# **Cortix Documentation**

Release 0.1.0

Valmor F. de Almeida

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**CHAPTER** 

ONE

**SRC** 

# 1.1 application

Application class for Cortix.

Cortix: a program for system-level modules coupling, execution, and analysis.

Bases: object

An Application is a singleton class composed of Module objects, and Network objects; the latter involve Module objects in various combinations. Each combination is assigned to a Network object.

```
get_module (name)
```

Returns a module with a given name. None if the name doesn't exist.

```
get network(name)
```

Returns a network with a given name. None if the name doesn't exist.

### modules

list(str) - List of names of Cortix module objects

### networks

list(str) – List of names of network objects

# 1.2 cortix\_driver\_template

Cortix driver for guest modules. Module developers must implement the public methods in this driver. Ideally, this implementation should be minimal. Developers should use this class to wrap their module (MyModule) implemented in a file named my\_module.py. This file will be placed inside the developer's module directory which is pointed to in the Cortix config.xml file.

Bases: object

Cortix driver for guest modules.

```
call_ports (cortix_time=0.0)
Call all ports at cortix_time

execute (cortix_time=0.0, timeStep=0.0)
Evolve system from cortix_time to cortix_time + timeStep
```

# 1.3 cortix main

The Cortix class definition.

Cortix: a program for system-level modules coupling, execution, and analysis.

```
class cortix_main.Cortix(name=None, config_file='cortix-config.xml')
    Bases: object
```

The main Cortix class definition. This class encapsulates the concepts of simulations, tasks, and modules, for a given application providing the user with an interface to the simulations.

```
run_simulations (task_name=None)
```

This method runs every simulation defined by the Cortix object. At the moment this is done one simulation at a time.

# 1.4 cortix module template

Simple MyModule module template for developers.

### 1.5 launcher

Launcher functionality of the Cortix Class.

Cortix: a program for system-level modules coupling, execution, and analysis.

The Launcher class handles the main funcitonality of stepping through the simulation time, and monitoring its progress.

```
run()
```

Function used to timestep through the modules. Runs the simulation from start to end, and monitors its progress at each time step.

1.3. cortix main 2

### 1.6 module

```
Cortix Module class defintion.
Cortix: a program for system-level modules coupling, execution, and analysis.
class module .Module (parent_work_dir=None,
                                                      library_name=None,
                                                                              library_parent_dir=None,
                           mod_config_node=<cortix.src.utils.configtree.ConfigTree object>)
     Bases: object
     The Module class encapsulates a computational module of some scientific domain.
     execute (slot_id, runtime_cortix_param_file, runtime_cortix_comm_file)
          Spawns a worker process to execute the module.
     get_port_mode (port_name)
          Returns the port mode specified by port_name
     get_port_type (port_name)
          Returns the port type specified by port_name
     has port name(port name)
          Returns true if a port with the name port_name is available in the module.
     library_name
          str – Module library name
     library parent dir
          str – Library parent directory
     name
          str – Module name
     port_names
          list(tuple) – List of names of module's ports
     ports
          list(tuple) – Module's ports
1.7 network
Network class for the Cortix project. A network defines the connectivity between Cortix modules.
Cortix: a program for system-level modules coupling, execution, and analysis.
class network .Network (net_config_node=<cortix.src.utils.configtree.ConfigTree object>)
     Bases: object
```

# Cortix network class definition. Defines the manner in which Modules interact. \_\_repr\_\_() Network to string conversion \_\_str\_\_() Network to string conversion connectivity list(dict) - List of the network connectivity get\_runtime\_cortix\_comm\_file(slot\_name) Returns the cortix comm file that corresponds to slot\_name. None if otherwise.

1.6. module 3

### 1.8 simulation

Simulation class of Cortix.

Cortix: a program for system-level modules coupling, execution, and analysis.

Bases: object

Cortix Simulation element as defined in the Cortix config.

```
execute (task_name=None)
```

This method allows for the execution of a simulation by executing each task, if any. Execution proceeds one task at a time.

### 1.9 task

```
Valmor F. de Almeida dealmeidav@ornl.gov; vfda
```

Cortix: a program for system-level modules coupling, execution, and analysis.

A Task is work done by a Simulation handled by Cortix. A Task will use a given Application.

```
evolve_time
    float - Task final time

evolve_time_unit
    str - Task final time unit

execute (application)
    This method is used to execute (accomplish) the given task.

name
    str - Task name

runtime_cortix_param_file
    str - Task's config file

set_runtime_cortix_param_file (full_path)
```

Sets the task config file to the specified file.

1.8. simulation 4

```
start_time
    float - Task initial time

start_time_unit
    str - Task initial time unit

time_step
    float - Magnitude of incremental step in the task's time

time_step_unit
    str - Time step unit

work_dir
    str - Working directory of task
```

### 1.10 utils

### 1.10.1 configtree

This file contains the class definition of ConfigTree, which aids in parsing the XML configuration files used within the Cortix project.

Cortix: a program for system-level modules coupling, execution, and analysis.

```
class configtree.ConfigTree (config_tree_node=None, config_file_name=None)
    Bases: object
```

This class generates objects that hold an ElementTree node of an XML tree structure. The level of the node depends on the argument passed when creating the object. If a node is passed, that node and all its subnodes are held. If a filename is passed, instead, an XML file is read and the root node is held at the top of the tree.

```
get all sub nodes(tag)
```

Returns a list of all nodes in the element tree that contain a given tag.

```
get_node_children()
```

Returns a list of all the nodes in the element tree.

```
get_node_name()
```

Returns the name associated with the root node of the element tree

```
get_node_tag()
```

Returns the tag associated with the root node of the element tree.

```
get_node_type()
```

Returns the type associated with the root node of the element tree.

```
get_root_node()
```

Returns the Element tree's root node

```
get sub node (tag)
```

Returns a subnode of the element tree specified by the parameter tag.

### 1.10.2 set logger level

This file contains a helper function used by functions across the Cortix project to set the level of the logger.

```
set_logger_level.set_logger_level (handler, handler_name, handler_level)
```

This is a helper function that takes in a file/console handler and sets its logger level accordingly.

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**CHAPTER** 

**TWO** 

### **MODULIB**

# 2.1 pyplot

### 2.1.1 cortix driver

Cortix driver for the PyPlot module.

### 2.1.2 pyplot

PyPlot module.

```
Author: Valmor F. de Almeida dealmeidav@ornl.gov; vfda Tue Jun 24 01:03:45 EDT 2014

class pyplot.PyPlot (slot_id, input_full_path_file_name, work_dir, ports=[], cortix_start_time=0.0, cortix_final_time=0.0, cortix_time_step=0.0, cortix_time_unit=None)

Bases: object

call_ports (cortix_time=0.0)

Transfer data at cortix_time

execute (cortix_time=0.0, time_step=0.0)
```

### 2.1.3 time\_sequence

Valmor F. de Almeida dealmeidav@ornl.gov; vfda

Pyplot module.

This class manages time-sequence data in XML or tabular formats. It is a helper for reading and manipulating stored file data in Cortix. The XML data is a ElementTree object.

Sat Jul 19 12:13:05 EDT 2014

2.1. pyplot 7

**CHAPTER** 

THREE

### **EXAMPLES**

# 3.1 console\_run

### 3.1.1 droplet\_run

Cortix: a program for system-level modules coupling, execution, and analysis.

```
droplet_run.run()
```

Run the Cortix Droplet example. If Cortix and its dependencies are installed, this program should be executed at the command prompt inside the directory this program resides, namely, cortix/cortix/example/console\_run/directory.

### 3.1.2 main executor

Cortix: a program for system-level modules coupling, execution, and analysis.

Cortix is a library and it is used by means of a driver. This file is a simple example of a driver. Many Cortix objects can be ran simultaneously; a single object may be sufficient since many simulation/tasks can be ran via one object.

As Cortix evolves additional complexity may be added to this driver and/or other driver examples can be created.

```
main_executor.main()
```

### 3.1.3 main\_mpi

Cortix: a program for system-level modules coupling, execution, and analysis.

Cortix is a library and it is used by means of a driver. This file is a simple example of a driver. Many Cortix objects can be ran simultaneously; a single object may be sufficient since many simulation/tasks can be ran via one object.

As Cortix evolves additional complexity may be added to this driver and/or other driver examples can be created.

```
main_mpi.main()
```

### 3.1.4 main\_pthread

Cortix: a program for system-level modules coupling, execution, and analysis.

Cortix is a library and it is used by means of a driver. This file is a simple example of a driver. Many Cortix objects can be ran simultaneously; a single object may be sufficient since many simulation/tasks can be ran via one object.

As Cortix evolves additional complexity may be added to this driver and/or other driver examples can be created.

```
main_pthread.main()
```

### 3.2 modulib

### 3.2.1 droplet

### cortix\_driver

Cortix driver for the PyPlot module.

### droplet

Droplet module example in Cortix.

```
class droplet.Droplet (slot_id,
                                           input_full_path_file_name,
                                                                          work dir,
                                                                                        ports=[],
                                                                                                       cor-
                               tix_start_time=0.0, cortix_final_time=0.0, cortix_time_step=0.0,
                                                                                                      cor-
                               tix_time_unit=None)
     Bases: object
     Droplet module used example in Cortix.
     _Droplet__evolve (cortix_time=0.0, cortix_time_step=0.0)
           ODE IVP problem: Given the initial data at t=0, u_1(0)=x_0, u_2(0)=v_0=\dot{u}_1(0) solve \frac{du}{dt}=f(u)
           in the interval 0 \le t \le t_f. When u_1(t) is negative, bounce the droplet to a random height between 0 and
           1.2 x_0 with no velocity, and continue the time integration until t \le t_f.
     call ports (cortix time=0.0)
           Transfer data at cortix_time
     execute (cortix_time=0.0, cortix_time_step=0.0)
           Evolve system from cortix_time to cortix_time + cortix_time_step
```

### 3.2.2 wind

### cortix\_driver

Cortix driver for the PyPlot module.

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```
call_ports (cortix_time=0.0)
    Call all ports at cortix_time

execute (cortix_time=0.0, time_step=0.0)
    Evolve system from cortix_time to cortix_time + time_step
```

### wind

```
Wind module example in Cortix.
```

3.2. modulib

**CHAPTER** 

**FOUR** 

### **SUPPORT**

### 4.1 actor

This is a simple way to hide the name of species of interest in a simulation. The user would modify and copy this class into the Cortix module of interest and keep it private.

Author: Valmor de Almeida dealmeidav@ornl.gov; vfda Sat Aug 15 13:41:12 EDT 2015

```
class actor.Actor(name)
Bases: object
See atoms list in Specie.
atoms
formula
```

# 4.2 fuel bucket

Author: Valmor de Almeida dealmeidav@ornl.gov; vfda

```
class fuel_bucket.FuelBucket (specs=Empty DataFrame Columns: [] Index: [])
    Bases: object
    cladding_end_thickness
    cladding_mass
    cladding_phase
    cladding_volume
    cladding_wall_thickness
    fresh_u235_mass
    fresh_u238_mass
    fresh_u-mass
    fuel_enrichment
    fuel_mass
```

```
fuel_mass_unit
fuel_phase
fuel_radioactivity
fuel_volume
gamma_pwr
get_cladding_end_thickness()
get_cladding_mass()
get_cladding_phase()
get_cladding_volume()
get_cladding_wall_thickness()
get_fresh_u235_mass()
get_fresh_u238_mass()
get_fresh_u_mass()
get_fuel_enrichment()
get_fuel_mass()
get_fuel_mass_unit()
get_fuel_phase()
get_fuel_radioactivity()
get_fuel_volume()
get_gamma_pwr()
get_heat_pwr()
get_inner_slug_id()
get_inner_slug_od()
get_n_slugs()
get_name()
get_outer_slug_id()
get_outer_slug_od()
get_radioactivity()
get_slug_cladding_volume()
get_slug_fuel_volume()
get_slug_length()
get_slug_type()
heat_pwr
inner_slug_id
inner_slug_od
n_slugs
```

4.2. fuel bucket

```
name
outer_slug_id
outer_slug_od
radioactivity
set_cladding_phase(phase)
set_fuel_phase(phase)
set_slug_length(x)
slug_cladding_volume
slug_fuel_volume
slug_length
slug_type
```

# 4.3 fuel\_bundle

This FuelBundle class is a container for usage with other plant-level process modules. It is meant to represent a fuel bundle of an oxide fuel LWR reactor. There are three main data structures:

- 1. fuel bundle specs
- 2. solid phase
- 3. gas phase

The container user will have to provide all the data and from then on, this class will help acess the data. The printing methods reveal the contained data.

Author: Valmor de Almeida dealmeidav@ornl.gov; vfda Sun Dec 27 15:06:55 EST 2015

```
class fuel_bundle.FuelBundle(specs=Empty DataFrame Columns: [] Index: [])
    Bases: object
    fresh_u235_mass
    fresh_u238_mass
    fresh_u_mass
    fuel_enrichment
    fuel_mass
    fuel_mass_unit
    fuel_pin_length
    fuel_pin_radius
    fuel_pin_volume
    fuel_radioactivity
    fuel_rod_od
    fuel_volume
    gamma_pwr
```

4.3. fuel bundle

```
gas_mass
gas_phase
gas_radioactivity
get_fresh_U235_mass()
get_fresh_u238_mass()
get_fresh_u_mass()
get_fuel_enrichment()
get_fuel_mass()
get_fuel_mass_unit()
get_fuel_pin_length()
get_fuel_pin_radius()
get_fuel_pin_volume()
get_fuel_radioactivity()
get_fuel_rod_od()
get_fuel_volume()
get_gamma_pwr()
get_gas_mass()
get_gas_phase()
get_gas_radioactivity()
get_heat_pwr()
get_n_fuel_rods()
get_name()
get_radioactivity()
get_solid_phase()
heat_pwr
n_fuel_rods
name
radioactivity
set_fuel_pin_length(x)
set_gas_phase(phase)
set_solid_phase(phase)
solid_phase
```

4.3. fuel bundle

# 4.4 fuel\_segment

```
Fuel segment Author: Valmor de Almeida dealmeidav@ornl.gov; vfda Sat Jun 27 14:46:49 EDT 2015

class fuel_segment.FuelSegment (geometry=Series([], dtype: float64), species=[])

Bases: object

geometry

get_attribute (name, nuclide=None, series=None)

get_geometry()

get_specie (name)

get_species()

species
```

# 4.5 fuelsegmentsgroups

```
Author: Valmor de Almeida dealmeidav@ornl.gov; vfda

Fuel segment

VFdALib support classes

Sat Jun 27 14:46:49 EDT 2015

class fuelsegmentsgroups.FuelSegmentsGroups (key=None, fuelSegments=None)

Bases: object

AddGroup (key, fuelSegments=None)

GetAttribute (groupKey=None, attributeName=None, nuclideSymbol=None, nuclideSeries=None)

GetFuelSegments (groupKey=None)

HasGroup (key)

RemoveFuelSegment (groupKey, fuelSegment)
```

# 4.6 fuelslug

Author: Valmor de Almeida dealmeidav@ornl.gov; vfda Fuel slug

### 4.6.1 ATTENTION:

This container requires two Phase() containers which are by definition histories. The history is not checked. Therefore any inconsistency will be propagated forward. A fuel slug has two solid phases: cladding and fuel. The user will decide how to best use the underlying history data in the Phase() container of each phase.

VFdALib support classes

Thu Dec 15 16:18:39 EST 2016

4.4. fuel segment 15

### 4.7 nuclides

Author: Valmor de Almeida dealmeidav@ornl.gov; vfda

Nuclides container. The purpose of the this container is to store and query a table of nuclides. Typically the table is filled in with data from an ORIGEN calculation or some other fission/transmutation code.

VFdALib support classes

```
Sat Jun 27 14:46:49 EDT 2015
```

```
class nuclides.Nuclides(propertyDensities=Empty DataFrame Columns: [] Index: [])
    Bases: object
    GetAttribute(name, symbol=None, series=None)
```

# 4.8 periodictable

Properties of the chemical elements.

Each chemical element is represented as an object instance. Physicochemical and descriptive properties of the elements are stored as instance attributes.

```
Author Christoph Gohlke Version 2015.01.29
```

Radiochemical data (isotopes) has been added to this table (2015-2016) Origin: http://www.radiochemistry.org/ Valmor F. de Almeida: dealmeidavf@gmail.com; dealmeidav@ornl.gov

### 4.8.1 Requirements

• CPython 2.7 or 3.4

4.7. nuclides

### References

- 1. http://physics.nist.gov/PhysRefData/Compositions/
- 2. http://physics.nist.gov/PhysRefData/IonEnergy/tblNew.html
- 3. http://en.wikipedia.org/wiki/%(element.name)s
- 4. http://www.miranda.org/~jkominek/elements/elements.db

### **Examples**

```
>>> from elements import ELEMENTS
>>> len(ELEMENTS)
109
>>> str(ELEMENTS[109])
'Meitnerium'
>>> ele = ELEMENTS['C']
>>> ele.number, ele.symbol, ele.name, ele.eleconfig
(6, 'C', 'Carbon', '[He] 2s2 2p2')
>>> ele.eleconfig_dict
{(1, 's'): 2, (2, 'p'): 2, (2, 's'): 2}
>>> sum(ele.mass for ele in ELEMENTS)
14659.1115599
>>> for ele in ELEMENTS:
... ele.validate()
... ele = eval(repr(ele))
```

# 4.9 phase

Phase history container. When you think of a phase value, think of that value at a specific point in time.

ATTENTION: The species (list of Specie) AND quantities (list of Quantity) data members have ARBITRARY density values either at an arbitrary point in the history or at no point in the history. This needs to be removed in the future to avoid confusion.

To obtain history values, associated to the phase, at a particular point in time, use the GetValue() method to access the history data frame (pandas) via columns and rows. The corresponding values in species and quantities are OVERRI-DEN and NOT to be used through the phase interface.

Author: Valmor F. de Almeida dealmeidav@ornl.gov; vfda Sat Sep 5 01:26:53 EDT 2015

class phase.Phase (time\_stamp=None, species=None, quantities=None)
 Bases: object

AddQuantity (newQuant)

AddRow (try\_time\_stamp, row\_values)

AddSpecie (new\_specie)

ClearHistory (value=0.0)

GetActors ()

GetColumn (actor)

GetOuantities ()

4.9. phase 17

```
GetQuantity (name)
     GetRow (try_time_stamp=None)
     GetSpecie (name)
     GetSpecies()
     GetTimeStamps()
     GetValue (actor, try_time_stamp=None)
     ResetHistory (try_time_stamp=None, value=None)
     {\tt ScaleRow}\ (\textit{try\_time\_stamp}, \textit{value})
     {\tt SetSpecieId}\,(name, val)
     SetValue (actor, value, try_time_stamp=None)
     WriteHTML (fileName)
     quantities
     species
     timeStamps
4.10 quantity
Author: Valmor de Almeida dealmeidav@ornl.gov; vfda
This Quantity class is to be used with other classes in plant-level process modules.
For unit testing do at the linux command prompt: python quantity.py
Sat Sep 5 12:51:34 EDT 2015
class quantity.Quantity(name='null-quantity',
                                                       formalName='null-quantity',
                                                                                       value=0.0,
                               unit='null-unit')
     Bases: object
     GetFormalName()
     GetUnit()
     GetValue()
     SetFormalName(fn)
     SetName(n)
     SetUnit(f)
     SetValue(v)
     formalName
     get_name()
     name
     unit
```

4.10. quantity 18

value

# 4.11 specie

Author: Valmor de Almeida dealmeidav@ornl.gov; vfda

This Specie class is to be used with other classes in plant-level process modules.

**NB:** Species is always used either in singular or plural cases, the class named here reflects one species. If many species are used in an external context, the species object name can be used without conflict.

For unit testing do at the linux command prompt: python specie.py

**NB:** The Specie() class encapsulates either the molecular or empirical chemical formula of a compound. The definition of a chemical species here is extended to ficticious compounds. This is done as follows. Say MAO2 is either a molecular or empirical chemical formula of a ficticious compound denoting minor actinides dioxide. The list of atoms is given as follows:

```
['0.49*Np-237', '0.42*Am-241', '0.08*Am-243', '0.01*Cm-244', '2.0*O-16']
```

note the MA forming nuclides add to 1 = 0.49 + 0.42 + 0.08 + 0.01. Therefore the number of atoms in this compound is 3. 1 MA "atom" and 2 O. Note that the total number of "atoms" is obtained by summing all multipliers: 0.49 + 0.42 + 0.08 + 0.01 + 2.0. The nuclide is indicated by the element symbol followed by a dash and the atomic mass number. Here the number of nuclide types is 5 (self.\_nNuclideTypes).

The numbers preceding the nuclide symbol before the \* will be referred to as multipliers. The sum of the multipliers will add to the number of "atoms" in the formula. WARNING: a multiplier could be in the format 0.00e-00. In this case a hiphen may appear twice, e.g.: 1.549e-09\*U-233

Other forms can be used for common true species

```
['Np-237', '2.0*O-16'] or ['Np-237', 'O-16', 'O-16'] or ['2*H', 'O'] or ['H', 'O', 'H'] etc...
```

This code will calculate the molar mass of any species with a given valid atom list using a provided periodic table of chemical elements. The user can also reset the value of the molar mass with a setter method.

Sat May 9 21:40:48 EDT 2015 created; vfda

GetMolarCC()

GetMolarCCUnit()
GetMolarGammaPwr()

GetMolarHeatPwr()

GetMolarMass()

GetMolarGammaPwrUnit()

GetMolarHeatPwrUnit()

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```
GetMolarMassUnit()
GetMolarRadioactivity()
GetMolarRadioactivityFractions()
GetMolarRadioactivityUnit()
GetNAtoms()
GetNNuclideTypes()
GetName()
GetPhase()
SetAtoms (atoms)
SetFlag(f)
SetFormula (atoms)
SetFormulaName(f)
SetMassCC(v)
SetMassCCUnit(v)
SetMolarCC(v)
SetMolarCCUnit(v)
SetMolarGammaPwr(v)
{\tt SetMolarGammaPwrUnit}\;(v)
SetMolarHeatPwr(v)
SetMolarHeatPwrUnit(v)
SetMolarMass(v)
SetMolarMassUnit(v)
{\tt SetMolarRadioactivity}\,(v)
SetMolarRadioactivityFractions (fracs)
SetMolarRadioactivityUnit(v)
SetName(n)
SetPhase(p)
atoms
flag
formula
formulaName
massCC
massCCUnit
molarCC
molarCCUnit
molarGammaPwr
```

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```
molarGammaPwrUnit
molarHeatPwr
molarHeatPwrUnit
molarMass
molarMassUnit
molarRadioactivity
molarRadioactivityFractions
molarRadioactivityUnit
nAtoms
nNuclideTypes
name
phase
```

### 4.12 stream

```
Author: Valmor F. de Almeida dealmeidav@ornl.gov; vfda
Stream container
VFdALib support classes
Sat Aug 15 17:24:02 EDT 2015
class stream.Stream(timeStamp, species=None, quantities=None, values=0.0)
     Bases: object
     GetActors()
     GetQuantities()
     GetQuantity(name)
     GetRow (timeStamp=None)
     GetSpecie (name)
     GetSpecies()
     GetTimeStamp()
     GetValue (actor, timeStamp=None)
     SetSpecieId (name, val)
     SetValue (actor, value=None, timeStamp=None)
```

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