returns A float number

5.4 Module "server"

```
class tc_python.server.LoggingPolicy(value)
```

```
Bases: enum.Enum
```

Logging policy that determines how the TC-Python logs are presented to the user.

FILE = 1

Logging to a file.

NONE = 2

No logging at all.

```
SCREEN = 0
```

Logging to the screen.

class tc_python.server.MetallurgyCalculations(metallurgy_calculations)
 Bases: object

Provides access to the calculation objects for all Process Metallurgy calculations.

These are specialised calculations for working with metallurgical processes. Both equilibrium calculations and kinetic process simulations (Effective Equilibrium Reaction Zone model) are available.

with_adiabatic_equilibrium_calculation (*database:* tc_python.process_metallurgy.base.ProcessDatabase)

 \rightarrow tc_python.process_metallurgy.equilibrium.AdiabaticEquilibriumCa Creates an adiabatic equilibrium calculation for Process Metallurgy.

Parameters database - The thermodynamic database used in the calculation

Returns A new AdiabaticEquilibriumCalculation object

with_adiabatic_process_calculation (database: tc_python.process_metallurgy.base.ProcessDatabase) \rightarrow tc_python.process_metallurgy.process.ProcessSimulationCalculation

Creates an adiabatic kinetic process simulation (EERZ, i.e. Effective Equilibrium Reaction Zone model).

Parameters database – The thermodynamic database used in the calculation

Returns A new ProcessSimulationCalculation object

with_isothermal_equilibrium_calculation(*database*: tc_python.process_metallurgy.base.ProcessDatabase)

 \rightarrow tc_python.process_metallurgy.equilibrium.IsoThermalEquilibrium Creates an isothermal equilibrium calculation for Process Metallurgy.

Parameters database - The thermodynamic database used in the calculation

Returns A new IsoThermalEquilibriumCalculation object

class tc_python.server.ResultLoader(result_loader)

Bases: object

Contains methods for loading results from previously done calculations.

diffusion (*path: str*) \rightarrow *tc_python.diffusion.DiffusionCalculationResult* Loads a DiffusionCalculationResult from disc.

Parameters path – path to the folder where result was previously saved.

Returns A new DiffusionCalculationResult object which later can be used to get specific values from the calculated result

phase_diagram(path: str) → tc_python.step_or_map_diagrams.PhaseDiagramResult Loads a PhaseDiagramResult from disc.

Parameters path – path to the folder where result was previously saved.

Returns A new PhaseDiagramResult object which later can be used to get specific values from the calculated result

precipitation_TTT_or_CCT (path: str) → tc_python.precipitation.PrecipitationCalculationTTTorCCTResult Loads a PrecipitationCalculationTTTorCCTResult from disc.

Parameters path – path to the folder where result was previously saved.

Returns A new PrecipitationCalculationTTTorCCTResult object which later can be used to get specific values from the calculated result

precipitation_single (path: str) → tc_python.precipitation.PrecipitationCalculationSingleResult
Loads a PrecipitationCalculationSingleResult from disc.

Parameters path – path to the folder where result was previously saved.

Returns A new PrecipitationCalculationSingleResult object which later can be used to get specific values from the calculated result

property_diagram (*path: str*) → *tc_python.step_or_map_diagrams.PropertyDiagramResult* Loads a PropertyDiagramResult from disc.

Parameters path – path to the folder where result was previously saved.

Returns A new PropertyDiagramResult object which later can be used to get specific values from the calculated result

property_model (path: str) → tc_python.propertymodel.PropertyModelResult Loads a PropertyModelResult from disc.

Parameters path – path to the folder where result was previously saved.

Returns A new PropertyModelResult object which later can be used to get specific values from the calculated result

scheil (path: str) → tc_python.scheil.ScheilCalculationResult
Loads a ScheilCalculationResult from disc.

Parameters path – path to the folder where result was previously saved.

Returns A new ScheilCalculationResult object which later can be used to get specific values from the calculated result

single_equilibrium(path: str) → tc_python.single_equilibrium.SingleEquilibriumResult Loads a SingleEquilibriumResult from disc.

Parameters path – path to the folder where result was previously saved.

Returns A new SingleEquilibriumResult object which later can be used to get specific values from the calculated result

class tc_python.server.**SetUp**(*debug_logging=False*)

Bases: object

Starting point for all calculations.

Note: This class exposes methods that have no precondition, it is used for choosing databases and elements.

disable_caching()

A previously set cache folder is no longer used.

Note: Within the session, caching is activated and used through the default temporary directory.

Returns This *SetUp* object

get_database_info($database_short_name: str$) \rightarrow str

Obtains the short information available for the specified database.

Parameters database_short_name – The name of the database (i.e. "FEDEMO", ...)

Returns The short information about the database

Obtains the path to the database file on disk. *TCPATH* is a placeholder for the root path of the used Thermo-Calc installation.

Note: Encrypted databases (*.*TDC*) cannot be edited.

Parameters database_short_name – The name of the database (i.e. "FEDEMO", ...)

Returns The path to the database on disk

$\texttt{get_databases()} \rightarrow List[str]$

Obtains the short names of all databases available in the used Thermo-Calc installation.

Note: Only databases with a valid license are listed.

Returns List of the available databases

Lists the names of all Property Models in the specified directory.

If the directory is not specified, the Property Model folder used by the normal Thermo-Calc application is used.

Parameters path_to_models – The path where the Property Models are installed. If no value is entered, the Property Model folder used by the normal Thermo-Calc application is used.

Returns Set containing all Property Model names

 $load_result_from_disk() \rightarrow tc_python.server.ResultLoader$

Loads a previously calculated result from disk.

Note: This only works for results created by calling one of the save_result() methods on a *Result* class created from a calculation.

Returns A new ResultLoader object

```
select_database_and_elements(database_name: str, list_of_elements: List[str]) \rightarrow tc_python.system.SystemBuilder
```

Selects a first thermodynamic or kinetic database and selects the elements in it.

Parameters

- database_name The name of the database, for example "FEDEMO"
- **list_of_elements** The list of the selected elements in that database, for example ["Fe", "C"]

Returns A new SystemBuilder object

select_thermodynamic_and_kinetic_databases_with_elements(thermodynamic_db_name:

str, kinetic_db_name: str, list_of_elements: List[str]) \rightarrow

tc_python.system.MultiDatabaseSystemBuil

Selects the thermodynamic and kinetic database at once, guarantees that the databases are added in the correct order. Further rejection or selection of phases applies to both databases.

Parameters

- thermodynamic_db_name The thermodynamic database name, for example "FEDEMO"
- kinetic_db_name The kinetic database name, for example "MFEDEMO"
- **list_of_elements** The list of the selected elements in that database, for example ["Fe", "C"]

Returns A new MultiDatabaseSystemBuilder object

select_user_database_and_elements ($path_to_user_database$: str, $list_of_elements$: List[str]) $\rightarrow tc_python.system.SystemBuilder$

Selects a user-defined database and selects the elements in it.

Note: By using a r-literal, it is possible to use slashes on all platforms, also on Windows: *select_user_database_and_elements(r"my path/user_db.tdb", ["Fe", "Cr"]])*

Otherwise it is required to use **double** back-slashes on Windows as separator.

Note: On Linux and Mac the path is case-sensitive, also the file ending.

Parameters

- **path_to_user_database** The path to the database file ("database".TDB), defaults to the current working directory. Only filename is required if the database is located in the same folder as the script.
- list_of_elements The list of the selected elements in that database, for example ["Fe", "C"]

Returns A new SystemBuilder object

set_cache_folder (path: str = ", precision_for_floats: int = 12)

Sets a folder where results from calculations and state of systems are saved. If at any time a calculation is run which has the exact same setting as a previous, the calculation is not re-run. The result is instead loaded from this folder.

Note: The same folder can be used in several scripts, and it can even be shared between different users. It can be a network folder.

Parameters

- path path to the folder where results should be stored. It can be relative or absolute.
- **precision_for_floats** The number of significant figures used when comparing if the calculation has the same setting as a previous.

Returns This SetUp object

$set_ges_version (version: int = 6)$

Setting the version of the Gibbs Energy System (GES).

Parameters version – The GES-version (currently version 5 or 6)

Returns This *SetUp* object

set_log_level_to_debug() Sets log level to DEBUG

sets log level to DEBUG

Returns This *SetUp* object

```
set_log_level_to_info()
```

Sets log level to INFO

Returns This SetUp object

```
with_metallurgy() \rightarrow tc_python.server.MetallurgyCalculations
Provides access to the calculation objects for all Process Metallurgy calculations.
```

These are specialised calculations for working with metallurgical processes. Both equilibrium calculations and kinetic process simulations (Effective Equilibrium Reaction Zone model) are available.

Bases: object

Starting point of the API. Typical syntax:

```
with TCPython() as session:
    session.select_database_and_elements(...)
```

Note: Each usage of *with TCPython()* causes significant overhead (starting a new process, stopping the old one, cleaning up the temporary disk space). Usually it is recommendable to call *with TCPython()* only once for each process, even if working in a loop. Instead you should pass the session or calculator object into the loop and use them there.

If necessary, beginning from version 2019a it is however possible to call with TCPython safely multiple times.

Starts a process of the API server and sets up the socket communication with it.

Parameters

- **logging_policy** Determines if the TC-Python log output is sent to the screen (*LoggingPolicy.SCREEN*), to file (*LoggingPolicy.FILE*) or nothing is logged at all (*LoggingPolicy.NONE*) **Default**: *LoggingPolicy.SCREEN*. Note that the log-handlers can also be adapted through the *tc_python.LOGGER* object at any time.
- **log_file** The log-file relative to the current path or absolute, only relevant if *log-ging_policy=LoggingPolicy.FILE*. Log-output will be appended.
- debug_mode If *True* it is tried to open a connection to an already running API-server. This is only used for debugging the API itself.
- is_unittest Should be True if called by a unit test, only to be used internally for development.
- **do_throw_on_backend_hard_crash** If *True* an *UnrecoverableCalculationException* will be thrown if the Java-backend crashes hard, if *False* the application will simply crash with a FORTRAN-stacktrace. *If* `*True*` *the exception can be caught outside of the* `*with*`-clause and the application can continue, if `*False*` more information about the error is shown by the stacktrace.
- port_number The port number for the communication with the Java-backend server. This is not required to be changed by normal users.

Warning: Most users should use *TCPython* using a with-statement for automatic management of the resources (network sockets and temporary files). If you anyway need to use that method, make sure to call *stop_api_server()* in any case using the try-finally-pattern.

tc_python.server.stop_api_server(gateway_id: str = None)

Clears all resources used by the session (i.e. shuts down the API server and deletes all temporary files). The disk usage of temporary files might be significant.

Warning: Call this method only if you used *start_api_server()* initially. It should never be called when the API has been initialized in a with-statement using *TCPython*.

5.5 Module "quantity_factory"

class tc_python.quantity_factory.DiffusionQuantity

Bases: tc_python.quantity.AbstractQuantity

Factory class providing quantities used for defining diffusion simulations and their results.

Note: In this factory class only the most common quantities are defined, you can always use the *Console Mode* syntax strings in the respective methods as an alternative (for example: "NPM(*)").

```
classmethod activity_of_component (component: str, use_ser: bool = False) \rightarrow tc_python.quantity.ActivityOfComponent
```

Creates a quantity representing the activity of a component.

Parameters

- component The name of the component, use ALL_COMPONENTS to choose all components
- **use_ser** Use Stable-Element-Reference(SER). The user-defined reference state is be used if this setting is set to *False*.

Returns A new ActivityOfComponent object.

classmethod	chemical_	diffusion_	_coefficien	t (phase:	str,	diffus	ing_eleme	ent:
				str,	gradient_	element:		str,
				referenc	e_element:		str)	\rightarrow
_	_			tc_pythe	n.quantity.	Chemica	Diffusion	Coefficient

Creates a quantity representing the chemical diffusion coefficient of a phase [m^2/s].

Parameters

- **phase** The name of the phase
- diffusing_element The diffusing element
- gradient_element The gradient element
- reference_element The reference element (for example "Fe" in a steel)

Returns A new ChemicalDiffusionCoefficient object.

classmethod chemical_potential_of_component (component: str, use_ser: bool = False) \rightarrow tc_python.quantity.ChemicalPotentialOfComponent

Creates a quantity representing the chemical potential of a component [J].

Parameters

- component The name of the component, use ALL_COMPONENTS to choose all components
- **use_ser** Use Stable-Element-Reference(SER). The user-defined reference state is used if this setting is set to *False*.

Returns A new ChemicalPotentialOfComponent object.

classmethod distance (*region:* str = 'All') \rightarrow tc_python.quantity.Distance Creates a quantity representing the distance [m].

Parameters region – The name of the region or *All* to choose global.

classmethod intrinsic_diffusion_coefficient (*phase:* str, diffusing_element: str, gradient_element: str, reference_element: str) \rightarrow tc_python.quantity.IntrinsicDiffusionCoefficient

Creates a quantity representing the intrinsic diffusion coefficient of a phase [m²/s].

Parameters

- phase The name of the phase
- diffusing_element The diffusing element
- gradient_element The gradient element
- reference_element The reference element (for example "Fe" in a steel)

Returns A new IntrinsicDiffusionCoefficient object.

classmethod l_bis (*phase: str, diffusing_element: str, gradient_element: str, reference_element: str*) \rightarrow tc_python.quantity.Lbis

Creates a quantity representing \vec{L} ' of a phase $[m^2/s]$.

Parameters

- **phase** The name of the phase
- diffusing_element The diffusing element
- gradient_element The gradient element
- reference_element The reference element (for example "Fe" in a steel)

Returns A new Lbis object.

classmethod mass_fraction_of_a_component (component: str) \rightarrow tc_python.quantity.MassFractionOfAComponent

Creates a quantity representing the mass fraction of a component.

Parameters component – The name of the component or *ALL_COMPONENTS* to choose all components

Returns A new MassFractionOfAComponent object.

```
classmethod mass_fraction_of_a_phase (phase: str) -
tc_python.quantity.MassFractionOfAPhase
Creates a quantity representing the mass fraction of a phase.
```

Parameters phase – The name of the phase or *ALL_PHASES* to choose all phases.

Returns A new MassFractionOfAPhase object.

classmethod mobility_of_component_in_phase(phase: str, component: str) \rightarrow tc python.quantity.MobilityOfComponentInPhase Creates a quantity representing the mobility of a component in a phase [m²/Js]. **Parameters** • **phase** – The name of the phase • component – The name of the component Returns A new MobilityOfComponentInPhase object. classmethod mole_fraction_of_a_component (component: str) tc_python.quantity.MoleFractionOfAComponent Creates a quantity representing the mole fraction of a component. **Parameters** component – The name of the component or ALL_COMPONENTS to choose all components Returns A new MoleFractionOfAComponent object. classmethod mole_fraction_of_a_phase (phase: str) tc_python.quantity.MoleFractionOfAPhase Creates a quantity representing the mole fraction of a phase. Parameters phase – The name of the phase or ALL_PHASES to choose all phases Returns A new MoleFractionOfAPhase object. classmethod position_of_lower_boundary_of_region(region: str) tc_python.quantity.PositionOfLowerBoundaryOfRegion Creates a quantity representing the position of lower boundary of a region [m]. Parameters region – The name of the region **Returns** A new PositionOfLowerBoundaryOfRegion object. classmethod position_of_upper_boundary_of_region(region: str) tc_python.quantity.PositionOfUpperBoundaryOfRegion Creates a quantity representing the position of upper boundary of a region [m]. **Parameters** region – The name of the region **Returns** A new PositionOfUpperBoundaryOfRegion object. **classmethod temperature** () \rightarrow tc_python.quantity.Temperature Creates a quantity representing the temperature [K]. Returns A new Temperature object. gradiclassmethod thermodynamic_factor (phase: *diffusing_element:* str. str. ent element: str, reference element: str) tc_python.quantity.ThermoDynamicFactor Creates a quantity representing thermodynamic factor of a phase. **Parameters** • **phase** – The name of the phase diffusing_element – The diffusing element • gradient_element - The gradient element • reference_element – The reference element (for example "Fe" in a steel) Returns A new ThermoDynamicFactor object.

```
classmethod time () \rightarrow tc_python.quantity.Time
    Creates a quantity representing the time [s].
classmethod total_mass_fraction_of_component(component:
                                                                                 str)
                                                           tc_python.quantity.TotalMassFractionOfComponent
    Creates a quantity representing the total mass fraction of a component.
        Parameters component – The name of the component
         Returns A new TotalMassFractionOfComponent object.
classmethod total mass fraction of component in phase (phase:
                                                                                  str.
                                                                                        com-
                                                                       ponent:
                                                                                          \rightarrow
                                                                                   str)
                                                                       tc python.quantity.TotalMassFractionOfCompo
    Creates a quantity representing the total mass fraction of a component in a phase.
        Parameters
             • phase – The name of the phase
             • component – The name of the component
         Returns A new TotalMassFractionOfComponentInPhase object.
classmethod total_mass_fraction_of_phase(phase:
                                                                             str)
                                                     tc_python.quantity.TotalMassFractionOfPhase
    Creates a quantity representing the total mass fraction of a phase.
         Parameters phase – The name of the phase.
         Returns A new TotalMassFractionOfPhase object.
classmethod total_mole_fraction_of_component(component:
                                                                                 str)
                                                           tc_python.quantity.TotalMoleFractionOfComponent
    Creates a quantity representing the total mole fraction of a component.
        Parameters component - The name of the component
         Returns A new TotalMoleFractionOfComponent object.
classmethod total_mole_fraction_of_component_in_phase(phase:
                                                                                  str,
                                                                                        com-
                                                                       ponent:
                                                                                   str)
                                                                                          \rightarrow
                                                                       tc_python.quantity.TotalMoleFractionOfCompo
    Creates a quantity representing the total mole fraction of a component in a phase.
        Parameters
             • phase – The name of the phase
             • component – The name of the component
         Returns A new TotalMoleFractionOfComponentInPhase object.
classmethod total_volume_fraction_of_phase (phase:
                                                                              str)
                                                        tc_python.quantity.TotalVolumeFractionOfPhase
    Creates a quantity representing the total volume fraction of a phase.
        Parameters phase – The name of the phase.
        Returns A new TotalVolumeFractionOfPhase object.
classmethod tracer_diffusion_coefficient (phase: str, diffusing_element: str) \rightarrow
                                                      tc python.quantity.TracerDiffusionCoefficient
    Creates a quantity representing tracer diffusion coefficient of a phase [m^2/s].
        Parameters
             • phase – The name of the phase
```

• diffusing_element – The diffusing element

Returns A new TracerDiffusionCoefficient object.

classmethod u_fraction_of_a_component(component:

str)

tc_python.quantity.UFractionOfAComponent Creates a quantity representing the u-fraction of a component.

ties a quantity representing the a maction of a component.

Parameters component – The name of the component

Returns A new UFractionOfAComponent object.

classmethod user_defined_function (*expression: str*) \rightarrow tc_python.quantity.Function Creates a quantity representing a user-defined function.

Parameters expression – The function expression

Returns A new Function object

classmethod velocity_of_lower_boundary_of_region(region: str) -

tc_python.quantity.VelocityOfLowerBoundaryOfRegior Creates a quantity representing the velocity of lower boundary of a region [m/s].

Parameters region – The name of the region

Returns A new VelocityOfLowerBoundaryOfRegion object.

classmethod velocity_of_upper_boundary_of_region (region: str) \rightarrow

tc_python.quantity.VelocityOfUpperBoundaryOfRegior Creates a quantity representing the velocity of upper boundary of a region [m/s].

Parameters region – The name of the region

Returns A new VelocityOfUpperBoundaryOfRegion object.

classmethod width_of_region (*region: str*) \rightarrow tc_python.quantity.Function Creates a quantity representing the width of a region [m].

Parameters region – The name of the region

Returns A new WidthOfRegion object.

class tc_python.quantity_factory.IndependentVariable Bases: tc_python.quantity.AbstractQuantity

Factory class providing quantities used for defining the independent variable in general diffusion result querying.

classmethod distance (*region: str* = '*All*') \rightarrow tc_python.quantity.Distance Creates an independent variable representing the distance [m].

Returns A new Distance object

classmethod time () \rightarrow tc_python.quantity.Time Creates an independent variable representing the time [s].

Returns A new Time object

class tc_python.quantity_factory.PlotCondition

Bases: tc_python.quantity.AbstractQuantity

Factory class providing quantities used for defining the plot condition in general diffusion result querying.

Note: In this factory class only the most common quantities are defined, you can always use the *Console Mode* syntax strings in the respective methods as an alternative (for example: "time last").

classmethod distance (distance point: float, region: str = 'All') \rightarrow tc_python.quantity.DistanceCondition

Creates a plot condition representing the distance [m].

Change in version 2019b: Mandatory parameter distancepoint added

Parameters

- distancepoint The distance from the lower interface of the region
- **region** The name of the region or *All* to choose global.

Returns A new DistanceCondition object

classmethod integral () \rightarrow tc_python.quantity.IntegralCondition Creates an integral plot condition.

Returns A new IntegralCondition object

classmethod interface (*region: str, interface_position:* tc_python.utils.InterfacePosition) \rightarrow tc_python.quantity.InterfaceCondition

Creates a plot condition representing an interface between two regions.

Parameters

- **region** The name of the region used for defining the interface
- **interface_position** The position of the interface relative to that region (lower or upper)

Returns A new InterfaceCondition object

classmethod time (*timepoint: Union[float, str]* = 'Last') \rightarrow tc_python.quantity.TimeCondition Creates a plot condition representing the time [s].

Change in version 2019b: Lists of timepoints are no longer supported

Parameters timepoint – The timepoint. Optionally "Last" can be used for the end of the simulation

Returns A new TimeCondition object

class tc_python.quantity_factory.ScheilQuantity
 Bases: tc_python.quantity.AbstractQuantity

Factory class providing quantities used for defining a Scheil calculation result (*tc_python.scheil*. *ScheilCalculationResult*).

classmethod apparent_heat_capacity_per_gram() \rightarrow tc_python.quantity.ApparentHeatCapacityPerGram Creates a quantity representing the apparent heat capacity [J/g/K].

Returns A new ApparentHeatCapacityPerGram object.

classmethod apparent_heat_capacity_per_mole() \rightarrow tc_python.quantity.ApparentHeatCapacityPerMole Creates a quantity representing the apparent heat capacity [J/mol/K].

Returns A new ApparentHeatCapacityPerMole object.

classmethod apparent_volumetric_thermal_expansion_coefficient()

tc_python.quantity.ApparentVolume

Creates a quantity representing the apparent volumetric thermal expansion coefficient of the system [1/K].

Returns A new ApparentVolumetricThermalExpansionCoefficient object.

```
classmethod composition_of_phase_as_mole_fraction (phase: str, component: str) \rightarrow
```

tc_python.quantity.CompositionOfPhaseAsMoleFracti

Creates a quantity representing the composition of a phase [mole-fraction].

Parameters

- **phase** The name of the phase, use ALL_PHASES to choose all stable phases
- component The name of the component, use ALL_COMPONENTS to choose all components

Returns A new CompositionOfPhaseAsMoleFraction object.

classmethod composition_of_phase_as_weight_fraction (phase: str, component: str) \rightarrow

tc_python.quantity.CompositionOfPhaseAsWeight

Creates a quantity representing the composition of a phase [weight-fraction].

Parameters

- **phase** The name of the phase, use *ALL_PHASES* to choose all stable phases
- component The name of the component, use ALL_COMPONENTS to choose all components

Returns A new CompositionOfPhaseAsWeightFraction object.

classmethod density_of_phase (*phase*: *str*) \rightarrow tc_python.quantity.DensityOfPhase Creates a quantity representing the average density of a phase [g/cm^3].

Parameters phase – The name of the phase or ALL_PHASES to choose all phases

Returns A new DensityOfPhase object.

classmethod density_of_solid_phase (*phase*: *str*) \rightarrow tc_python.quantity.DensityOfSolidPhase Creates a quantity representing the average density of a solid phase [g/cm^3].

Note: Deprecated in version 2022a: This quantity has been renamed to *density_of_phase()*. It will be removed in release 2023a.

Parameters phase – The name of the phase or ALL_PHASES to choose all solid phases

Returns A new DensityOfSolidPhase object.

classmethod density_of_system() \rightarrow tc_python.quantity.DensityOfSystem Creates a quantity representing the average density of the system [g/cm^3].

Returns A new DensityOfSystem object.

 $\texttt{classmethod distribution_of_component_of_phase} (\textit{phase: str, component: str}) \rightarrow \texttt{classmethod distribution_of_component_of_phase} (\texttt{phase: str, component: str}) \rightarrow \texttt{classmethod distribution_of_component_of_phase} (\texttt{phase: str, component: str}) \rightarrow \texttt{classmethod distribution_of_component} (\texttt{phase: str, component: str}) \rightarrow \texttt{classmethod distribution_of_component_of_phase} (\texttt{phase: str, component: str}) \rightarrow \texttt{classmethod distribution_of_component_of_phase} (\texttt{phase: str, component: str}) \rightarrow \texttt{classmethod distribution_of_component: str})$

tc_python.quantity.DistributionOfComponentOfPhase Creates a quantity representing the (molar) fraction of the specified component being present in the specified phase compared to the overall system [-]. This corresponds to the degree of segregation to that phase.

Parameters

- **phase** The name of the phase
- **component** The name of the component

Returns A new DistributionOfComponentOfPhase object.

classmethod heat_per_gram() \rightarrow tc_python.quantity.HeatPerGram

Creates a quantity representing the total heat release from the liquidus temperature down to the current temperature [J/g].

Note: The total or apparent heat release during the solidification process consists of two parts: one is the so-called latent heat, i.e. heat due to the liquid -> solid phase transformation (*latent_heat_per_mole(*) and *latent_heat_per_gram(*)), and the other is the heat related to the specific heat of liquid and solid phases (*heat_per_mole(*) and *heat_per_gram(*)).

Returns A new HeatPerGram object.

classmethod heat_per_mole() \rightarrow tc_python.quantity.HeatPerMole

Creates a quantity representing the total heat release from the liquidus temperature down to the current temperature [J/mol].

Note: The total or apparent heat release during the solidification process consists of two parts: one is the so-called latent heat, i.e. heat due to the liquid -> solid phase transformation (*latent_heat_per_mole(*) and *latent_heat_per_gram(*)), and the other is the heat related to the specific heat of liquid and solid phases (*heat_per_mole(*) and *heat_per_gram(*)).

Returns A new HeatPerMole object.

$\texttt{classmethod latent_heat_per_gram()} \rightarrow tc_python.quantity.LatentHeatPerGram$

Creates a quantity representing the cumulated latent heat release from the liquidus temperature down to the current temperature [J/g].

Note: The total or apparent heat release during the solidification process consists of two parts: one is the so-called latent heat, i.e. heat due to the liquid -> solid phase transformation (*latent_heat_per_mole(*) and *latent_heat_per_gram(*)), and the other is the heat related to the specific heat of liquid and solid phases (*heat_per_mole(*) and *heat_per_gram(*)).

Returns A new LatentHeatPerGram object.

classmethod latent_heat_per_mole() \rightarrow tc_python.quantity.LatentHeatPerMole

Creates a quantity representing the cumulated latent heat release from the liquidus temperature down to the current temperature [J/mol].

Note: The total or apparent heat release during the solidification process consists of two parts: one is the so-called latent heat, i.e. heat due to the liquid -> solid phase transformation (*latent_heat_per_mole(*) and *latent_heat_per_gram(*)), and the other is the heat related to the specific heat of liquid and solid phases (*heat_per_mole(*) and *heat_per_gram(*)).

Returns A new LatentHeatPerMole object.

Parameters phase - The name of the phase or ALL_PHASES to choose all solid phases

Returns A new MassFractionOfASolidPhase object.

classmethod mass_fraction_of_all_liquid() \rightarrow tc_python.quantity.MassFractionOfAllLiquid Creates a quantity representing the total mass fraction of all the liquid phase.

Returns A new MassFractionOfAllLiquid object.

classmethod mass_fraction_of_all_solid_phases()

tc_python.quantity.MassFractionOfAllSolidPhase Creates a quantity representing the total mass fraction of all solid phases.

Returns A new MassFractionOfAllSolidPhase object.

classmethod molar_volume_of_phase (*phase: str*) \rightarrow tc_python.quantity.MolarVolumeOfPhase Creates a quantity representing the molar volume of a phase [m^3/mol].

Parameters phase – The name of the phase or ALL_PHASES to choose all phases

Returns A new MolarVolumeOfPhase object.

classmethod molar_volume_of_system () \rightarrow tc_python.quantity.MolarVolumeOfSystem Creates a quantity representing the molar volume of the system [m³/mol].

Returns A new MolarVolumeOfSystem object.

 $\texttt{classmethod mole_fraction_of_a_solid_phase} (phase: str) \rightarrow$

tc_python.quantity.MoleFractionOfASolidPhase Creates a quantity representing the molar fraction of a solid phase.

Parameters phase - The name of the phase or ALL_PHASES to choose all solid phases

Returns A new MoleFractionOfASolidPhase object.

classmethod mole_fraction_of_all_liquid () \rightarrow tc_python.quantity.MoleFractionOfAllLiquid Creates a quantity representing the total molar fraction of all the liquid phase.

Returns A new MoleFractionOfAllLiquid object.

classmethod mole_fraction_of_all_solid_phases()

tc_python.quantity.MoleFractionOfAllSolidPhases Creates a quantity representing the total molar fraction of all solid phases.

Returns A new MoleFractionOfAllSolidPhases object.

classmethod site_fraction_of_component_in_phase ((phase:	str,	comp	onent:
	str,	sub_latti	ce_ordina	al_no:
	int	=	0)	\rightarrow
	tc pythe	on.quantity	.SiteFrac	tionOfComponentInPhase

Creates a quantity representing the site fractions [-].

Parameters

- phase The name of the phase, use ALL_PHASES to choose all stable phases
- **component** The name of the component, use *ALL_COMPONENTS* to choose all components
- **sub_lattice_ordinal_no** The ordinal number (i.e. 1, 2, ...) of the sublattice of interest, use None to choose all sublattices

Note: Detailed information about the sublattices can be obtained by getting the *Phase* object of a phase from the *System* object using tc_python.system.get_phase_in_system. For each phase the sublattices are obtained by using tc_python.system.Phase.get_sublattices. The

order in the returned list is equivalent to the sublattice ordinal number expected, **but note that the ordinal numbers start with 1**.

Returns A new SiteFractionOfComponentInPhase object.

classmethod temperature () \rightarrow tc_python.quantity.Temperature Creates a quantity representing the temperature [K].

Returns A new Temperature object.

class tc_python.quantity_factory.ThermodynamicQuantity
 Bases: tc_python.quantity.AbstractQuantity

Factory class providing quantities used for defining equilibrium calculations (single equilibrium, property and phase diagrams, \ldots) and their results.

Note: In this factory class only the most common quantities are defined, you can always use the *Console Mode* syntax strings in the respective methods as an alternative (for example: "NPM(*)").

classmethod activity_of_component (component: str, use_ser: bool = False) \rightarrow tc_python.quantity.ActivityOfComponent

Creates a quantity representing the activity of a component [-].

Parameters

- component The name of the component, use ALL_COMPONENTS to choose all components
- **use_ser** Use Stable-Element-Reference(SER). The user-defined reference state is used if this setting is set to *False*.

Returns A new ActivityOfComponent object.

classmethod chemical_diffusion_coefficient (phase: str, diffusing_element: str, gradient_element: str, reference_element: str) \rightarrow tc_python.quantity.ChemicalDiffusionCoefficient

Creates a quantity representing the chemical diffusion coefficient of a phase $[m^2/s]$.

Parameters

- **phase** The name of the phase
- diffusing_element The diffusing element
- gradient_element The gradient element
- **reference_element** The reference element (for example "Fe" in a steel)

Returns A new ChemicalDiffusionCoefficient object.

classmethod chemical_potential_of_comp	onent (component:	str,	use_ser:	
	bool =	False)	\rightarrow	

tc_python.quantity.ChemicalPotentialOfComponent

Creates a quantity representing the chemical potential of a component [J].

Parameters

component – The name of the component, use ALL_COMPONENTS to choose all components

• **use_ser** – Use Stable-Element-Reference(SER). The user-defined reference state is used if this setting is set to *False*.

Returns A new ChemicalPotentialOfComponent object.

```
classmethod composition_of_phase_as_mole_fraction (phase: str, compo-
```

nent: $str = 'All' \rightarrow$ tc_python.quantity.CompositionOfPhaseAsMoleFraction

Creates a quantity representing the composition of a phase [mole-fraction].

Parameters

- **phase** The name of the phase, use *ALL_PHASES* to choose all stable phases
- component The name of the component, use ALL_COMPONENTS to choose all components

Returns A new CompositionOfPhaseAsMoleFraction object.

classmethod composition_of_phase_as_weight_fraction(phase: str, com-

ponent: str) \rightarrow

tc_python.quantity.CompositionOfPhaseAsWeightl

Creates a quantity representing the composition of a phase [weight-fraction].

Parameters

- **phase** The name of the phase, use *ALL_PHASES* to choose all stable phases
- component The name of the component, use ALL_COMPONENTS to choose all components

Returns A new CompositionOfPhaseAsWeightFraction object.

classmethod gibbs_energy_of_a_phase (*phase:* str, use_ser: bool = False) \rightarrow tc_python.quantity.GibbsEnergyOfAPhase Creates a quantity representing the Gibbs energy of a phase [J].

Parameters

- phase The name of the phase or ALL_PHASES to choose all phases
- **use_ser** Use Stable-Element-Reference(SER). The user-defined reference state will be used when this setting is set to False.

Returns A new GibbsEnergyOfAPhase object.

classmethod mass_	_fraction_of_a	_component	(component:		str)	\rightarrow
				 -		

tc_python.quantity.MassFractionOfAComponent Creates a quantity representing the mass fraction of a component.

Parameters component – The name of the component or *ALL_COMPONENTS* to choose all components

Returns A new MassFractionOfAComponent object.

 $\begin{array}{cl} \texttt{classmethod mass_fraction_of_a_phase} (phase: & str) & \rightarrow \\ & & & \\ & & & \\ & & & \\ & &$

Parameters phase – The name of the phase or *ALL_PHASES* to choose all phases.

Returns A new MassFractionOfAPhase object.

classmethod mole_fraction_of_a_component (*component:* str) \rightarrow tc_python.quantity.MoleFractionOfAComponent Creates a quantity representing the mole fraction of a component.

Parameters component – The name of the component or *ALL_COMPONENTS* to choose all components

Returns A new MoleFractionOfAComponent object.

classmethod mole_fraction_of_a_phase (*phase*: str) tc_python.quantity.MoleFractionOfAPhase Creates a quantity representing the mole fraction of a phase.

Parameters phase – The name of the phase or *ALL_PHASES* to choose all phases

Returns A new MoleFractionOfAPhase object.

classmethod normalized_driving_force_of_a_phase (*phase*: str) \rightarrow tc_python.quantity.NormalizedDrivingForceOfAPhase

Creates a quantity representing normalized driving force of a phase [-].

Warning: A driving force calculation requires that the respective phase has been set to the state *DORMANT*. The parameter *All* is only reasonable if all phases have been set to that state.

Parameters phase – The name of the phase or ALL_PHASES to choose all phases

Returns A new DrivingForceOfAPhase object.

classmethod pressure () \rightarrow tc_python.quantity.Pressure Creates a quantity representing the pressure [Pa].

Returns A new Pressure object.

classmethod system_size () \rightarrow tc_python.quantity.SystemSize Creates a quantity representing the system size [mol].

Returns A new SystemSize object.

classmethod temperature () \rightarrow tc_python.quantity.Temperature Creates a quantity representing the temperature [K].

Returns A new Temperature object.

classmethod tracer_diffusion_coefficient (*phase: str, diffusing_element: str*) \rightarrow tc python.quantity.TracerDiffusionCoefficient

Creates a quantity representing tracer diffusion coefficient of a phase [m²/s].

Parameters

- **phase** The name of the phase
- diffusing_element The diffusing element

Returns A new TracerDiffusionCoefficient object.

classmethod u_fraction_of_a_component(component: str) -

tc_python.quantity.UFractionOfAComponent

Creates a quantity representing the u-fraction of a component.

Parameters component – The name of the component

Returns A new UFractionOfAComponent object.

classmethod user_defined_function (*expression: str*) \rightarrow tc_python.quantity.Function Creates a quantity representing a user-defined function.

Parameters expression – The function expression

Returns A new Function object

classmethod volume_fraction_of_a_phase (*phase:* str) \rightarrow tc_python.quantity.VolumeFractionOfAPhase Creates a quantity representing the volume fraction of a phase.

Parameters phase – The name of the phase or ALL_PHASES to choose all phases

Returns A new VolumeFractionOfAPhase object.

5.6 Module "utils"

```
class tc_python.utils.CompositionType (value)
     Bases: enum.Enum
     The type of composition.
     COMPONENT = 1
         Composition given per component, this will be identical to ELEMENT in case of metals.
     ELEMENT = 0
         Composition given per element.
class tc_python.utils.CompositionUnit(value)
     Bases: enum.Enum
     The composition unit.
     MASS FRACTION = 1
         Mass fraction.
     MASS PERCENT = 0
         Mass percent.
     MOLE FRACTION = 3
         Mole fraction.
     MOLE PERCENT = 2
         Mole percent.
class tc_python.utils.ConversionUnit(value)
     Bases: enum. Enum
     The composition unit used in a conversion.
     MOLE FRACTION = 0
         Mole fraction.
     MOLE PERCENT = 1
         Mole percent.
     WEIGHT_FRACTION = 2
         Weight fraction.
     WEIGHT_PERCENT = 3
         Weight percent.
class tc_python.utils.GasAmountUnit(value)
     Bases: enum.Enum
     The amount of a gas.
```

KILOGRAM = 1 Kilogram.

NORM_CUBIC_METER = 0

Norm cubic meter (according to ISO 2533, p=101325 Pa, T=288.15 K, typically used for trading of gas bottles). Other definitions vary only slightly.

```
class tc_python.utils.GasCompositionUnit(value)
```

Bases: enum.Enum

The composition unit for a gas.

```
MASS_FRACTION = 1
```

Mass fraction.

MASS_PERCENT = 0 Mass percent.

MOLE_FRACTION = 3 Mole fraction.

MOLE_PERCENT = 2 Mole percent.

VOLUME_FRACTION = 5 Volume fraction.

VOLUME_PERCENT = 4 Volume percent.

```
class tc_python.utils.GasRateUnit(value)
    Bases: enum.Enum
```

The rate of a gas flow.

```
KILOGRAM_PER_SEC = 1
```

Kilogram per second.

NORM_CUBIC_METER_PER_SEC = 0

Norm cubic meter per second (according to ISO 2533, p=101325 Pa, T=288.15 K, typically used for trading of gas bottles). Other definitions vary only slightly.

```
class tc_python.utils.InterfacePosition(value)
```

Bases: enum.Enum

The position of an interface relative to its region. Only used for diffusion simulations.

LOWER = 0

The interface is on the lower side of its region.

UPPER = 1

The interface is on the upper side of its region.

class tc_python.utils.PhaseUnit(value)

Bases: enum.Enum

The units available for a phase fraction.

MASS_FRACTION = 1 Mass fraction.

MOLE_FRACTION = 0 Mole fraction.

```
VOLUME_FRACTION = 2
Volume fraction.
```

```
class tc_python.utils.ResultValueGroup (result_line_group_java)
Bases: object
```

A x-y-dataset representing a line data calculation result (i.e. a Thermo-Calc quantity 1 vs. quantity 2).

Warning: Depending on the calculator, the dataset might contain *NaN*-values to separate the data between different subsets.

Variables

- label a str describing what the data corresponds to
- **x** list of floats representing the first quantity ("x-axis")
- **y** list of floats representing the second quantity ("y-axis")

```
get_label() \rightarrow str
Accessor for the line label :return the line label
```

```
get_x () \rightarrow List[float]
Accessor for the x-values :return the x values
```

- $\texttt{get}_y () \rightarrow \texttt{List[float]}$ Accessor for the y-values :return the y values
- class tc_python.utils.TemperatureProfile
 Bases: object

Represents a time-temperature profile used by non-isothermal calculations.

Note: The total simulation time can differ from the defined temperature profile. Constant temperature is assumed for any timepoint after the end of the defined profile.

add_time_temperature (*time: float*, *temperature: float*) Adds a time-temperature point to the non-isothermal temperature profile.

Parameters

- time The time [s]
- temperature The temperature [K]

Returns This TemperatureProfile object

5.7 Module "propertymodel_sdk"

class tc_python.propertymodel_sdk.CCTResult (quantity_id: str, description: str)
 Bases: tc_python.propertymodel_sdk.ResultQuantity

Represents a Continuous Cooling (CCT) result.

Parameters

quantity_id – The id of this result

- **description** The description of this result
- add_time_temperature (time_temperature_id: str, description: str)

Adds a time-temperature pair to the result.

Parameters

- time_temperature_id The id of the time-temperature pair
- **description** The description of the time-temperature pair

temperature_suffix = '(T)'
The temperature suffix of a CCTResult

time_suffix = ' (t) '
The time suffix of a CCTResult

Bases: object

Represents Continuous Cooling (CCT) result values.

Parameters

- cooling_rate The cooling rate [K/s]
- cooling_rate_start_temperature The start temperature of cooling [K]
- cooling_rate_end_temperature The end temperature of cooling [K]

set_result_time_temperature (time_temperature_id: str, time: float, temperature: float)
Sets a time-temperature pair of the result.

Parameters

- time_temperature_id The id of the time-temperature pair
- time The time [s]
- temperature The temperature [K]

class tc_python.propertymodel_sdk.CalculationContext(system:

tc_python.system.System, model_utils=None)

Bases: object

Represents the interface of the Property Model with the Thermo-Calc application and the rest of the TC-Python functionality.

Parameters

- **system** The system object of this calculation
- model_utils The model utils object

```
\texttt{get\_argument\_ids()} \rightarrow Set[str]
```

Returns a list with the arguments of the models. Including arguments created from dynamic parameters.

```
get_dependent_component() \rightarrow str
```

Obtains the dependent component from the UI

Note: The dependent component is that which has no composition specified explicitly, typically this is the major element of the material (such as Fe, Al, Ni, ...)

Returns The dependent component

$\texttt{get_mass_fractions()} \rightarrow Dict[str, float]$

Obtains the current composition from the UI as mass-fraction.

Note: In case of stepping over one or multiple axis, the returned data will represent the composition at the current step.

Returns The composition (key: component, value: content) [mass-fraction]

```
get_mass_percents() \rightarrow Dict[str, float]
```

Obtains the current composition from the UI in mass-percent.

Note: In case of stepping over one or multiple axis, the returned data will represent the composition at the current step.

Returns The composition (key: component, value: content) [mass-percent]

```
get_mole_fractions () \rightarrow Dict[str, float]
```

Obtains the current composition from the UI as mole-fraction.

Note: In case of stepping over one or multiple axis, the returned data will represent the composition at the current step.

Returns The composition (key: component, value: content) [mole-fraction]

```
get_mole_percents() \rightarrow Dict[str, float]
```

Obtains the current composition from the UI in mole-percent.

Note: In case of stepping over one or multiple axis, the returned data will represent the composition at the current step.

Returns The composition (key: component, value: content) [mole-percent]

```
get_temperature() \rightarrow float
```

Obtains the current temperature from the UI.

Returns The temperature [K]

get_ui_boolean_value (*component_id: str*) \rightarrow bool Obtains the value from the specified checkbox UI component.

Parameters component_id - Id of the checkbox

Returns The setting of the checkbox

- get_ui_condition_list (component_id: str) → tc_python.propertymodel_sdk.ConditionListEntry Used to get the selected condition from components of type UIConditionListComponent :param component id: Id of the list UI component :return: The selected condition
- get_ui_float_value (component_id: str) \rightarrow float Obtains the value from the specified UI component.

Parameters component_id - Id of the UI component

Returns The value

- $\texttt{get_ui_list_value}(\mathit{component_id: str}) \rightarrow \texttt{str}$
 - Obtains the selected entry from a UI component list. If a special element (such as ANY, NONE, ...) is selected, the corresponding locale-independent placeholder is provided.

Parameters component_id – Id of the list UI component

Returns The selected entry

- $get_ui_string_value(component_id: str) \rightarrow str$
 - Obtains the selected entry from a UI component text field.

Parameters component_id - Id of the string UI component

Returns The selected entry

 $get_ui_temperature_value(component_id: str) \rightarrow float$

Obtains the temperature from the specified temperature UI component.

Parameters component_id - Id of the temperature UI component

Returns The temperature [K], note that input unit of the UI is specified in the model panel. If required, the temperature is automatically converted to K.

set_result_cct_values (quantity_id: str, r: tc_python.propertymodel_sdk.CCTResultValues)
Sets the value of a previously defined result quantity (of type CCTResultValues) for further usage in
the Thermo-Calc application for plotting, etc.

Parameters

- **quantity_id** unique id of the result quantity
- **r** the CCTResultValues to be set
- set_result_quantity_value (quantity_id: str, value: float, parameter: str = ")

Sets the value of a previously defined result quantity for further usage in the Thermo-Calc application for plotting, etc.

Note: Any result quantity that remains unset is automatically set to *NaN*.

Parameters

- quantity_id Unique id of the result quantity
- parameter Use if result is parameterized. f.i. "per phase"
- **value** The value to be set

set_single_equilibrium_result (quantity_id: str, r: tc_python.single_equilibrium.SingleEquilibriumResult)

```
class tc_python.propertymodel_sdk.ConditionListEntry
    Bases: object
```

Used in combination with components of type UIConditionListComponent.

Contains the element, if the selected condition is a composition Contains the Console Mode syntax of the selected condition. Contains the unit of the selected condition

class tc_python.propertymodel_sdk.PropertyModel (_locale: str = 'en-US')
Bases: object

The abstract base class for all property models.

Note: Every Property Model needs to implement most of the abstract methods of this class. However, some abstract methods are optional and should only be implemented if required.

Note: If overwriting the constructor in a Property Model, the constructor of the implemented class must have the identical signature and should pass the parameters to this base class constructor.

Tip: It is possible to switch off **internal INFO-log messages coming from the calculation engine** by changing the log-level on the TC-Python log object like this: *logging.getLogger("tc_python").setLevel(logging.ERROR)*.

Parameters _locale – The locale to be used, this is an internal parameter and is of no meaning to the end-user

Variables logger – logger object that is connected to the Thermo-Calc UI (INFO- and WARN-ING-level will be printed as INFO, ERROR-level as ERROR), it can be accessed like this: self.logger.info("Some message")

abstract add_button_callback (component_id: str, ui_component_ids: Dict[str, int]) \rightarrow List[tc_python.propertymodel_sdk.UIComponent]

Implement this method if you have one or more UI components on which you called UIComponent. enable_add_button(), which adds a + button next to the component.

This method will be executed when you press any such + button.

This method is typically used to add more UI components dynamically and the method must return a list of the UI components to be added.

This method can optionally be implemented by a Property Model.

Parameters

- component_id The id of the UI component next to the pressed + button
- **ui_component_ids** dict with the id:s and the index of the current ui components of the model. Including arguments created in previous calls to add_button_callback.

Returns A list of UIComponent objects to be added

abstract after_evaluations()

Called by the Thermo-Calc application immediately after the last model evaluation (using the method *PropertyModel.evaluate_model()*). Use this method for any required cleanup.

This method can optionally be implemented by a Property Model.

abstract before_evaluations (*context:* tc_python.propertymodel_sdk.CalculationContext) Called by the Thermo-Calc application immediately before the first model evaluation (using the method *PropertyModel.evaluate_model()*). Use this method for any required preparations.

This method can optionally be implemented by a Property Model.

Parameters context – The calculation context

- abstract evaluate_model (*context*: tc_python.propertymodel_sdk.CalculationContext) Called by the Thermo-Calc application when the model should be actually calculated. This is the main
 - method of the Property Model that contains the actual calculation code.

This method needs to be implemented by all property models.

Parameters context – The calculation context, this provides access to the Thermo-Calc application and all other TC-Python modules

abstract get_license_key() \rightarrow str

Provides the license key of the model.

This method can optionally be implemented by a Property Model.

abstract provide_calculation_result_quantities()

List[tc_python.propertymodel_sdk.ResultQuantity]

Called by the Thermo-Calc application when the model should provide its result quantity objects.

This method needs to be implemented by all property models.

Returns Result quantity objects of the model (to be filled later with results in the method *PropertyModel.evaluate_model()*)

$\texttt{abstract provide_model_category()} \rightarrow List[str]$

Called by the Thermo-Calc application when the model should provide its category (shown in the Thermo-Calc model tree).

This method needs to be implemented by all property models.

Returns Category of the model, it may be present in several categories

<code>abstract provide_model_description()</code> \rightarrow str

Called by the Thermo-Calc application when the model should provide its detailed description.

This method needs to be implemented by all property models.

Returns Description text for the model

abstract provide_model_name() \rightarrow str

Called by the Thermo-Calc application when the model should provide its name (shown in the Thermo-Calc model tree).

This method needs to be implemented by all property models.

Returns Name of the model

abstract provide_model_parameters() \rightarrow Dict[str, float]

Called by the Thermo-Calc application when the model should provide all model parameters and their current values.

This method can optionally be implemented by a Property Model.

Note: These are internal variables of the Property Model that are intended to be modified from the outside. Typically this is used to adjust their values in a optimizer during the development of the model.

Returns The model parameter ids and their current values [unit according to the parameter meaning]

abstract provide_ui_panel_components() \rightarrow List[*tc_python.propertymodel_sdk.UIComponent*] Called by the Thermo-Calc application when the model should provide its UI components for the model panel to be plotted. This happens also whenever a model gets checked in the model tree.

This method needs to be implemented by all property models.

Returns Model UI panel components in the order to be presented in the model panel

 $\texttt{abstract remove_button_callback} (\textit{component_id: str, ui_component_ids: Dict[str, int]}) \rightarrow \texttt{abstract remove_button_callback} (\texttt{component_id: str, ui_component_ids: Dict[str, int]}) \rightarrow \texttt{abstract remove_button_callback} (\texttt{abstract remove_button_callbac$

List[str]

Implement this method if you have one or more UI components on which you called UIComponent. enable_remove_button(), which adds a - button next to the component.

This method will be executed when you press any such - button.

This method is typically used to remove UI components dynamically and the method must return a list of the ids of the components that are going to be removed.

This method can optionally be implemented by a Property Model.

Parameters

- component_id the id of the UI component next to the pressed button
- **ui_component_ids** list with the id:s and the index of the current ui components of the model. Including arguments created in previous calls to add_button_callback.

Returns a list of UI component ids that are required to be removed

abstract set_model_parameter(model_parameter_id: str, value: float)

Called by the Thermo-Calc application when a model parameter should be reset.

This method can optionally be implemented by a Property Model.

Note: These are internal variables of the Property Model that are intended to be modified from the outside. Typically this is used to adjust their values in a optimizer during the development of the model.

Parameters

- model_parameter_id The parameter id
- **value** The value [unit according to the parameter meaning]

```
class tc_python.propertymodel_sdk.ResultQuantity(quantity_id: str, descrip-
tion: str, quantity_type:
tc_python.propertymodel_sdk.ResultQuantityType)
```

Bases: object

Defines a calculation result quantity of a Property Model that is identified by a unique id.

Parameters

- quantity_id Unique id of the quantity
- description Description of the quantity (shown in the Thermo-Calc UI)
- **quantity_type** Type of the quantity (defines the unit)

```
get_description() \rightarrow str
```

Obtains the description of the quantity.

Returns Description of the quantity

$\texttt{get_id}() \rightarrow str$

Obtains the id of the quantity.

Returns Unique id of the quantity

```
get\_type() \rightarrow tc\_python.propertymodel\_sdk.ResultQuantityType
Obtains the type of quantity.
```

Returns Type of the quantity

```
class tc_python.propertymodel_sdk.ResultQuantityType(value)
    Bases: enum.Enum
```

Defining the type of a result quantity.

```
CCT_QUANTITY = 5
A cct quantity
```

```
ENERGY_QUANTITY = 2
An energy quantity
```

GENERAL_QUANTITY = 0 A general quantity

LENGTH_QUANTITY = 7 A length in quantity

```
SINGLE_EQUILIBRIUM_QUANTITY = 6
A cct quantity
```

```
STRENGTH_QUANTITY = 8
A strength quantity
```

```
SURFACE_ENERGY_QUANTITY = 3
A surface energy quantity
```

```
TEMPERATURE_QUANTITY = 1
A temperature quantity
```

TIME_QUANTITY = 4 A time quantity

class tc_python.propertymodel_sdk.SpecialListMarkers Bases: object

Placeholders for special list elements that are locale-dependent. They will be provided by UI list components if a special marker has been selected.

```
ANY_LIST_MARKER = 'ANY'
Marker that represents "Any"
```

```
NONE_LIST_MARKER = 'NONE'
Marker that represents "None"
```

Bases: tc_python.propertymodel_sdk.UIComponent

Checkbox UI component of the model panel.

Parameters

- component_id Unique id of the component
- name Name of the component, will be presented in the model panel

- description Additional description of the component
- **setting** Initial setting of the checkbox

connect_component_visibility (dependent_component_id: str)

Connects the visibility of any other UI component of the model panel to the value of this boolean component.

Parameters dependent_component_id – Id of the UI element to be dependent on this boolean component

enable_add_button()

Adds a + button to the right of the UI component.

Returns This UI component

enable_remove_button()

Adds a - button to the right of the UI component.

Returns This UI component

get_dependent_components() \rightarrow List[str]

Obtains a list containing all UI elements currently connected regarding their visibility.

Returns A list with the component id of all UI elements currently connected

get_setting() \rightarrow bool

Obtains the setting of the checkbox.

Returns The setting of the checkbox

remove_component_visibility (dependent_component_id: str)

Removes the visibility connection to a UI component that has been previously connected.

Parameters dependent_component_id - Id of the previously connection UI element

set_index (index: int = - 1)

Sets the position in the graphical user interface.

Parameters index – The position

Returns This UI component

class tc_python.propertymodel_sdk.UIComponent(component_id: str, name: str, description: str)

Bases: object

Abstract Base class for all UI components of the model panel.

Never make an instance of UIComponent, always use the sub-classes. For instance UIStringComponent.

Parameters

- component_id Unique id of the component
- **name** Name of the component, will be presented in the model panel
- description Additional description of the component

get_description() \rightarrow str

Obtains the additional description of the component.

Returns Additional description of the component

 $get_id() \rightarrow str$

Obtains the unique id of the component.

Returns Unique id of the component

get_name () \rightarrow str Obtains the name of the component.

Returns Name of the component, will be presented in the model panel

class tc_python.propertymodel_sdk.**UIConditionListComponent**(*component_id: str*,

name: str, description: str)

Bases: tc_python.propertymodel_sdk.UIComponent

System condition list UI component of the model panel.

Parameters

- component_id Unique id of the component
- name Name of the component, will be presented in the model panel
- **description** Additional description of the component

class tc_python.propertymodel_sdk.UIFloatComponent (component_id: str, name: str, de-

scription: str, value: float)

Bases: tc_python.propertymodel_sdk.UIComponent

General real value text field UI component of the model panel.

Parameters

- component_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- **value** Initial setting of the text field

enable_add_button()

Adds a + button to the right of the UI component.

Returns This UI component

enable_remove_button () Adds a - button to the right of the UI component.

Returns this UI component

 $\begin{array}{l} \textbf{get_value} () \ \rightarrow float\\ Obtains the setting of the text field. \end{array}$

Returns The setting of the text field

set_index (index: int = - 1)
Sets the position in the graphical user interface.

Parameters index – The position

Returns This UI component

<pre>class tc_python.propertymodel_sdk.UIGeneralListComponent</pre>	. (compone	ent_id:	str,
	name:	str,	descrip-
	tion:	str,	content:
	List[Tup	le[str,	<i>str]]</i> ,
	selected_entry: $str = "$)		
Bases: ta puthan proportymodal adk UICampanant			

Bases: tc_python.propertymodel_sdk.UIComponent

General list UI component of the model panel that can contain any strings.

Parameters

- component_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- content Entries of the list, they need to contain a locale-independent id and a localized content string, for example: [("ENTRY_1_ID", "entry 1"), (ENTRY_2_ID", "entry 2")]
- **selected_entry** Entry to be initially selected. If omitted, by default the first element is selected.

connect_component_visibility(dependent_component_id: str, selected_item_to_set_visible:

str)

Connects the visibility of any other UI component of the model panel to the selection of a certain entry of the list.

Parameters

- dependent_component_id Id of the UI element to be dependent on the chosen element
- **selected_item_to_set_visible** Entry (locale independent id) of the list to be chosen to set the dependent component visible

enable_add_button()

Adds a + button to the right of the UI component.

Returns This UI component

enable_remove_button()

Adds a - button to the right of the UI component.

Returns This UI component

 $\texttt{get_content}() \rightarrow List[Tuple[str, str]]$

Obtains the entries of the list.

Returns Entries of the list, they need to contain a locale-independent id and a localized content string, for example: [("ENTRY_1_ID", "entry 1"), (ENTRY_2_ID", "entry 2")]

get_dependent_components() \rightarrow Dict[str, List[str]]

Obtains a dictionary containing all UI elements currently connected regarding their visibility.

Returns All UI elements currently connected (key: dependent component id, value: required list entries to set it visible)

$\texttt{get_selected_entry()} \rightarrow str$

Obtains the initially selected entry.

Returns Initially selected entry. If empty, the first element is selected.

```
remove_component_visibility(dependent_component_id: str)
```

Removes the visibility connection to a UI component that has been previously connected.

Parameters dependent_component_id - Id of the previously connection UI element

set_index (index: int = - 1)

Sets the position in the graphical user interface.

Parameters index – The position

Returns This UI component

Bases: tc_python.propertymodel_sdk.UIComponent

Phase list UI component of the model panel.

Parameters

- component_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- default_phase Default phase, if omitted no default phase is chosen and only initially the first element of the list is selected. If an ANY-marker is added, this is chosen as the default element.
- **any_marker_setting** Defines if an entry "ANY PHASE" should be added to the phase list, if set to true this overrides any default phase setting

enable_add_button()

Adds a + button to the right of the UI component.

Returns This UI component

enable_remove_button()

Adds a - button to the right of the UI component.

Returns This UI component

get_any_marker_setting() \rightarrow bool

Obtains the setting if any entry "ANY PHASE" is added to the phase list.

Returns If an entry "ANY PHASE" is added to the phase list, if set to true this overrides any default phase setting

$\texttt{get_default_phase()} \rightarrow str$

Obtains the default phase.

Returns Default phase, if omitted no default phase is chosen and only initially the first element of the list is selected. If an ANY-marker is added, this is chosen as the default element.

set_index (index: int = - 1)

Sets the position in the graphical user interface.

Parameters index – The position

Returns This UI component

Bases: tc_python.propertymodel_sdk.UIComponent

General text field UI component of the model panel.

Parameters

- component_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component

- string Initial setting of the text field
- enable_add_button() Adds a + button to the right of the UI component.

Returns This UI component

enable remove button() Adds a - button to the right of the UI component.

Returns This UI component

get_value() \rightarrow str Obtains the setting of the text field.

Returns The setting of the text field

set_index (index: int = - 1) Sets the position in the graphical user interface.

Parameters index – The position

Returns This UI component

```
class tc_python.propertymodel_sdk.UITemperatureComponent (component_id:
                                                                                      str.
                                                                    name: str, description:
                                                                    str, temp: float)
```

Bases: tc_python.propertymodel_sdk.UIComponent

Temperature value text field UI component of the model panel.

Parameters

- component_id Unique id of the component
- **name** Name of the component, will be presented in the model panel
- description Additional description of the component
- temp Initial temperature to be set in the text field (unit defined by the user in the Thermo-Calc system)

enable_add_button()

Adds a + button to the right of the UI component.

Returns This UI component

enable remove button()

Adds a - button to the right of the UI component.

Returns This UI component

get_temp() \rightarrow float

Obtains the temperature set in the text field.

Returns The temperature to be set in the text field (unit defined by the user in the Thermo-Calc system)

set_index (index: int = - 1)

Sets the position in the graphical user interface.

Parameters index – The position

Returns This UI component

tc_python.propertymodel_sdk.create_boolean_ui_component (component_id: str, name:

str, description: str,

```
initial setting: bool) \rightarrow
```

tc_python.propertymodel_sdk.UIBooleanComponer

Creates a UI checkbox component for a boolean value. The value of that component can later be accessed during the model evaluation.

Parameters

- component_id Unique id of the component
- **name** Name of the component, will be presented in the model panel
- description Additional description of the component
- **initial_setting** Initial setting of the checkbox

Returns The created component

tc_python.propertymodel_sdk.create_condition_list_ui_component(component_id:

str, name: str, description: str) → tc python.propertymodel sdk.UICondition

Creates a UI list component for all conditions defined in the system. The value of that component can later be accessed during the model evaluation.

Parameters

- component_id Unique id of the component
- **name** Name of the component, will be presented in the model panel
- description Additional description of the component

Returns The created component

<pre>tc_python.propertymodel_sdk.create_energy_quantity(quantity_id:</pre>	str,	de-
scription:	str)	\rightarrow
tc_python.propert		

Creates a UI energy result quantity (in J). When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

Parameters

- quantity_id Unique id of the result quantity
- **description** Additional description of the result quantity

Returns The created result quantity

```
tc_python.propertymodel_sdk.create_float_ui_component(component_id: str,
name: str, description:
str, value: float) →
tc python.propertymodel sdk.UIFloatComponent
```

Creates a UI text field component for a real number. The value of that component can later be accessed during the model evaluation.

Parameters

- component_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component

• value - Initial setting of the text field

Returns The created component

<pre>tc_python.propertymodel_sdk.create_general_quantity(quantity_id:</pre>	str,	de-
scription:	str)	\rightarrow

tc_python.propertymodel_sdk.ResultQuantity

Creates a general result quantity that can contain any type of result (without a unit). When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

Parameters

- quantity_id Unique id of the result quantity
- description Additional description of the result quantity

Returns The created result quantity

tc_python.propertymodel_sdk.ResultQuantity

Creates a length result quantity. When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

Parameters

- **quantity_id** Unique id of the result quantity
- description Additional description of the result quantity

Returns The created result quantity

tc_python.propertymodel_sdk.create_list_ui_component(component_id: str, name:

str, description: str, entry_list: List[Tuple[str, str]], selected_entry: str = ") \rightarrow

tc_python.propertymodel_sdk.UIGeneralListComponen

Creates a UI list component for string entries. The value of that component can later be accessed during the model evaluation.

Parameters

- component_id Unique id of the component
- **name** Name of the component, will be presented in the model panel
- description Additional description of the component
- **entry_list** Entries of the list, they need to contain a locale-independent id and a localized content string, for example: [("ENTRY_1_ID", "entry 1"), (ENTRY_2_ID", "entry 2")]
- **selected_entry** Entry to be initially selected. If omitted, by default the first element is selected.

Returns The created component

<pre>tc_python.propertymodel_sdk.create_phase_list_ui_component</pre>	(component_id: str,
	name: str, de-
	scription: str, de-
	fault_phase: str
	= ", any_marker:
	$bool = False) \rightarrow$
	tc_python.propertymodel_sdk.UIPhaseListCom
	a anti-at a default where that is

Creates a UI list component for all phases defined in the system. It is possible to select a default phase that is

supposed to be the **expected phase selection** for that list. The value of that component can later be accessed during the model evaluation.

A **default** phase is the phase that is initially selected and re-selected as soon as a currently selected phase is removed. If the default phase is not available, a "NONE"-marker will be created and used instead of the default phase. A typical use case for the default phase setting is a phase list that expects to contain the LIQUID-phase of a system.

Parameters

- component_id Unique id of the component
- **name** Name of the component, will be presented in the model panel
- description Additional description of the component
- **default_phase** Default phase, if omitted no default phase is chosen and only initially the first element of the list is selected. If an ANY-marker is added, this is chosen as the default element.
- **any_marker** Defines if an entry "ANY PHASE" should be added to the phase list, if set to true this overrides any default phase setting

Returns The created component

Creates a strength result quantity. When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

Parameters

- quantity_id Unique id of the result quantity
- description Additional description of the result quantity

Returns The created result quantity

```
tc_python.propertymodel_sdk.create_string_ui_component(component_id: str,
```

name: str, description:

str, string: str) \rightarrow *tc_python.propertymodel_sdk.UIStringComponent*

Creates a UI text field component. The value of that component can later be accessed during the model evaluation.

Parameters

- **component_id** Unique id of the component
- **name** Name of the component, will be presented in the model panel
- description Additional description of the component
- string Initial setting of the text field

Returns The created component

tc_python.propertymodel_sdk.create_surface_energy_quantity(quantity_id: str,

description: str) \rightarrow

tc_python.propertymodel_sdk.ResultQuantity

Creates an energy result quantity (in J). When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

Parameters

- quantity_id Unique id of the result quantity
- description Additional description of the result quantity

Returns The created result quantity

```
tc_python.propertymodel_sdk.create_temperature_quantity(quantity_id: str, de-
```

scription: str)

 \rightarrow

tc_python.propertymodel_sdk.ResultQuantity Creates a temperature result quantity (in K). When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

Parameters

- quantity_id Unique id of the result quantity
- description Additional description of the result quantity

Returns The created result quantity

tc_python.propertymodel_sdk.create_temperature_ui_component (component_id:

str, name: str, description: str, initial temp: float) \rightarrow

tc python.propertymodel sdk.UITemperature

Creates a UI text field component for a temperature value. The value of that component can later be accessed during the model evaluation.

Parameters

- component_id Unique id of the component
- **name** Name of the component, will be presented in the model panel
- description Additional description of the component
- **initial_temp** Initial temperature to be set in the text field. (The unit of initial_temp is Kelvin. The value in the text field will be automatically converted using the unit chosen by the user.)

Returns The created component

```
tc_python.propertymodel_sdk.create_time_quantity(quantity_id: str, description: str) \rightarrow
```

tc_python.propertymodel_sdk.ResultQuantity

Creates a time result quantity (in s). When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

Parameters

- quantity_id Unique id of the result quantity
- **description** Additional description of the result quantity

Returns The created result quantity

5.8 Module "exceptions"

exception tc_python.exceptions.**APIServerException** Bases: tc_python.exceptions.GeneralException

An exception that occurred during the communication with the API-server. It is normally not related to an error in the user program.

exception tc_python.exceptions.CalculationException
Bases: tc_python.exceptions.TCException

An exception that occurred during a calculation.

exception tc_python.exceptions.ComponentNotExistingException
Bases: tc_python.exceptions.GeneralException

The selected component is not existing.

exception tc_python.exceptions.DatabaseException
Bases: tc_python.exceptions.CalculationException

Error loading a thermodynamic or kinetic database, typically due to a misspelled database name or a database missing in the system.

exception tc_python.exceptions.DegreesOfFreedomNotZeroException
Bases: tc_python.exceptions.CalculationException

The degrees of freedom in the system are not zero, i.e. not all required conditions have been defined. Please check the conditions given in the exception message.

exception tc_python.exceptions.**EquilibriumException** Bases: tc_python.exceptions.CalculationException

An equilibrium calculation has failed, this might happen due to inappropriate conditions or a very difficult problem that can not be solved.

exception tc_python.exceptions.GeneralCalculationException
Bases: tc_python.exceptions.CalculationException

General error occurring while a calculation is performed.

exception tc_python.exceptions.GeneralException
Bases: tc_python.exceptions.TCException

A general exception that might occur in different situations.

exception tc_python.exceptions.InvalidCalculationConfigurationException
Bases: tc_python.exceptions.CalculationException

Thrown when errors are detected in the configuration of the calculation.

exception tc_python.exceptions.InvalidCalculationStateException
Bases: tc_python.exceptions.CalculationException

Trying to access an invalid calculation object that was invalidated by calling invalidate on it.

exception tc_python.exceptions.InvalidNumberOfResultGroupsException
 Bases: tc_python.exceptions.ResultException

A calculation result contains several result groups, which is not supported for the used method.

exception tc_python.exceptions.InvalidResultConfigurationException
Bases: tc_python.exceptions.ResultException

A calculation result configuration is invalid.

```
exception tc_python.exceptions.InvalidResultStateException
    Bases: tc_python.exceptions.CalculationException
```

Trying to access an invalid result (for example a SingleEquilibriumTempResult object that got already invalidated by condition changes or a result that was invalidated by calling *invalidate* on it).

exception tc_python.exceptions.LicenseException
Bases: tc_python.exceptions.GeneralException

No valid license for the API or any Thermo-Calc product used by it found.

exception tc_python.exceptions.NoDataForPhaseException
 Bases: tc_python.exceptions.ResultException

There is no result data available for a selected phase.

exception tc_python.exceptions.NotAllowedOperationException
Bases: tc_python.exceptions.CalculationException

The called method or operation is not allowed in the current mode of operation (i.e. debug or production mode). *Production mode* means that the Property Model is only present as an *.*py.encrypted*-file, while in *debug mode* it is available as *.*py*-file. Certain methods for obtaining internal model parameters are not available for encrypted models.

exception tc_python.exceptions.PhaseNotExistingException
Bases: tc_python.exceptions.GeneralException

The selected phase is not existing, so no data can be provided for it.

exception tc_python.exceptions.ResultException
Bases: tc_python.exceptions.TCException

An exception that occurred during the configuration of a calculation result.

exception tc_python.exceptions.SyntaxException
Bases: tc_python.exceptions.CalculationException

Syntax error in a Console Mode expression.

exception tc_python.exceptions.TCException
 Bases: Exception

The root exception of TC-Python.

exception tc_python.exceptions.UnrecoverableCalculationException
Bases: tc_python.exceptions.CalculationException

The calculation reached a state where no further actions are possible, this happens most often due to a FORTRAN- hard crash in the API server backend.

Note: It is possible to catch that exception outside of the *with*-clause context and to continue by setting up a new context (i.e. by a new *with TCPython() as session*).

tc_python.exceptions.handle_exception(e)

5.9 Module "abstract_base"

```
class tc_python.abstract_base.AbstractCalculation(calculator)
    Bases: object
```

Abstract base class for calculations.

```
get_configuration_as_string() \rightarrow str
```

Returns detailed information about the current state of the calculation object.

Warning: The structure of the calculator objects is an implementation detail and might change between releases without notice. **Therefore do not rely on the internal object structure**.

$\texttt{get_system_data()} \rightarrow \textit{tc_python.abstract_base.SystemData}$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with_system_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

Returns The system data

invalidate()

Invalidates the object and frees the disk space used by it. *This is only required if the disk space occupied by the object needs to be released during the calculation*. No data can be retrieved from the object afterwards.

with_system_modifications (system_modifications: tc_python.abstract_base.SystemModifications)
 Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

Note: This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a * .tdb-file.

Parameters system_modifications - The system modification to be performed

Returns

```
class tc_python.abstract_base.AbstractResult(result)
```

Bases: object

Abstract base class for results. This can be used to query for specific values .

invalidate()

Invalidates the object and frees the disk space used by it. *This is only required if the disk space occupied by the object needs to be released during the calculation*. No data can be retrieved from the object afterwards.

class tc_python.abstract_base.PhaseParameter(parameter_name: Union[str, object])
 Bases: object

Database phase parameter expression used by *SystemModifications.set()*.

Parameters parameter_name – The phase parameter name

```
get_intervals() \rightarrow List[tc_python.abstract_base.TemperatureInterval]
Returns the list of all defined intervals.
```

Returns The defined temperature intervals

get_lower_temperature_limit () \rightarrow float Returns the lower temperature limit.

Returns The lower temperature limit in K

get_name() \rightarrow str

Returns the name of the phase parameter.

Returns The name of the phase parameter.

remove_all_intervals()

Removes all previously defined temperature intervals.

Returns This *PhaseParameter* object

remove_interval_with_upper_limit (upper_temperature_limit: float)

Removes a previously defined temperature interval with matching upper temperature limit.

If no such interval exists, an exception is thrown.

Returns This *PhaseParameter* object

Adds/overwrites a parameter expression for a temperature interval.

Default value of the upper limit of the interval: 6000 K

Note: The lower temperature limit is either defined by the lower temperature limit given with *PhaseParameter.set_lower_temperature_limit()* or by the upper temperature limit of the adjacent interval.

Note: If there is an existing interval with exactly the same *upper_temperature_limit*, that interval is overwritten, otherwise the interval is added.

Parameters

- **parameter_expression** The parameter expression, example: +V34*T*LN(T)+V35*T**2+V36*T**(-1)+V37*T**3")
- **upper_temperature_limit** The upper temperature limit for which the expression should be used

Returns This *PhaseParameter* object

set_interval (*interval*: tc_python.abstract_base.TemperatureInterval) Adds/overwrites a temperature interval.

Note: The lower temperature limit is either defined by the lower temperature limit given with *PhaseParameter.set_lower_temperature_limit()* or by the upper temperature limit of the adjacent interval.

Note: If there is an existing interval with exactly the same *upper_temperature_limit*, that interval is overwritten, otherwise the interval is added.

Returns This *PhaseParameter* object

set_lower_temperature_limit (*lower_temperature_limit: float = 298.15*) Sets the lower temperature limit of the phase parameter.

Default: 298.15 K

Parameters lower_temperature_limit – The lower temperature limit in K

Returns This *PhaseParameter* object

class tc_python.abstract_base.**SystemData**(*system_data*) Bases: object

Provides information about the parameters and functions of a user database. The obtained objects can be used to modify the database using with_system_modifications() of all calculators.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as *.tdb-file.

get_phase_parameter (*parameter: str*) \rightarrow *tc_python.abstract_base.PhaseParameter* Returns a phase parameter.

Example:

system_data.get_phase_parameter('G(HCP_A3,FE:VA;0)')

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as a *.tdb-file.

Note: For details about the syntax search the Thermo-Calc help for *GES* (the name for the Gibbs Energy System module in Console Mode).

Parameters parameter – The name of the phase parameter (for example: "G(LIQUID,FE;0)")

Returns The phase parameter

$\texttt{get_phase_parameter_names()} \rightarrow List[str]$

Returns all phase parameters present in the current system.

Returns The list of phase parameters

get_system_function (f: str) \rightarrow tc_python.abstract_base.SystemFunction Returns a system function.

Note: The parameter 'f' was previously called 'function' but was renamed.

Example:

system_data.get_system_function('GHSERCR')

Note: Functions can only be read from unencrypted (i.e. user) databases loaded as a *.tdb-file.

Note: For details about the syntax search the Thermo-Calc help for *GES* (the name for the Gibbs Energy System module in Console Mode).

Parameters f – The name of the system function (for example: "GHSERCR")

Returns The system function

```
get_system_function_names() \rightarrow List[str]
```

Returns all system functions present in the current system.

Returns The list of system functions

```
class tc_python.abstract_base.SystemFunction(function_name: Union[str, object])
Bases: object
```

Database function expression used by SystemModifications.set().

Parameters function_name – The function name

get_intervals () \rightarrow List[*tc_python.abstract_base.TemperatureInterval*] Returns the list of all defined intervals.

Returns The defined temperature intervals

get_lower_temperature_limit () \rightarrow float Returns the lower temperature limit.

Returns The lower temperature limit in K

get_name() \rightarrow str

Returns the name of the system function.

Returns The name of the system function

remove_all_intervals()

Removes all previously defined temperature intervals.

Returns This SystemFunction object

remove_interval_with_upper_limit (upper_temperature_limit: float)
Removes a previously defined temperature interval with matching upper temperature limit.

If no such interval exists, an exception is thrown.

Returns This SystemFunction object

set_expression_with_upper_limit (function_expression: str, upper_temperature_limit: float = 6000.0)

Adds/overwrites a function expression for a temperature interval.

Default value of the upper limit of the interval: 6000 K

Note: The lower temperature limit is either defined by the lower temperature limit given with *SystemFunction.set_lower_temperature_limit()* or by the upper temperature limit of the adjacent interval.

Note: If there is an existing interval with exactly the same *upper_temperature_limit*, that interval is overwritten, otherwise the interval is added.

Parameters

- function_expression The function expression, example: +V34*T*LN(T)+V35*T**2+V36*T**(-1)+V37*T**3")
- **upper_temperature_limit** The upper temperature limit for which the expression should be used

Returns This SystemFunction object

set_interval (*interval*: tc_python.abstract_base.TemperatureInterval) Adds/overwrites a temperature interval.

Note: The lower temperature limit is either defined by the lower temperature limit given with *SystemFunction.set_lower_temperature_limit()* or by the upper temperature limit of the adjacent interval.

Note: If there is an existing interval with exactly the same *upper_temperature_limit*, that interval is overwritten, otherwise the interval is added.

Returns This SystemFunction object

set_lower_temperature_limit (lower_temperature_limit: float = 298.15)
Sets the lower temperature limit of the system function.

Default: 298.15 K

Parameters lower_temperature_limit - The lower limit in K

Returns This SystemFunction object

class tc_python.abstract_base.SystemModifications
 Bases: object

Functionality to modify a user database during a calculation by changing phase parameters and system functions.

The actual changes are only applied by using tc_python.abstract_base.AbstractCalculation. with_system_modifications() on a calculator object.

run_ges_command(ges_command: str)

Sends a GES-command. This is actually applied when running `with_system_modifications` on a calculator object.

Example: *run_ges_command*("*AM-PH-DE FCC_A1 C_S 2 Fe:C*") for adding a second composition set to the FCC_A1 phase with *Fe* as major constituent on first sublattice and *C* as major constituent on second sublattice.

Note: For details about the syntax search the Thermo-Calc help for *GES* (the name for the Gibbs Energy System module in Console Mode).

Note: It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

Warning: As this method runs raw GES-commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten parenthesis, ...).

Parameters ges_command – The GES-command (for example: "AM-PH-DE FCC_A1 C_S 2 Fe:C")

Returns This SystemModifications object

set (parameter_or_function:

Union[tc_python.abstract_base.PhaseParameter,

per temperature limit: float)

tc_python.abstract_base.SystemFunctionJ) Overwrites or creates a phase parameter or system function.

Example: system_modifications.set(PhaseParameter('G(LIQUID,FE;0)').set_expression_with_upper_limit('+1.2*GFELIQ

Example: system_modifications.set(SystemFunction("DGDEF").set_expression_with_upper_limit(+10.0-R*T', 1000).set_expression_with_upper_limit(+20.0-R*T', 3000))

Note: The old parameter/function is overwritten and any temperature intervals not defined are lost.

Note: Please consult the Thermo-Calc GES-system documentation for details about the syntax.

Returns This SystemModifications object

class tc_python.abstract_base.TemperatureInterval(expression: Union[str, object], up-

Bases: object

Temperature interval expression used by the classes *SystemFunction* and *PhaseParameter*.

Parameters

- **expression** The temperature function expressed in Thermo-Calc database syntax.
- upper_temperature_limit The upper temperature limit in K

get_expression() \rightarrow str

Returns the function expression of this temperature interval.

Returns The temperature function expression

get_upper_temperature_limit () \rightarrow float Returns the upper limit of this temperature interval.

Returns The upper temperature limit in K

set_expression(expression: str)

Sets the function expression of this temperature interval.

Parameters expression – The temperature function expression

set_upper_temperature_limit (upper_temperature_limit: float)
Sets the upper limit of this temperature interval.

Parameters upper_temperature_limit – The upper temperature limit in K

CHAPTER

TROUBLESHOOTING

This section provides an FAQ for common problems that occur when using TC-Python.

6.1 Diagnostics script

If you have problems running TC-Python, run the diagnostics script below.

On Linux you can alternatively download the script directly into your current working directory by:

curl -0 https://www2.thermocalc.com/downloads/support/diagnostics-py/tc-python-

```
→diagnostic-script-2022b.py
.....
Run this script when troubleshooting TC-Python
It is important to run this script EXACTLY the same way as you run your TC-Python.
⇔script
(In the same IDE, same project, same Python environment, same Jupyter notebook e.t.c)
.....
version = '2022b'
print('Testing TC-Python version: ' + version)
print ('Please make sure that the variable "version" above, matches the release that_
→you want to test, if not change it and re-run this script.')
# below this line, nothing needs to be manually updated.
import sys
print('')
print('Python version: (should be at least 3.5 and can NOT be older than 3.0)')
print(str(sys.version_info[0]) + '.' + str(sys.version_info[1]))
if sys.version_info[0] < 3 or sys.version_info[1] < 5:</pre>
   print('Wrong version of Python !!!!!')
print('')
print ('Python executable path: (gives a hint about the used virtual / conda_
\hookrightarrowenvironment, in case of Anaconda the corresponding \n'
      'environment name can be found by running `conda env list` on the Anaconda.
→command prompt, '
      'TC-Python must be installed into \nEACH separate environment used!)')
print(sys.executable)
```

(continues on next page)

```
(continued from previous page)
```

```
import os
print('')
print ('Thermo-Calc ' + version + ' installation directory: (must be a valid path to a_

complete installation of ' + version + ')')

tc_env_variable = 'TC' + version[2:].upper() + '_HOME'
try:
   print(os.environ[tc_env_variable])
except:
   print('No Thermo-calc environment variable for ' + version + ' was found. (' + tc_
print('')
print('Url of license server: (if license server is NO-NET, you need a local license.
→file)')
try:
   print(os.environ['LSHOST'])
except:
   print ('No Thermo-calc license server url was found. (LSHOST)')
print('')
print ('Path to local license file: (only necessary if not using license server)')
try:
   print(os.environ['LSERVRC'])
except:
   print ('No path to local license file was found. (LSERVRC)')
import tc_python
numerical_version = version[:-1]
if version[-1] == 'a':
   numerical_version += '.1.*'
elif version[-1] == 'b':
   numerical_version += '.2.*'
print('')
print('TC-Python version: (needs to be ' + numerical_version + ')')
print(tc_python.__version__)
with tc_python.TCPython() as session:
   print('')
   print('Lists the databases: (should be a complete list of the installed databases_
↔that you have license for or do not require license)')
   print(session.get_databases())
```

6.2 "No module named tc_python" error on first usage

This problem occurs because your used Python interpreter cannot find the TC-Python package. We expect that you have installed the TC-Python package in your **Python system interpreter** following the instructions in the *Installation Guide*.

Normally the error message "*No module named tc_python*" is caused by unintentionally configuring a PyCharm project to use a so-called **Virtual Environment**. This happens unfortunately by default when creating a new PyCharm project with not changing the default settings.

Note: A Virtual Environment is basically a separate and completely independent copy of the system-wide Python interpreter. It does not contain any packages.

On Windows systems we recommend to use the Anaconda Python Distribution as Python interpreter. However, the instructions given here are valid for any operating system and distribution.

Since TC-Python 2018b we do recommend to **not use Virtual Environments** unless there is a reasonable use case for that.

There are two possible solutions to fix the problem:

1. The quick fix for your problem is to run

pip install <path to the TC-Python folder>/TC_Python-<version>-py3-none-any.whl

within the *Terminal window* of the opened PyCharm project. This *Terminal window* automatically runs within the *Virtual Environment* configured for the project (if any). You can see the name of the *Virtual Environment* at the beginning of each command prompt line (here it is called *venv*):

```
Microsoft Windows [Version 10.0.16299.431]
(c) 2017 Microsoft Corporation. All rights reserved.
(venv) C:\Users\User\Documents\>
```

The command will consequently install TC-Python also within that Virtual Environment automatically. The Terminal window can be found at the bottom of the IDE. Note that it might be necessary to enable these buttons first by selecting the menu entry **View** \rightarrow **Tool Buttons**.

- 2. The better fix is to change your project **to use the system interpreter**. This is described in detail in the section *Fixing potential issues with the environment* in Step 5 of the *Installation Guide*.
 - It is recommendable to use that approach also for all your future projects.

Both fixes will only change the configuration of the opened project. Further useful information can be found in the section *Python Virtual Environments*.

6.3 "pip install" fails with "Failed to establish a new network connection" or similar

If *pip install* fails with a network related error (might also be "*socket not available*", "*retrying after connection broken*", ...) it is often due to the computer being behind a proxy-server, this is common in large organizations. Of course also the network connection might be broken.

TC-Python has dependencies to a few other packages:

- *py4j*
- jproperties
- six (transient dependency of jproperties)
- 1. The recommended approach is to simply use *pip*. It will resolve the dependencies automatically by downloading them from the *PyPI*-repository server (https://pypi.org). If your computer is located behind a proxy-server, the connection to the repository will fail. In that case it is necessary to configure *pip* with the detailed configuration of the proxy server:

pip install -proxy user:password@proxy_ip:port py4j jproperties

2. Another alternative is to manually download the latest *.whl-file of each dependency from the repository server (https://pypi.org -> Search projects) and to install it manually using:

```
pip install py4j-#.#.#-py2.py3-none-any.whl
...
```

The actual actual version number needs to be inserted into the file name. The downside of this approach is that updates to that package have to be fully manual also in the future. Additionally it is also necessary to install all transient dependencies in that way.

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t

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