
Coarse-grained OpenMM Documentation

Release 0.0.1

Shirts research group

**Garrett A. Meek
Lenny T. Fobe
Michael R. Shirts**

**Dept. of Chemical and Biological Engineering
University of Colorado Boulder**

May 12, 2019

CONTENTS

1	OpenMM Simulation() protocols for coarse grained modeling	2
2	Indices and tables	5
	Python Module Index	6
	Index	7

This documentation is generated automatically using Sphinx, which reads all docstring-formatted comments from Python functions in the ‘cg_openmm’ repository. (See `cg_openmm/doc` for Sphinx source files.)

OPENMM SIMULATION() PROTOCOLS FOR COARSE GRAINED MODELING

This page details the functions in `cg_openmm/src/cg_mm_tools/cg_openmm.py`.

```
cg_mm_tools.cg_openmm.build_mm_force(sigma, epsilon, charge, num_beads,
                                         cutoff=Quantity(value=1,
                                                             unit=nanometer))
```

Build an OpenMM ‘Force’ for the non-bonded interactions in our model.

sigma: Non-bonded bead Lennard-Jones interaction distances, (float * simtk.unit.distance)

epsilon: Non-bonded bead Lennard-Jones interaction strength, (float * simtk.unit.energy)

charge: Charge for all beads (float * simtk.unit.charge)

cutoff: Cutoff distance for nonbonded interactions (float * simtk.unit.distance)

num_beads: Total number of beads in our coarse grained model (integer)

```
cg_mm_tools.cg_openmm.build_mm_simulation(topology,                system,
                                             positions,              tempera-
                                             ture=Quantity(value=300.0,
                                                             unit=kelvin),      simula-
                                             tion_time_step=Quantity(value=0.002,
                                                             unit=picosecond),    to-
                                             tal_simulation_time=Quantity(value=1.0,
                                                             unit=picosecond),    out-
                                             put_pdb='output.pdb',
                                             output_data='output.dat',
                                             print_frequency=100)
```

Construct an OpenMM simulation object for our coarse grained model.

topology: OpenMM topology object

system: OpenMM system object

positions: Array containing the positions of all beads in the coarse grained model (np.array('num_beads' x 3 , (float * simtk.unit.distance)))

temperature: Simulation temperature (float * simtk.unit.temperature)

simulation_time_step: Simulation integration time step (float * simtk.unit.time)

total_simulation_time: Total simulation time (float * simtk.unit.time)

output_data: Name of output file where we will write the data from this simulation (string)

print_frequency: Number of simulation steps to skip when writing data to 'output_data' (integer)

`cg_mm_tools.cg_openmm.build_mm_system`(*box_size*, *mass*, *num_beads*,
sigma, *epsilon*, *charge*)

Construct an OpenMM system for our coarse grained model

box_size: Simulation box length (float * simtk.unit.length)

mass: Coarse grained particle mass (float * simtk.unit.length)

num_beads: Total number of beads in our coarse grained model (int)

sigma: Non-bonded bead Lennard-Jones interaction distances, (float * simtk.unit.distance)

epsilon: Non-bonded bead Lennard-Jones interaction strength, (float * simtk.unit.energy)

charge: Charge for all beads (float * simtk.unit.charge)

`cg_mm_tools.cg_openmm.build_mm_topology`(*polymer_length*,
backbone_length,
sidechain_length)

Construct an OpenMM topology for our coarse grained model

polymer_length: Number of monomers in our coarse grained model (integer)

backbone_length: Number of backbone beads on individual monomers in our coarse grained model, (integer)

sidechain_length: Number of sidechain beads on individual monomers in our coarse grained model, (integer)

`cg_mm_tools.cg_openmm.distance`(*positions_1*, *positions_2*)

Construct a matrix of the distances between all particles.

positions_1: Positions for a particle (np.array(length = 3))

positions_2: Positions for a particle (np.array(length = 3))

distance (float * unit)

`cg_mm_tools.cg_openmm.get_box_vectors`(*box_size*)

Assign all side lengths for simulation box.

box_size: Simulation box length (float * simtk.unit.length)

`cg_mm_tools.cg_openmm.set_box_vectors`(*system*, *box_size*)

Build a simulation box.

system: OpenMM system object

box_size: Simulation box length (float * simtk.unit.length)

INDICES AND TABLES

- genindex
- modindex
- search

PYTHON MODULE INDEX

C

`cg_mm_tools.cg_openmm`, [2](#)

INDEX

B

`build_mm_force()` (in module
 [cg_mm_tools.cg_openmm](#)), 2
`build_mm_simulation()` (in module
 [cg_mm_tools.cg_openmm](#)), 2
`build_mm_system()` (in module
 [cg_mm_tools.cg_openmm](#)), 3
`build_mm_topology()` (in module
 [cg_mm_tools.cg_openmm](#)), 3

C

`cg_mm_tools.cg_openmm(module)`, 2

D

`distance()` (in module
 [cg_mm_tools.cg_openmm](#)), 3

G

`get_box_vectors()` (in module
 [cg_mm_tools.cg_openmm](#)), 3

S

`set_box_vectors()` (in module
 [cg_mm_tools.cg_openmm](#)), 3