

- “Reporters” store data about the simulation
- “StateData” gives various options listed in the check boxes
- “DCD” is a binary file format for molecular dynamics trajectories
- “Report Interval” is how often the data are stored
- “Equilibration” is the number of steps before data is stored
- “Production” is the number of steps the simulation is run
- “Minimize” will minimize the energy before running the simulation.
- Let’s set the options as shown on the right

General	System	Integrator	Simulation
Reporters	StateData, DCD		
Report Interval	100		
Equilibration steps	0		
Production steps	1000		
Minimize?	True		
Max minimize steps			
StateData options	<input checked="" type="checkbox"/> Step index <input type="checkbox"/> Time <input checked="" type="checkbox"/> Speed <input checked="" type="checkbox"/> Progress <input checked="" type="checkbox"/> Potential energy <input type="checkbox"/> Kinetic energy <input type="checkbox"/> Total energy <input checked="" type="checkbox"/> Temperature <input type="checkbox"/> Volume <input type="checkbox"/> Density		

- Your script from the OpenMM script builder should look like the window on the right
- Create a directory called “OpenMM” on the Desktop.
- Save the script as “MD_ubq.py” and move it into the “OpenMM” directory.
- Copy “1ubq.pdb” from the VMD tutorial into the same “OpenMM” directory.

```
#####
# this script was generated by openmm-builder. to customize it further,
# you can save the file to disk and edit it with your favorite editor.
#####

from __future__ import print_function
from simtk.openmm import app
import simtk.openmm as mm
from simtk import unit
from sys import stdout

pdb = app.PDBFile('1ubq.pdb')
forcefield = app.ForceField('amber99sbildn.xml', 'amber99_osc.xml')

system = forcefield.createSystem(pdb.topology,
                                nonbondedMethod=app.CutoffNonPeriodic, nonbondedCutoff=1.0*unit.nanometers,
                                constraints=app.HBonds, rigidWater=True)
integrator = mm.LangevinIntegrator(300*unit.kelvin, 1.0/unit.picoseconds,
                                   2.0*unit.femtoseconds)
integrator.setConstraintTolerance(0.00001)

platform = mm.Platform.getPlatformByName('CPU')
simulation = app.Simulation(pdb.topology, system, integrator, platform)
simulation.context.setPositions(pdb.positions)

print('Minimizing...')
simulation.minimizeEnergy()

simulation.context.setVelocitiesToTemperature(300*unit.kelvin)
simulation.reporters.append(app.DCDReporter('trajectory.dcd', 100))
simulation.reporters.append(app.StateDataReporter(stdout, 100, step=True,
                                                  time=True, potentialEnergy=True, temperature=True, progress=True,
                                                  remainingTime=True, speed=True, totalSteps=1000, separator='\t'))

print('Running Production...')
simulation.step(1000)
print('Done!')
```

OpenMM Script Builder

Get Help

MD_ubq.py

Save Script

General

System

Integrator

Simulation

Integrator

Langevin

```
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```