RayPyNG

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raypyng provides a simple API to work with RAY-UI, a software for optical simulation of synchrotron beamlines and x-ray systems developed by Helmholtz-Zentrum Berlin.

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CHAPTER

ONE

INSTALLATION

raypyng will work only if using a Linux or a macOS distribution.

1.1 Install RAY-UI

Download the RAY-UI installer from this link, and run the installer.

1.2 Install xvfb

xvfb is a virtual X11 framebuffer server that let you run RAY-UI headless Install xvfb:

```
sudo apt install xvfb
```

Note: xvfb-run script is a part of the xvfb distribution and runs an app on a new virtual X11 server.

1.3 Install raypyng

• You will need Python 3.8 or newer. From a shell ("Terminal" on OSX), check your current Python version.

```
python3 --version
```

If that version is less than 3.8, you must update it.

We recommend installing raypyng into a "virtual environment" so that this installation will not interfere with any existing Python software:

```
python3 -m venv ~/raypyng-tutorial
source ~/raypyng-tutorial/bin/activate
```

Alternatively, if you are a conda user, you can create a conda environment:

```
conda create -n raypyng-tutorial "python>=3.8" conda activate raypyng-tutorial
```

• Install the latest versions of raypyng and ophyd. Also, install IPython (a Python interpreter designed by scientists for scientists).

```
python3 -m pip install --upgrade raypyng ipython
```

• Start IPython:

```
ipython --matplotlib=qt5
```

The flag --matplotlib=qt5 is necessary for live-updating plots to work.

Or, if you wish you use raypyng from a Jupyter notebook, install a kernel like so:

```
ipython kernel install --user --name=raypyng-tutorial --display-name "Python...

→ (raypyng)"
```

You may start Jupyter from any environment where it is already installed, or install it in this environment along-side raypyng and run it from there:

```
pip install notebook
jupyter notebook
```

CHAPTER

TWO

TUTORIAL

2.1 Simulations

2.1.1 Perform Simulations

raypyng is not able to create a beamline from scratch. To do so, use RAY-UI, create a beamline, and save it. What you save is .rml file, which you have to pass as an argument to the Simulate class. In the following example, we use the file for a beamline called *elisa*, and the file is saved in rml/elisa.rml. The hide parameter can be set to true only if *xvfb* is installed.

```
from raypyng import Simulate
rml_file = 'rml/elisa.rml'

sim = Simulate(rml_file, hide=True)
elisa = sim.rml.beamline
```

The elements of the beamline are now available as python objects, as well as their properties. If working in ipython, tab autocompletion is available. For instance to access the source, a dipole in this case:

```
# this is the dipole object
elisa.Dipole
# to acess its parameter, for instance, the photonFlux
elisa.Dipole.photonFlux
# to access the value
elisa.Dipole.photonFlux.cdata
# to modify the value
elisa.Dipole.photonFlux.cdata = 10
```

To perform a simulation, any number of parameters can be varied. For instance, one can vary the photon energy of the source, and set a a certain aperture of the exit slits:

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```
#and then plug them into the Simulation class sim.params=params
```

It is also possible to define coupled parameters. If for instance, one wants to increase the number of rays with the photon energy

The simulations files and the results will be saved in a folder called *RAYPy_simulation_* and a name of your choice, that can be set. This folder will be saved, by default, in the folder where the program is executed, but it can eventually be modified

```
sim.simulation_folder = '/home/raypy/Documents/simulations'
sim.simulation_name = 'test'
```

This will create a simulation folder with the following path and name

```
/home/raypy/Documents/simulations/RAYPy_simulation_test
```

Sometimes, instead of using millions of rays, it is more convenient to repeat the simulations and average the results We can set which parameters of which optical elements can be exported. The number of rounds of simulations can be set like this:

```
# repeat the simulations as many times as needed
sim.repeat = 1
```

One can decide whether want RAY-UI or raypyng to do a preliminary analysis of the results. To let RAY-UI analyze the results, one has to set:

```
sim.analyze = True # let RAY-UI analyze the results
```

In this case, the following files are available to export:

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```
print(sim.possible_exports)
> ['AnglePhiDistribution',
> 'AnglePsiDistribution',
> 'BeamPropertiesPlotSnapshot',
> 'EnergyDistribution',
> 'FootprintAbsorbedRays',
> 'FootprintAllRays',
> 'FootprintOutgoingRays',
```

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```
> 'FootprintPlotSnapshot',
> 'FootprintWastedRays',
> 'IntensityPlotSnapshot',
> 'IntensityX',
> 'IntensityYZ',
> 'PathlengthDistribution',
> 'RawRaysBeam',
> 'RawRaysIncoming',
> 'RawRaysOutgoing',
> 'ScalarBeamProperties',
> 'ScalarElementProperties']
```

To let raypyng analyze the results set:

```
sim.analyze = False # don't let RAY-UI analyze the results
sim.raypyng_analysis=True # let raypyng analyze the results
```

In this case, only these exports are possible

```
print(sim.possible_exports_without_analysis)
> ['RawRaysIncoming', 'RawRaysOutgoing']
```

The exports are available for each optical element in the beamline, ImagePlanes included, and can be set like this:

Finally, the simulations can be run using

```
sim.run(multiprocessing=5, force=True)
```

where the *multiprocessing* parameter can be set either to False or to an int, corresponding to the number of parallel instances of RAY-UI to be used. Generally speaking, the number of instances of RAY-UI must be lower than the number of cores available. If the simulation uses many rays, monitor the RAM usage of your computer. If the computation uses all the possible RAM of the computer the program may get blocked or not execute correctly.

2.1.2 Simulation Output

Expect this folders and subfolders to be created:

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Analysis performed by RAY-UI

If you decided to let RAY-UI do the analysis, you should expect the following files to be saved in your simulation folder:

- one file for each parameter you set with the values that you passed to the program. If for instance, you input the Dipole numberRays, you will find a file called <code>input_param_Dipole_numberRays.dat</code>
- one folder called *round_n* for each repetition of the simulations. For instance, if you set sim.repeat=2 you will have two folders *round_0* and *round_1*
- inside each *round_n* folder you will find the beamline files modified with the parameters you set in *sim.params*, these are the .*rml* files, that can be opened by RAY-UI.
- inside each *round_n* folder you will find your exported files, one for each simulation. If for instance, you exported the *ScalarElementProperties* of the Dipole, you will have a list of files *0_Dipole-ScalarElementProperties.csv*
- *looper.csv* each simulation and its parameters.

Analysis performed by raypyng

If you decided to let raypyng do the analysis, you should expect the following files to be saved in your simulation folder:

- one file for each parameter you set with the values that you passed to the program. If for instance, you input the Dipole numberRays, you will find a file called <code>input_param_Dipole_numberRays.dat</code>
- one folder called *round_n* for each repetition of the simulations. For instance, if you set sim.repeat=2 you will have two folders *round_0* and *round_1*
- inside each *round_n* folder you will find the beamline files modified with the parameters you set in *sim.params*, these are the .*rml* files, that can be opened by RAY-UI.
- inside each *round_n* folder you will find your exported files, one for each simulation. If for instance, you exported the *RawRaysOutgoing* of the Dipole, you will have a list of files *0_Dipole-RawRaysOutgoing.csv*
- for each *RawRaysOutgoing* file, raypyng calculates some properties, and saves a corresponding file, for instance *0_Dipole_analyzed_rays.dat*. Each of these files contains the following information:
 - SourcePhotonFlux
 - NumberRaysSurvived
 - PercentageRaysSurvived
 - PhotonFlux
 - Bandwidth
 - HorizontalFocusFWHM
 - VerticalFocusFWHM

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- In the simulation folder, all the for each exported element is united (and in case of more rounds of simulations averaged) in one single file. For the dipole, the file is called *Dipole.dat*
- looper.csv each simulation and its parameters.

2.2 Recipes

raypyng provides some recipes to make simulations, that simplify the syntax in the script. Two recipes are provided, one to make Resolving Power simulations, one to make Flux simulations.

2.3 List of available examples

In the example folder, the following examples are available:

- example_rml.py in this example is shown how to read, manipulate and save an rml file.
- example_runner.py in this example is shown how to use the RAY-UI API to start RAY-UI, load a file, trace it and export the desired results.
- example_simulation_analyze.py simulate a beamline, let Ray-UI do the analysis
- example_simulation_noanalyze.py simulate a beamline, let raypyng do the analysis
- example_eval_noanalyze_and_analyze.py plots the results of the two previous simulations
- example_simulation_Flux.py simulations using the flux recipe, useful if you intend to simulate the flux of your beamline
- example_simulation_RP.py simulations using the resolving power (RP) recipe, useful if you intend to simulate the RP of your beamline. The reflectivity of every optical element is switched to 100% and not calculated using the substrate and coating(s) material(s). The information about the Flux of the beamline is therefore not reliable.
- example_beamwaist.py raypyng can plot the beam waist of the x-rays across your beamline. It performs simulations using the beam waist recipe, and it exports the RawRaysOutgoing file from each optical element. It then uses a simple geometrical x-ray tracer to propagate each ray until the next optical element and plots the results (both top view and side view). This is still experimental and it may fail.

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CHAPTER

THREE

HOW TO GUIDES

To simplify the scripting, especially when repetitive, there is the possibility to write recipe for raypyng, to perform simulations, and automatize some tasks.

3.1 Recipe

3.1.1 Recipe Template

This the template to use to write a recipe. At the beginning of the file import SimulationRecipe from raypyng and define a the Simulation class as an empty dummy. This will ensure that you have access to all the methods of the Simulation class.

A recipe should containe at least the __init__() method and three more methods: params(), and simulation_name(), and they must have as an argument the simulate class.

Compose the simulation parameters in the params method: The simulation parameter must return a list of dictionaries, where the keys of the dictionaries are parameters of on abject present in the beamline, instances of ParamElement class. The items of the dictionary must be the values that the parameter should assume for the simulations.

Compose the simulation parameters in the params() method: The params() method must return a list of dictionaries. The keys of the dictionaries are parameters of on abject present in the beamline, instances of ParamElement class. The items of the dictionary must be the values that the parameter should assume for the simulations.

Compose the export parameters in the exports() method: The The exports() method must return a list of dictionaries, method must return a list of dictionaries. The keys of the dictionaries are parameters of on abject present in the beamline, instances of ParamElement class. The items of the dictionary is the name of the file that you want to export (print the output of Simulation.possible_exports and possible_exports_without_analysis.

Define the name to give to the simulation folder in simulation_name()

```
from raypyng.recipes import SimulationRecipe

class Simulate: pass

class MyRecipe(SimulationRecipe):
    def __init__(self):
        pass

    def params(self,sim:Simulate):

        params = []
```

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```
return params

def exports(self,sim:Simulate):
    exports = []
    return exports

def simulation_name(self,sim:Simulate):
    self.sim_folder = ...
    return self.sim_folder
```

3.1.2 How To Write a Recipe

An example of how to write a recipe that exports file for each element present in the beamline automatically.

```
class ExportEachElement(SimulationRecipe):
"""At one defined energy export a file for each
optical elements
def __init__(self, energy:float,/,nrays:int=None,sim_folder:str=None):
    Args:
        energy_range (np.array, list): the energies to simulate in eV
        nrays (int): number of rays for the source
        sim_folder (str, optional): the name of the simulation folder. If None,
→ the rml filename will be used. Defaults to None.
    .....
    if not isinstance(energy, (int,float)):
       raise TypeError('The energy must be an a int or float, while it is a', ...
→type(energy))
    self.energy = energy
    self.nrays = nrays
    self.sim_folder = sim_folder
def params(self,sim:Simulate):
   params = []
    # find source and add to param with defined user energy range
    found source = False
    for oe in sim.rml.beamline.children():
        if hasattr(oe, "photonEnergy"):
        self.source = oe
            found source = True
            break
    if found_source!=True:
```

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```
raise AttributeError('I did not find the source')
    params.append({self.source.photonEnergy:self.energy})
    # set reflectivity to 100%
    for oe in sim.rml.beamline.children():
            for par in oe:
                try:
                    params.append({par.reflectivityType:0})
                except:
                    pass
    # all done, return resulting params
   return params
def exports(self,sim:Simulate):
    # find all the elements in the beamline
   oe_list=[]
    for oe in sim.rml.beamline.children():
        oe_list.append(oe)
    # compose the export list of dictionaries
    exports = []
    for oe in oe_list:
        exports.append({oe:'RawRaysOutgoing'})
   return exports
def simulation_name(self,sim:Simulate):
    if self.sim_folder is None:
        return 'ExportEachElement'
    else:
        return self.sim_folder
if __name__ == "__main__":
    from raypyng import Simulate
    import numpy as np
    import os
    rml_file = ('rml_file.rml')
            = Simulate(rml_file, hide=True)
    sim.analyze = False
   myRecipe = ExportEachElement(energy=1000,nrays=10000,sim_folder=
→'MyRecipeTest')
    # test resolving power simulations
    sim.run(myRecipe, multiprocessing=5, force=True)
```

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API

4.1 Simulation

4.1.1 Simulate

class raypyng.simulate.Simulate(rml=None, hide=False, ray_path=None, **kwargs)

A class that takes care of performing the simulations with RAY-UI

Parameters

- **rml** (*RMLFile/string*, *optional*) string pointing to an rml file with the beamline template, or an RMLFile class object. Defaults to None.
- hide (bool, optional) force hiding of GUI leftovers, xvfb needs to be installed. Defaults to False.
- ray_path (str, optional) the path to the RAY-UI installation folder. If None, the program will look for RAY-UI in the standard installation paths.

property analyze

Turn on or off the RAY-UI analysis of the results. The analysis of the results takes time, so turn it on only if needed

Returns

True: analysis on, False: analysis off

Return type

bool

property exports

The files to export once the simulation is complete. for a list of possible files check self.possible_exports and self.possible_exports_without_analysis.

It is expected a list of dictionaries, and for each dictionary the key is the element to be exported and the valuee are the files to be exported

property params

The parameters to scan, as a list of dictionaries. For each dictionary the keys are the parameters elements of the beamline, and the values are the values to be assigned.

property path

The path where to execute the simlations

Returns

by default the path is the current path from which the program is executed

Return type

string

property possible_exports

A list of the files that can be exported by RAY-UI

Returns

list of the names of the possible exports for RAY-UI

Return type

list

property possible_exports_without_analysis

A list of the files that can be exported by RAY-UI when the analysis option is turned off

Returns

list of the names of the possible exports for RAY-UI when analysis is off

Return type

list

property raypyng_analysis

Turn on or off the RAYPyNG analysis of the results.

Returns

True: analysis on, False: analysis off

Return type

bool

reflectivity(reflectivity=True)

Switch the reflectivity of all the optical elements in the beamline on or off.

Parameters

reflectivity (*bool*, *optional*) – If True the reflectivity is switched on, if False the reflectivity is switched off. Defaults to True.

property repeat

The simulations can be repeated an arbitrary number of times If the statitcs are not good enough using 2 millions of rays is suggested to repeat them instead of increasing the number of rays

Returns

the number of repetition of the simulations, by default is 1

Return type

int

property rml

RMLFile object instantiated in init

rml_list()

This function creates the folder structure and the rml files to simulate. It requires the param to be set. Useful if one wants to create the simulation files for a manual check before starting the simulations.

run(recipe=None, /, multiprocessing=True, force=False)

This method starts the simulations. params and exports need to be defined.

Parameters

• recipe (SimulationRecipe, optional) – If using a recipee pass it as a parameter. Defaults to None.

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- multiprocessing (boolint, optional) If True all the cpus are used. If an integer n is provided, n cpus are used. Defaults to True.
- **force** (*bool*, *optional*) If True all the similations are performed, even if the export files already exist. If False only the similations for which are missing some exports are performed. Defaults to False.

save_parameters_to_file(dir)

save all the user input parameters to file. It takes the values from the SimulationParams class

Parameters

dir (*str*) – the folder where to save the parameters

property simulation_name

A string to append to the folder where the simulations will be executed.

4.1.2 SimulationParams

class raypyng.simulate.**SimulationParams**(rml=None, param_list=None, **kwargs)

The entry point of the simulation parameters.

A class that takes care of the simulations parameters, makes sure that they are written correctly, and returns the the list of simulations that is requested by the user.

Parameters

- rml (RMLFile/string, optional) string pointing to an rml file with the beamline template, or an RMLFile class object. Defaults to None.
- param_list (list, optional) list of dictionaries containing the parameters and values to simulate. Defaults to None.

property params

The parameters to scan, as a list of dictionaries. For each dictionary the keys are the parameters elements of the beamline, and the values are the values to be assigned.

property rml

RMLFile object instantiated in init

4.2 Recipes

4.2.1 Resolving Power

Recipe for resolving power simulations.

The eflectivity is automatically switched off for all elements, the source if automatically recognized.

Parameters

- energy_range (np.array, list) the energies to simulate in eV
- exported_object (ObjectElement) the object to export

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- **source** (ObjectElement, *optional*) the source object. If None is provided, an automatic recognition of the source will be tried. Defaults to None.
- **sim_folder** (*str*, *optional*) the name of the simulation folder. If None, the rml filename will be used. Defaults to None.

4.2.2 Flux

class raypyng.recipes.**Flux**(*energy_range: range, exported_object:* ObjectElement, /, *args, source:

Optional[ObjectElement] = None, sim_folder: Optional[str] = None)

Recipe for flux simulations.

The reflectivity is automatically switched on for all elements, and the source is automatically identified.

Parameters

- energy_range (np.array, list) the energies to simulate in eV
- exported_object (ObjectElement) the object to export
- **source** (ObjectElement, *optional*) the source object. If None is provided, an automatic recognition of the source will be tried. Defaults to None.
- **sim_folder** (*str*, *optional*) the name of the simulation folder. If None, the rml filename will be used. Defaults to None.

4.3 Process simulation files

4.3.1 PostProcess rays analyzed by raypyng

class raypyng.postprocessing.PostProcess

class to post-process the data.

It works only if the exported data are RawRaysOutgoing

cleanup($dir_path: Optional[str] = None, repeat: int = 1, exp_elements: Optional[list] = None)$

Reads all the results of the postprocessing process and summarize them in a single file for each exported object.

This functions reads all the temporary files created by self.postptocess_RawRays() saves one file for each exported element in dir_path, and deletes the temporary files. If more than one round of simulations was done, the values are averaged.

Parameters

- **dir_path** (*str*, *optional*) The path to the folder to cleanup. Defaults to None.
- repeat(int, optional) number of rounds of simulations. Defaults to 1.
- **exp_elements** (*list*, *optional*) the exported elements names as str. Defaults to None.

extract_nrays_from_source(rml_filename)

Extract photon flux from rml file, find source automatically

Parameters

rml_filename (str) – the rml file to use to extract the photon flux

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Returns

the photon flux

Return type

str

PostProcess rountine of the RawRaysOutgoing extracted files.

The method looks in the folder dir_path for a file with the filename: filename = os.path.join(dir_path,sim_number+exported_element + '-' + exported_object+'.csv') for each file it calculates the number of rays, the bandwidth, and the horizontal and vertical focus size, it saves it in an array that is composed by [n_rays,bandwidth,hor_focus,vert_focus], that is then saved to os.path.join(dir_path, sim_number+exported_element+'_analyzed_rays.npy')

Parameters

- **exported_element** (*list*, *optional*)—a list of containing the exported elements name as str. Defaults to None.
- **exported_object** (*str*, *optional*) the exported object, tested only with RawRaysOutgoing. Defaults to None.
- dir_path (str, optional) the folder where the file to process is located. Defaults to None.
- **sim_number** (*str*, *optional*) the prefix of the file, that is the simulation number with a _prepended, ie θ _. Defaults to None.

4.3.2 PostProcess rays analyzed by RAY-UI

class raypyng.postprocessing.PostProcessAnalyzed

class to analyze the data exported by RAY-UI

$moving_average(x, w)$

Computes the morivng average with window w on the array x

Parameters

- **x** (array) the array to average
- w (int) the window for the moving average

Returns

the x array once the moving average was applied

Return type

np.array

retrieve_bw_and_focusSize(folder_name: str, oe: str, nsimulations: int, rounds: int)

Extract the bandwidth and focus size from ScalarBeamProperties of an object.

Parameters

- **folder_name** (str) the path to the folder where the simulations are
- **oe** (*str*) the optical element name
- **nsimulations** (*int*) the number of simulations

• rounds (int) – the number of rounds of simulations

Returns

the bandwidth foc_x np.array: the horizontal focus foc_y np.array: the vertical focus

Return type

bw np.array

retrieve_flux_beamline(folder_name, source, oe, nsimulations, rounds=1, current=0.3)

Extract the flux from object ScalarBeamProperties and from source ScalarElementProperties.

This function takes as arguments the name of the simulation folder, the exported objet in RAY-UI and there number of simulations and returns the flux at the optical element in percentage and in number of photons, and the flux produced by the dipole. It requires ScalarBeamProperties to be exported for the desired optical element, if the source is a dipole it requires ScalarElementProperties to be exported for the Dipole

Parameters

- **folder_name** (str) the path to the folder where the simulations are
- **source** (*str*) the source name
- **oe** (*str*) the optical element name
- **nsimulations** (*int*) the number of simulations
- rounds (int) the number of rounds of simulations
- **current** (*float*, *optional*) the ring current in Ampere. Defaults to 0.3.

Returns

photon_flux (np.array)

[the photon flux at the optical element] flux_percent (np.array): the photon flux in percentage relative to the source_Photon_flux (np.array): the photon flux of the source

else:

flux_percent (np.array): the photon flux in percentage relative to the source

Return type

if the source is a Dipole

4.4 RAY-UI API

4.4.1 RayUIRunner

class raypyng.runner.**RayUIRunner**(ray_path=None, ray_binary='rayui.sh', background=True, hide=False)
RayUIRunner class implements all logic to start a RayUI process, load and rml file, trace and export.

Parameters

- ray_path (str, optional) the path to the RAY-UI installation folder. Defaults to config.ray_path, that will look for the ray_path in the standard installation folders.
- ray_binary (_type_, optional) the binary file of RAY-UI. Defaults to "rayui.sh".
- $\bullet \ \ background\ (bool\ , \ optional) activate\ background\ mode.\ Defaults\ to\ True.$
- hide (bool, optional) Hide the RAY-UI graphical instances. Available only if xvfb is installed. Defaults to False.

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```
property isrunning
           Check weather a process is running and rerutn a boolean
               Returns
                   returns True if the process is running, otherwise False
               Return type
                   bool
     kill()
          kill a RAY-UI process
     property pid
           Get process id of the RayUI process
               Returns
                   PID of the process if it running, None otherwise
               Return type
                   int
     run()
           Open one instance of RAY-UI using subprocess
               Raises
                   RayPyRunnerError – if the RAY-UI executable is not found raise an error
4.5 RML
4.5.1 RMLFile
class raypyng.rml.RMLFile(filename: Optional[str] = None, /, template: Optional[str] = None)
     Read/Write wrapper for the Ray RML files
           Parameters
                 • filename (str, optional) – path to rml file. Defaults to None.
                 • template (str, optional) – path to rml file to use as template. Defaults to None.
     read(file: Optional[str] = None)
           Read rml file
               Parameters
                   file (str, optional) - file name to read. If set to None will use template file name defined
                   during initilizatino of the class. Defaults to None.
     write(file: Optional[str] = None)
           Write the rml to file
               Parameters
```

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file (str, optional) – filename. Defaults to None.

4.5.2 BeamlineElement

```
class raypyng.rml.BeamlineElement(name: str, attributes: dict, **kwargs)
     add_cdata(cdata)
          Store cdata
     add_child(element)
          Store child elements.
     get_attribute(key)
          Get attributes by key
     get_elements(name=None)
          Find a child element by name
     get_full_path()
          Returns the full path of the xml object
                  path of the xml object
              Return type
     resolvable_name()
          Returns the name of the objects, removing lab.beamline.
              Returns
                  name of the object
              Return type
                  str
4.5.3 ObjectElement
```

```
class raypyng.rml.ObjectElement(name: str, attributes: dict, **kwargs)
     add_cdata(cdata)
          Store cdata
     add_child(element)
           Store child elements.
     get_attribute(key)
          Get attributes by key
     get_elements(name=None)
          Find a child element by name
     get_full_path()
           Returns the full path of the xml object
                  path of the xml object
               Return type
                   str
```

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resolvable_name()

Returns the name of the objects, removing lab.beamline.

Returns

name of the object

Return type

str

4.5.4 ParamElement

```
class raypyng.rml.ParamElement(name: str, attributes: dict, **kwargs)
     add_cdata(cdata)
          Store cdata
     add_child(element)
          Store child elements.
     get_attribute(key)
          Get attributes by key
     get_elements(name=None)
          Find a child element by name
     get_full_path()
          Returns the full path of the xml object
               Returns
                   path of the xml object
              Return type
     resolvable_name()
          Returns the name of the objects, removing lab.beamline.
               Returns
                  name of the object
              Return type
                   str
```

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