

ESPResSo++ Documentation

Release 1.9.3

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Welcome to the homepage of the ESPResSo++ project

ESPResSo++ is an extensible, flexible, fast and parallel simulation software for soft matter research. It is a highly versatile software package for the scientific simulation and analysis of coarse-grained atomistic or bead-spring models as they are used in soft matter research.

ESPResSo and ESPResSo++ have common roots and share parts of the developer/user community. However their development is independent and they are different software packages.

ESPResSo++ is free, open-source software published under the GNU General Public License (GPL).

Please cite this, if you used ESPResSo++ in your research J. D. Halverson, T. Brandes, O. Lenz, A. Arnold, S. Bevc, V. Starchenko, K. Kremer, T. Stuehn, D. Reith, "ESPResSo++: A Modern Multiscale Simulation Package for Soft Matter Systems", Computer Physics Communications, 184 (2013), pp. 1129-1149 DOI: 10.1016/j.cpc.2012.12.004 Online access: http://dx.doi.org/10.1016/j.cpc.2012.12.004

Recent publications where ESPResSo++ was used

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CHAPTER

ONE

OVERVIEW

- highly modularized object oriented and efficient C++ code
- parallelized with MPI
- python user interface
- classical MD simulations with short and long ranged pair, angular or dihedral interactions
- efficient Adaptive Resolution Scheme (AdResS) implementation
- multisystem integrator (e.g. for parallel tempering)
- reads input files of GROMACS, LAMMPS, and ESPResSo

INSTALLATION

The first step in the installation of ESPResSo++ is to download the latest release from the following location:

https://github.com/espressopp/espressopp/releases

On the command line type:

```
tar -xzf espressopp-1.9.3.tgz
```

This will create a subdirectory espressopp-1.9.3

Enter this subdirectory

```
cd espressopp-1.9.3
```

Create the Makefiles using the cmake command. If you don't have it yet, you have to install it first. It is available for all major Linux distributions and also for Mac OS X. (ubuntu,debian: "apt-get install cmake" or get it from http://www.cmake.org)

```
cmake .
```

(the space and dot after *cmake* are necessary)

If cmake doesn't finish successfully (e.g. it didn't find all the libraries) you can tell cmake manually, where to find them by typing:

```
ccmake .
```

This will open an interactive page where all configuration information can be specified. Alternatively, if cmake . complains on missing BOOST or MPI4PY libraries and you had not installed them, you can try

```
cmake . -DEXTERNAL_BOOST=OFF -DEXTERNAL_MPI4PY=OFF
```

In this case, ESPResSo++ will try to use internal Boost and mpi4py libraries.

After successfully building all the Makefiles you should build ESPResSo++ with:

make

(This will take several minutes)

Before beeing able to use the espressopp module in Python you need to source the ESPRC file:

```
source ESPRC
```

(This sets all corresponding environment variables to point to the module, e.g. PYTHONPATH) You have to source this file every time you want to work with espressopp. It would advisable to e.g. source the file in your .bashrc file ("source <path to espressopp>/ESPRC")

In order to use matplotlib.pyplot for graphical output get the open source code from:

http://sourceforge.net/projects/matplotlib

and follow the installation instructions of your distribution.

CHAPTER

THREE

TUTORIAL

3.1 Basic System Setup

ESPResSo++ is implemented as a python module that has to be imported at the beginning of every script:

```
>>> import espressopp
```

ESPResSo++ uses an object called *System* to store some global variables and is also used to keep the connection between some other important modules. We create it with:

```
>>> system = espressopp.System()
```

Starting a new simulation with ESPResSo++ we should have an idea about what we want to simulate. E.g. how big should the simulation box be or what is the density of the system or what are the interactions and the interaction ranges between our particles.

Let us start with the size of the simulation box:

```
>>> box = (10, 10, 10)
```

In many cases you will need a random number generator (e.G. to couple to a temperature bath or to randomly position particles in the simulation box). ESPResSo++ provides its own random number generator (for the experts: see boost/random.hpp) so let's use it:

```
>>> rng = espressopp.esutil.RNG()
```

Our simulation box needs some boundary conditions. We want to use periodic boundary conditions:

```
>>> bc = espressopp.bc.OrthorhombicBC(rng, box)
```

We tell our system object about this:

```
>>> system.bc = bc
>>> system.rng = rng
```

Now we need to decide which parallelization scheme for the particle storage we want to use. In the current version of ESPResSo++ there is only one storage scheme implemented which is *domain decomposition*. Further parallelized storages (e.g. *atom decomposition* or *force decomposition*) will be implemented in future versions.

The domain decomposition storage needs to know how many CPUs (or cores, if there are multicore CPUs) are available for the simulation and how to assign the CPUs to the different domains of our simulation box. Moreover the storage needs to know the maximum interaction range of the particles. In a simple Lennard-Jones fluid this could for example be $r_{cut}=2^{\frac{1}{6}}$. This value together with the *skin* value determines the minimal size for the so called *linked cells* which are used to speed up Verlet list rebuilds (see Frenkel&Smit or Allen&Tildesley for the details).

```
>>> maxcutoff = pow(2.0, 1.0/6.0)
>>> skin = 0.4
```

Tell the system about it:

```
>>> system.skin = skin
```

In the most simple case, if you want to use only one CPU, the nodeGrid and the cellGrid could look like this:

```
>>> nodeGrid = (1,1,1)
>>> cellGrid = (2,2,2)
```

In general you don't need to take care of that yourself. Just use the corresponding ESPResSo++ routines to calculate a reasonable *nodeGrid* and *cellGrid*:

```
>>> nodeGrid = espressopp.tools.decomp.nodeGrid(espressopp.MPI.COMM_WORLD.

→size)
>>> cellGrid = espressopp.tools.decomp.cellGrid(box, nodeGrid, maxcutoff, 
→skin)
```

Now we have all the ingredients we need for the *domain decomposition* storage of our system:

```
>>> ddstorage = espressopp.storage.DomainDecomposition(system, nodeGrid, _ cellGrid)
```

We initialized the DomainDecomposition object with a pointer to our system. We also have to inform the system about the DomainDecomposition storage:

```
>>> system.storage = ddstorage
```

The next module we need is the *integrator*. This object will do the actual work of integrating Newtons equations of motion. ESPResSo++ implements the well known *velocity Verlet* algorithm (see for example Frenkel&Smit):

```
>>> integrator = espressopp.integrator.VelocityVerlet(system)
```

We have to tell the integrator about the basic time step:

```
>>> dt = 0.005
>>> integrator.dt = dt
```

Let's do some math in between:

Note: For 3D vectors like positions, velocities or forces ESPResSo++ provides a so called *Real3D* type, which simplifies handling and arithmetic operations with vectors. 3D coordinates would typically be defined like this:

```
>>> a = espressopp.Real3D(2.0, 5.0, 6.0)
>>> b = espressopp.Real3D(0.1, 0.0, 0.5)
```

Now you could do things like:

In order to make defining vectors even more simple include the line

```
>>> from espressopp import Real3D
```

just at the beginning of your script. This allows to define vectors as:

```
>>> vec = Real3D(2.0, 1.5, 5.0)
```

Back to our simulation:

The most simple simulation we can do is integrating Newtons equation of motion for one particle without any external forces. So let's simply add one particle to the storage of our system. Every particle in ESPResSo++ has a unique particle id and a position (this is obligatory).

Of course nothing will happen when we integrate this. The particle will stay where it is. Add some initial velocity to the particle by adding the follow line to the script:

```
>>> system.storage.modifyParticle(pid, 'v', Real3D(1.0, 0, 0))
```

After particles have been modified make sure that this information is distributed to all CPUs:

```
>>> system.storage.decompose()
```

Now we can propagate the particle by calling the integrator:

```
>>> integrator.run(100)
```

Check the result with:

```
>>> print "The new particle position is: ", system.storage.getParticle(pid).pos
```

Let's add some more particles at random positions with random velocities and random mass and random type 0 or 1. The boundary condition object knows about how to create random positions within the simulation box. We can add all the particles at once by creating a particle list first:

```
>>> particle_list = []
>>> num_particles = 9
>>> for k in range(num_particles):
    pid = 2 + k
>>>
    pos = system.bc.getRandomPos()
>>>
         = Real3D(system.rng(), system.rng(), system.rng())
>>>
     V
     mass = system.rng()
     type = system.rng(2)
     part = [pid, pos, type, v, mass]
     particle_list.append(part)
>>> system.storage.addParticles(particle_list, 'id', 'pos', 'type', 'v', 'mass')
>>> # don't forget the decomposition
>>> system.storage.decompose()
```

To have a look at the overall system there are several possibilities. The easiest way to get a nice picture is by writing out a PDB file and looking at the configuration with some visualization programm (e.g. VMD):

```
>>> filename = "myconf.pdb"
>>> espressopp.tools.pdb.pdbwrite(filename, system)
```

or (if vmd is in your search PATH) you could directly connect to VMD by:

```
>>> espressopp.tools.vmd.connect(system)
```

or you could print all particle information to the screen:

```
>>> for k in range(10):
>>>    p = system.storage.getParticle(k+1)
>>>    print p.id, p.type, p.mass, p.pos, p.v, p.f, p.q
```

3.2 Simple Lennard Jones System

Lets just copy and paste the beginning from the "System Setup" tutorial:

```
>>> import espressopp
>>> from espressopp import Real3D
>>>
>>> system
                 = espressopp.System()
>>> box
                 = (10, 10, 10)
>>> rng
                 = espressopp.esutil.RNG()
>>> bc
                 = espressopp.bc.OrthorhombicBC(rng, box)
>>> system.bc
                 = bc
>>> system.rng
                 = rng
>>> maxcutoff
                = pow(2.0, 1.0/6.0)
>>> skin
                 = 0.4
>>> system.skin = skin
\rightarrow \rightarrow nodeGrid = (1,1,1)
>>> cellGrid
                 = (1, 1, 1)
                 = espressopp.tools.decomp.nodeGrid(espressopp.MPI.COMM_WORLD.
>>> nodeGrid
⇔size)
>>> cellGrid = espressopp.tools.decomp.cellGrid(box, nodeGrid, maxcutoff,
⇔skin)
>>> ddstorage
                 = espressopp.storage.DomainDecomposition(system, nodeGrid,_
→cellGrid)
>>> system.storage = ddstorage
>>> integrator
                 = espressopp.integrator.VelocityVerlet(system)
>>> dt.
                  = 0.005
>>> integrator.dt = dt
```

And lets add some random particles:

```
>>> num_particles = 20
>>> particle_list = []
>>> for k in range(num_particles):
>>> pid = k + 1
>>> pos = system.bc.getRandomPos()
>>> v = Real3D(0,0,0)
>>> mass = system.rng()
>>> type = 0
>>> part = [pid, pos, type, v, mass]
>>> particle_list.append(part)
>>> system.storage.addParticles(particle_list, 'id', 'pos', 'type', 'v', 'mass')
>>> system.storage.decompose()
```

All particles should interact via a Lennard Jones potential:

```
>>> LJPot = espressopp.interaction.LennardJones(epsilon=1.0, sigma=1.0, 

-cutoff=maxcutoff, shift='auto')
```

shift=True means that the potential will be shifted at the cutoff so that potLJ(cutoff)=0 Next we create a VerletList which will than be used in the interaction: (the Verlet List object needs to know from which system to get its particles and which cutoff to use)

```
>>> verletlist = espressopp.VerletList(system, cutoff=maxcutoff)
```

Now create a non bonded interaction object and add the Lennard Jones potential to that:

Tell the system about the newly created NonBondedInteraction object:

```
>>> system.addInteraction(NonBondedInteraction)
```

We should set the langevin thermostat in the integrator to cool down the random particle system:

```
>>> langevin = espressopp.integrator.LangevinThermostat(system)
>>> langevin.gamma = 1.0
>>> langevin.temperature = 1.0
>>> integrator.addExtension(langevin)
```

and finally let the system run and see how it relaxes or explodes:

```
>>> espressopp.tools.analyse.info(system, integrator)
>>> for k in range(100):
>>> integrator.run(10)
>>> espressopp.tools.analyse.info(system, integrator)
```

Due to the random particle positions it may happen, that two or more particles are very close to each other and the resulting repulsive force between them are so high that they 'shoot off' in different directions with very high speed. Usually the numbers are then larger than the computer can deal with. A typical error message you get could look like this:

Note: ERROR: particle 5 has moved to outer space (one or more coordinates are nan)

In order to prevent this, systems that are setup in a random way and thus have strong overlaps between particels have to be "warmed up" before they can be equilibrated.

In ESPResSo++ there are several possible ways of warming up a system. As a first approach one could simply constrain the forces in the integrator. For this purpose ESPResSo++ provides an integrator Extension named CapForces. The two parameters of this Extension are the system and the maximum force that a particle can get. The following python code shows how CapForces can be used. Add it to your Lennard-Jones example just after adding the Langevin Extension:

After the warmup the time step of the integrator can be set to a larger value. The CapForce extension can be disconnected after the warmup to get the original full Lennard-Jones potential back.

```
>>> integrator.dt = 0.005
>>> integrator.step = 0
>>> force_capping.disconnect()
>>> print "warmup finished - force capping switched off."
```

3.2.1 Task 1:

write a python script that creates a random configuration of 1000 Lennard Jones particles with a number density of 0.85 in a cubic simulation box. Warm up and equilibrate this configuration. Examine the output of the command

```
>>> espressopp.tools.analyse.info(system, integrator)
```

after each integration step. How fast is the energy of the system going down? How long do you have to warmup? What are good parameters for dt, force_capping and number of integration steps?

3.3 Advanced Lennard Jones System

This tutorial needs the matplotlib.pyplot and numpy libraries and also VMD to be installed.

```
>>> import espressopp
```

After importing espressopp we import several other Python packages that we want to use for graphical output of some system parameters (e.g. temperature and energy)

```
>>> import math
>>> import time
>>> import matplotlib
>>> matplotlib.use('TkAgg')
>>> import matplotlib.pyplot as plt
>>> plt.ion()
```

We setup a standard Lennard-Jones system with 1000 particles and a density of 0.85 in a cubic siomulation box. ESPResSo++ provides a "shortcut" to setup such a system:

```
>>> N = 1000
>>> rho = 0.85
>>> L = pow(N/rho, 1.0/3)
>>> system, integrator = espressopp.standard_system.LennardJones(N, (L, L, L), dt=0.

$\infty$0001)
```

We also add a Langevin thermostat:

```
>>> langevin = espressopp.integrator.LangevinThermostat(system)
>>> langevin.gamma = 1.0
>>> langevin.temperature = 1.0
>>> integrator.addExtension(langevin)
```

We do a very short warmup in the beginning to get rid of "extremely" high forces

```
>>> force_capping = espressopp.integrator.CapForce(system, 1000000.0)
>>> integrator.addExtension(force_capping)
>>> espressopp.tools.analyse.info(system, integrator)
>>> for k in range(10):
>>> integrator.run(100)
>>> espressopp.tools.analyse.info(system, integrator)
```

Now let's initialize a graph. So that we can have a realtime-view on what is happening in the simulation:

```
>>> plt.figure()
```

We want to observe temperature and energy of the system:

```
>>> T = espressopp.analysis.Temperature(system)
>>> E = espressopp.analysis.EnergyPot(system, per_atom=True)
```

x will be the x-axixs of the graph containg the time. yT and yE will be temperature and energy as y-axes in 2 plots:

```
>>> x = []
>>> yT = []
>>> yE = []
>>> yEmin = 0.0
>>> x.append(integrator.dt * integrator.step)
>>> yT.append(T.compute())
>>> yE.append(E.compute())
>>> yTmax = max(yT)
>>> yEmax = max(yE)
```

Initialize the two graphs ('ro' means red circles, 'go' means green cirlces, see also pyplot documentation)

```
>>> plt.subplot(211)
>>> gT, = plt.plot(x, yT, 'ro')
>>> plt.subplot(212)
>>> gE, = plt.plot(x, yE, 'go')
```

We also want to observe the configuration with VMD. So we have to connect to vmd. This command will automatically start vmd (vmd has to be found in your PATH environment for this to work)

```
>>> sock = espressopp.tools.vmd.connect(system)
>>> for k in range(200):
>>> integrator.run(1000)
>>> espressopp.tools.vmd.imd_positions(system, sock)
```

Update the x-, and y-axes:

```
>>> x.append(integrator.dt * integrator.step)
>>> yT.append(T.compute())
>>> yE.append(E.compute())
>>> yTmax = max(yT)
>>> yEmax = max(yE)
```

Plot the temperature graph

```
>>> plt.subplot(211)
>>> plt.axis([x[0], x[-1], yTmin, yTmax*1.2])
>>> gT.set_ydata(yT)
>>> gT.set_xdata(x)
>>> plt.draw()
```

Plot the energy graph

```
>>> plt.subplot(212)
>>> plt.axis([x[0], x[-1], yEmin, yEmax*1.2])
>>> gE.set_ydata(yE)
>>> gE.set_xdata(x)
>>> plt.draw()
```

In the end save the equilibrated configurations as .eps and .pdf files

```
>>> plt.savefig('mypyplot.eps')
>>> plt.savefig('mypyplot.pdf')
```

3.4 Polymer Melt

We first import espressopp and then define all the parameters of the simulation:

3.4. Polymer Melt

```
>>> import espressopp
                      = 10
>>> num_chains
>>> monomers_per_chain = 10
>>> L
                      = 10
>>> box
                      = (L, L, L)
                      = 0.97
>>> bondlen
                      = pow(2, 1.0/6.0)
>>> rc
                      = 0.3
>>> skin
>>> dt
                      = 0.005
>>> epsilon
                      = 1.0
>>> sigma
                      = 1.0
```

Like in the simple Lennard Jones tutorial we setup the system and the integrator. First the system with the excluded volume interaction (WCA, Lennard Jones type)

Then the integrator with the Langevin extension

```
>>> integrator = espressopp.integrator.VelocityVerlet(system)
>>> integrator.dt = dt
>>> thermostat = espressopp.integrator.LangevinThermostat(system)
>>> thermostat.gamma = 1.0
>>> thermostat.temperature = temperature
>>> integrator.addExtension(thermostat)
```

Know we add the particles. Keep in mind that we want to create a polymer melt. This means that particles are "bonded" in chains. We setup each polymer chain as a random walk.

```
>>> props = ['id', 'type', 'mass', 'pos', 'v']
>>> vel_zero = espressopp.Real3D(0.0, 0.0, 0.0)
```

In providing bonding information for the particles we "setup" the bonded chains. For this we use the FixedPairList object that needs to know where and in which storage the particles can be found:

```
>>> bondlist = espressopp.FixedPairList(system.storage)
>>> pid = 1
>>> type = 0
>>> mass = 1.0
>>> chain = []
```

ESPResSo++ provides a function that will return position and bond information of a random walk. You have to provide a start ID (particle id) and a starting position which we will get from the random position generator of the boundary condition object:

```
port k in range(monomers_per_chain):
    part = [pid + k, type, mass, positions[k], vel_zero]
    chain.append(part)

pid += monomers_per_chain

type += 1

system.storage.addParticles(chain, *props)

system.storage.decompose()

chain = []

bondlist.addBonds(bonds)
```

Note: try out the command

```
>>> espressopp.tools.topology.polymerRW(pid, startpos, monomers_per_chain, bondlen)
```

to see what it returns

Don't forget to distribute the particles and the bondlist to the CPUs in the end:

```
>>> system.storage.decompose()
```

Finally add the information about the bonding potential. In this example we are using a FENE-potential between the bonded particles.

```
>>> potFENE = espressopp.interaction.FENE(K=30.0, r0=0.0, rMax=1.5)
>>> interFENE = espressopp.interaction.FixedPairListFENE(system, bondlist, potFENE)
>>> system.addInteraction(interFENE)
```

Start the integrator and observe how the system explodes. Like in the random Lennard Jones system, we have the same problem here: particles can strongly overlap and thus will get very high forces accelerating them to infinite (for the computer) speed.

```
>>> espressopp.tools.analyse.info(system, integrator)
>>> for k in range(nsteps):
>>> integrator.run(isteps)
>>> espressopp.tools.analyse.info(system, integrator)
>>> espressopp.tools.analyse.info(system, integrator)
```

3.4.1 Task 2:

Try to warmup and equilibrate a dense polymer melt (density=0.85) by using the warmup methods that you have learned in the Lennard Jones tutorial.

3.4.2 Hint:

During warmup you can slowly switch on the excluded volume interaction by starting with a small epsilon and increasing it during integration: You can do this by continuously overwriting the interaction potential after some time interval.

```
>>> potLJ = espressopp.interaction.LennardJones(new_epsilon, sigma, rc)
>>> interaction.setPotential(type1=0, type2=0, potential=potLJ)
```

3.5 AddNewPotential

The aim of the tutorial is to implement a new interaction potential in ESPResSo++. We start with the Gromos fourth-power bond-stretching potential, because its functional form is simple and its implementation is somewhat

3.5. AddNewPotential 15

similar to other potentials already implemented in ESPResSo++. Everything you learn in this tutorial will then be relevent for implementing any other more complicated potential.

Make sure you have a working, compiled version of ESPResSo++ before starting the tutorial.

For those who are not so familiar with C++ or interfacing python and C++, you will find some helpful notes in the appendix.

3.5.1 Steps for adding a new interaction potential

- 1. Choose the potential and derive the force.
- 2. Choose the appropriate interaction template from those in \$ESPRESSOHOME/src/interaction, e.g. VerletListInteractionTemplate.hpp, FixedTripleListInteractionTemplate.hpp
- 3. Create the .cpp, .hpp and .py files for your potential, place them in \$ESPRESSOHOME/src/interaction and modify \$ESPRESSOHOME/src/interaction/bindings.cpp and \$ESPRESSOHOME/src/interaction/__init__.py
- 4. Compile.

These steps are described in more detail below for our tutorial example potential.

3.5.2 Today's tutorial exercise

Step 1

The potential we are implementing today is a two-body bonded potential with the form

$$V(r_{ij}) = \frac{1}{4}k_{ij}(r_{ij}^2 - r_{0,ij}^2)^2$$

noindent where r_{ij} is the distance between particles i and j. The potential has two input parameters r_0 and k. Derive the force.

Step 2

This is a 2-body interaction between a predefined (fixed) list of atom pairs. What is the appropriate interaction template to use? Choose one in \$ESPRESSOHOME/src/interaction

Open the interaction template file. (When you close the file later, close it without saving, or else later on your compile time will be very long, because of the number of dependencies on the interaction template!) Identify the functions addForces() and computeEnergy(). Many interaction templates also contain functions such as computeVirial(), computeVirialX() (for calculating the virial in slabs along the x-direction) etc.

Find the function calls:

```
potential->_computeForce(force, dist)
```

in addForces() and

```
potential->_computeEnergy(r21)
```

in computeEnergy().

An interaction template can be combined with many different potentials (e.g. harmonic potential, Lennard Jones potential, etc.) Each potential will have its own C++ class containing functions to compute the energy and forces for that particular potential (see e.g. Harmonic.cpp/hpp, LennardJones.cpp/hpp) In turn, each potential can be combined with many different interaction templates.

You don't need to modify the interaction template file today. (Close it without saving!)

Step 3

In this step we create the .cpp, .hpp and .py files for our potential. Let's call the potential FourthPower. The FourthPower.py file will contain the end-user python interface, and in the FourthPower.cpp and FourthPower.hpp files we will create a C++ class for our potential. We will also write a wrapper which will allow the user to call the C++ code from the python interface.

3(a) Interfacing potential class and interaction template

In many cases, it's not necessary to understand the contents of this section in order to implement a new potential. If you like, you can skip directly to Section 3(b) Creating the new potential class.

Now we need to understand how the interaction template will interface with our new class. This is done via a class template, e.g. in Potential.hpp, AnglePotential.hpp, DihedralPotential.hpp etc.

Still in \$ESPRESSOHOME/src/interaction, open the file Potential.hpp. (When you close the file later, close it without saving, or else later on your compile time will be very long, because of the number of dependencies on the file!)

Find the functions _computeForce(Real3D& force, const Real3D& dist) and _computeEnergy(real dist) which you identified in the interaction template. Note _computeForce(Real3D& force, const Real3D& dist) calls function _computeForceRaw(force,dist,distSqr) and computeEnergy(real dist) _computeEnergySqr(dist*dist) which calls _computeEnergySqrRaw(distSqr). The functions _computeForceRaw() and _computeEnergySqrRaw() are the new functions we need to write for our new potential. They will be member methods of our new C++ class FourthPower.

You don't need to modify anything in Potential.hpp today. (Close it without saving!)

3(b) Creating the new potential class

An easy way to implement the new C++ class is to identify a previously implemented potential which somewhat resembles your new potential, e.g. here we could take the Harmonic potential, which is also a 2-body potential, and which has also been interfaced with the FixedPairListInteractionTemplate.

Still in \$ESPRESSOHOME/src/interaction, copy the files Harmonic.py, Harmonic.cpp and Harmonic.hpp to new files FourthPower.py, FourthPower.cpp and FourthPower.hpp. In the new files, find and replace all occurences of 'Harmonic' with 'FourthPower', and 'HARMONIC' with 'FOURTH-POWER'.

First modify FourthPower.hpp.

Note the #include statement for FixedPairListInteractionTemplate.hpp and Potential.hpp, the files you examined in $Step\ 2$ and $Step\ 3(a)$ Interfacing potential class and interaction template.

The Harmonic potential had parameters called K and r0. You can reuse these for the FourthPower potential, along with the setters and getters setK, getK, setR0 and getR0. For better efficiency, you could also create a new variable which contains the square of r0.

Now we need functions _computeForceRaw() and _computeEnergySqrRaw(), as explained in Step 3(a) Interfacing potential class and interaction template. Modify these functions to use the functional form of the fourth power potential as derived in Step 1. Note that Real3D dist, which contains the vector between the two particles, has been defined as $r_{p1} - r_{p2}$ (see addForces() in FixedPairListInteractionTemplate.hpp).

Next open Harmonic.py and FourthPower.py.

Here is an example of an end-user's python script to add an interaction using the harmonic potential:

3.5. AddNewPotential 17

```
harmonicbondslist = espresso.FixedPairList(system.storage)
harmonicbondslist.addBonds(bond_list) #bond_list is a list of tuples_

if (particleindex_i, particleindex_j),...]
harmonic_potential = espresso.interaction.Harmonic(K=10.0, r0=1.0, cutoff = 5.0,_

if t = 0.0)
harmonic_interaction = espresso.interaction.FixedPairListHarmonic(system,_

if harmonicbondslist, potential=harmonic_potential)
system.addInteraction(harmonic_interaction)
```

Compare this to the contents of Harmonic.py to understand the python source code.

Our new potential FourthPower can be called by the end-user in a similar way. Since the Harmonic and Fourth-Power potentials have similar input parameters (K, r0) and both use the FixedPairListInteractionTemplate, you don't need to make any further modifications to the file FourthPower.py, besides replacing 'Harmonic' with 'FourthPower'.

Next open FourthPower.cpp.

Here you will find the C++/python interface, in the function registerPython(). If you want to understand this function, you will find details in *Exposing a C++ class or struct to python using boost*. You don't need to make any further modifications to this file, besides replacing 'Harmonic' with 'FourthPower'.

3(c) Including the new class in espressopp

Finally, update the files \$ESPRESSOHOME/src/interaction/bindings.cpp and \$ESPRESSOHOME/src/interaction/__init__.py (for example by copying and modifying all the lines referring to the Harmonic potential so that they now refer to the FourthPower potential). You need to make three modifications: to include the new .hpp file, to call the new registerPython() wrapper, and to import everything in the new python module.

Step 4

Move to the directory \$ESPRESSOHOME. Update the makefiles and compile using the commands:

```
cmake . make
```

3.5.3 Advanced exercise

For an interaction potential of your choosing, follow the steps above to implement it, e.g. a non-bonded two-body interaction, probably using VerletListInteractionTemplate and based on the LennardJones potential, or a bonded three-body interaction, probably using FixedTripleListInteractionTemplate.hpp and based on the AngularHarmonic potential.

You will probably have to write setters and getters for the parameters in your potential in your .hpp file, and make the corresponding modifications to the function registerPython() in the .cpp file and the python user interface in the .py file.

3.6 Appendices

3.6.1 Exposing a C++ class or struct to python using boost

(See http://www.boost.org/doc/libs/1_56_0/libs/python/doc/tutorial/doc/html/python/exposing.html)

Say we have a C++ struct called World:

Now we write the C++ class wrapper for struct World to expose the constructor and the functions greet and set to python:

```
{
    class_<World>("World", init<std::string>())
      .def("greet", &World::greet)
      .def("set", &World::set)
   ;
}
```

If there are additional constructors we can also expose them using \mathtt{def} (), e.g. for an additional constructor which takes two doubles:

```
class_<World>("World", init<std::string>())
   .def(init<double, double>())
   .def("greet", &World::greet)
   .def("set", &World::set)
;
```

We can also expose the data members of the C++ class or struct and the associated access (getter and setter) functions using add_property(), e.g. for the variable myValue with access functions getMyValue and setMyValue:

```
.add_property("myValue",&World::getMyValue,&World::setMyValue)
```

C++ classes and structs may be derived from other classes. Say we have the C++ struct myDerivedStruct which is derived from the struct myBaseStruct:

```
struct myBaseStruct { virtual ~myBaseStruct(); };
struct myDerivedStruct : myBaseStruct {};
```

We can wrap the base class myBaseStruct as explained above:

```
<Base>("Base")
    /*...*/
;
```

Now when we want to wrap the class myDerivedStruct, we tell boost that it is derived from the base class my-BaseStruct:

```
class_<myDerivedStruct, bases<myBaseStruct> > ("myDerivedStruct")
    /*...*/
;
```

3.6.2 C++ templates

See http://www.cplusplus.com/doc/oldtutorial/templates/

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3.6.3 typedef

typedef declaration allows you to create an alias that can be used anywhere in place of a (possibly complex) type name

```
typedef DataType AliasName;
```

3.6.4 Python notes

Syntax for classes in python

(See also https://docs.python.org/2/tutorial/classes.html)

Here is a python class called DerivedClassName which is derived from two other base classes (BaseClassName1 and BaseClassName1), is initialised with two variables x and y which have default values 1 and 2, and contains a function myFunction.

PMI

PMI = parallel method invocation. For more details see the file \$ESPRESSOHOME/src/pmi.py

3.7 Adaptive Resolution Simulations

3.7.1 Theory and Background

ESPResSo++ provides functionality to run adaptive resolution simulations using the Adaptive Resolution Simulation Scheme (AdResS). In AdResS molecules in different regions in a simulation box are described by different non-bonded force fields, typically atomistic (AT) and coarse-grained (CG). These different subregions are interfaced and coupled via a hybrid region, where the interaction smoothly changes. Molecules can diffuse between the different regions and change their interaction on the fly.

There are two different AdResS approaches: The force-based scheme, in which forces are interpolated, as well as the energy-based scheme (Hamiltonian AdResS or H-AdResS) which interpolates on the level of potential energies. In force-based AdResS (see, for example, Praprotnik et al., J. Chem. Phys. 123, 224106 (2005) as well as Annu. Rev. Phys. Chem. 59, 545 (2008)), we have for the net force between the molecules α and β

$$\mathbf{F}_{\alpha|\beta} = \lambda(\mathbf{R}_{\alpha})\lambda(\mathbf{R}_{\beta})\mathbf{F}_{\alpha|\beta}^{AT} + (1 - \lambda(\mathbf{R}_{\alpha})\lambda(\mathbf{R}_{\beta}))\,\mathbf{F}_{\alpha|\beta}^{CG},$$

where $\mathbf{F}_{\alpha|\beta}^{AT}$ is an AT force field based on the individual atoms belonging to the molecules α and β and λ is a position dependent resolution function smoothly changing from 1 in the AT region to 0 in the CG region via the hybrid buffer region. It is evaluated based on the molecules' center of mass positions \mathbf{R}_{α} . Note that there can of course also be bonded interactions, but these are typically not interpolated, as they are computationally usually much cheaper to evaluate than the non-bonded forces. For the sake of clarity, we omit them here.

In H-AdResS (see Potestio et al., Phys. Rev. Lett. 110, 108301 (2013)), interpolation is performed directly on potential energies in the Hamiltonian as

$$H = \sum_{\alpha} \sum_{i \in \alpha} \frac{\mathbf{p}_{\alpha i}^2}{2m_{\alpha i}} + \sum_{\alpha} \left\{ \lambda(\mathbf{R}_{\alpha}) V_{\alpha}^{\mathsf{AT}} + \left(1 - \lambda(\mathbf{R}_{\alpha})\right) V_{\alpha}^{\mathsf{CG}} \right\},$$

where the first term corresponds to the kinetic energy and we again omitted intramolecular interactions. The forces obtained from this Hamiltonian are

$$\begin{split} \mathbf{F}_{\alpha i} &= \sum_{\beta \neq \alpha} \sum_{j \in \beta} \left\{ \frac{\lambda_{\alpha} + \lambda_{\beta}}{2} \mathbf{F}^{\mathrm{AT}}_{\alpha i | \beta j} + \left(1 - \frac{\lambda_{\alpha} + \lambda_{\beta}}{2} \right) \mathbf{F}^{\mathrm{CG}}_{\alpha i | \beta} \right\} \\ &- \left[V^{\mathrm{AT}}_{\alpha} - V^{\mathrm{CG}}_{\alpha} \right] \nabla_{\alpha i} \lambda_{\alpha}. \end{split}$$

The last term, the so-called drift force, comes from applying the position gradient on the position-dependent resolution function λ . It acts only in the hybrid region and unphysically pushes molecules from one region to the other. Therefore, it needs to be corrected. On the other hand, force-based AdResS, contrary to H-AdResS, does not allow a Hamiltonian formulation at all.

Usually, the force fields used in the different regions of the adaptive simulation setup have significantly different pressures given the same temperature and particle density. This pressure gradient leads, in addition to the drift force in H-AdResS, to particles being pushed across the hybrid region. Eventually, the system would evolve to an equilibrium state with a inhomogeneous density profile across the simulation box. Therefore, correction forces needs to be applied in the hybrid region to counter these effects. In H-AdResS one can use a so-called free energy correction (FEC), which on average cancels the drift force in the hybrid region (see Potestio et al., Phys. Rev. Lett. 110, 108301 (2013)). The FEC corresponds to the free energy difference between the subsystems and can, for instance, be derived from Kirkwood thermodynamic integration. An alternative approach which is typically used to cancel the pressure gradient in force-based AdResS is the so-called thermodynamic force (see Fritsch et al., Phys. Rev. Lett. 108, 170602 (2012)). It is derived by constructing the correction directly from the distorted density profile which is obtained without any correction and then refined iteratively.

3.7.2 ESPResSo++ code

Several measures had to be taken to implement adaptive resolution simulations in ESPResSo++. On top of the normal particles, which serve as the CG particles in AdResS, another layer of extra AT particles is introduced such that one has access to both atomistic and CG particles throughout the whole system. A mapping between the two defines which atoms belong to which CG bead. The resolution function λ is implemented as a particle property of the CG particles that is updated after each integration step based on the new positions. This happens in an extension to the Velocity Verlet integrator. The actual adaptive resolution scheme is then implemented via new interaction templates that define how forces and energies are computed in force-based and energy-based AdResS. These templates use for particle pairs in the atomistic region the actual atoms, this is the AT particles, for the force and energy computation while in the CG region they use the CG particles. In the hybrid region, both are used, as defined in the equations above. The drift term of H-AdResS is implemented similarly. Furthermore, the AdResS integrator extension makes sure that the atomistic particles in the CG region travel along with the CG particles and that similarly the CG particles in the AT region are properly updated according to the new atomistic positions after each integration step. The FEC as well as a module to apply the Thermodynamic Force are implemented as integrator extensions.

In the following, we explain the new features step by step (more details about parameters etc. can be found in the documention of the different classes).

Adress Domain Decomposition

When setting up the storage we have to use an appropriate domain decomposition that accommodates storage and proper interprocessor communication of both AT and CG particles.

```
# (H-)AdResS domain decomposition
system.storage = espressopp.storage.DomainDecompositionAdress(system, nodeGrid, ______
cellGrid)
```

Atomistic and Coarse-Grained particles

When adding particles to the storage, we have to define them as atomistic or coarse-grained. This has been implemented as the particle property "adrat". If it is 0, the particle is coarse-grained. If it is 1, it is an atomistic particle.

When adding the particles as above, it is important that a set of atomistic particles belonging to one CG particle appears in the list of particles allParticles always after the corresponding CG particle.

Next, the FixedTupleListAdress defines which atomistic particles belong to which coarse-grained particles.

```
# create FixedTupleList object and add the tuples
ftpl = espressopp.FixedTupleListAdress(system.storage)
ftpl.addTuples(tuples)
system.storage.setFixedTuplesAdress(ftpl)
```

In this example, tuples is a list of tuples, where each tuple itself is another short list in which the first element is the CG particle and the other elements are the AT particles belonging to it. Note that in ESPResSo++ the CG particle is positioned always in the center of mass of its atoms.

Having set up the FixedTupleList, we can also set up an AdResS fixed pair list that defines bonds between AT particles within individual molecules. This is done in the following way:

```
# add bonds between AT particles
fpl = espressopp.FixedPairListAdress(system.storage, ftpl)
fpl.addBonds(bonds)
```

where bonds is a list of bonds between AT particles within CG molecules. Similarly, triple lists for angles, quadruple lists for dihedrals etc. are set up. Compared to conventional bonds, angles, etc. between different normal CG particles one just adds the suffix Adress to the appropriate list object and provides it also with the FixedTupleList (ftpl in the example). Note that you can define several different such fixed pair lists and you can, for example, also in AdResS simulations still use the normal FixedPairList to define bonds between regular CG particles.

AdResS Verlet List

Next, we construct the AdResS Verlet list object for non-bonded interacting particle pairs:

We have to provide the cutoffs of the list as well as the sizes of the atomistic and hybrid regions. The parameter <code>cutoff</code> corresponds to the cutoff used for CG particle pairs with both particles being in the CG region, while <code>adrcut</code> is the cutoff for all other particle pairs (at least one particle of the pair is in the AT or hybrid region). We want to stress that this pair list is build based on the CG particles' positions. Hence, for the AT and hybrid region one needs in some situations to provide a Verlet list cutoff (adrcut) slightly larger than the actual maximum interaction range of the potential, in order to not lose interactions between some atom pairs. Let us clarify this with an example: Thinking of a pair of water molecules, both coarse-grained into single beads, these CG beads could be farther apart than the interaction cutoff. Two hydrogen atoms pointing towards each other, however, could in fact still be in interaction range. Therefore, an appropriate buffer needs to be provided.

The sphereAdr flag decides how to geometrically set up the change in resolution. If it's true, the AT region is a spherical region positioned at adrCenter with radius dEx. If sphereAdr is false, the resolution changes along the x-axis of the system and dEx corresponds to half the width of the AT region. dHy always is the full width of the hybrid region. Instead of providing a 3D position for adrCenter as above, one can also provide a particle ID of a CG particle. In this case, the atomistic region will follow the movement of the particle. This should be only done, however, for force-based AdResS, since it would break the Hamiltonian character of H-AdResS, and also only when using a spherical adaptive geometry. Then, however, it is even possible to provide a list of particle IDs, in which case the AT region corresponds to the overlap of the spherical regions defined by the individual particles provided in the list. It will deform accordingly while these particle move.

Interactions

When adding interactions to the system we have to use the corresponding interaction templates. Here is how to set up a non-bonded interaction in a H-AdResS system:

First, we define the appropriate interaction type, in H-AdResS this is <code>VerletListHadressLennardJones</code>. Next we define the actual potentials. Then we associate them with the H-AdResS interaction and add the interaction to the system. For force-based AdResS the only change required would be to use the <code>VerletListAdressLennardJones</code> interaction.

Note that the here used interaction, <code>VerletListHadressLennardJones</code>, couples only Lennard-Jones-type potentials with tabulated ones. However, there exist more such interaction templates for other potentials and potential combinations.

AdResS Integrator Extension

Finally, we have to set up the AdResS integrator extension:

```
# AdResS integrator extension
adress = espressopp.integrator.Adress(system, verletlist, ftpl, regionupdates = 1)
integrator.addExtension(adress)
```

It takes as arguments the Verlet list and the fixed tuple list. Additionally, for the case of a moving and/or deforming AdResS region based on one or more particles, the parameter regionupdates specifies how regularly we want to update the shape of the AdResS region in number of steps. This is to avoid as much as possible of the additional communication required to inform different processors of the change of the AdResS region. The parameter defaults to 1 and is not used at all for static AdResS regions.

Having set up the AdResS extension, we can distribute all particles in the box and place the CG molecules in the centers of mass of the atoms which they belong to. This can be done conveniently via

```
# distribute atoms and CG molecules according to AdResS domain decomposition,
# place CG molecules in the center of mass
espressopp.tools.AdressDecomp(system, integrator)
```

Free Energy Compensation

When using H-AdResS, we probably want to also employ a FEC. This can be done as follows:

```
# set up FEC

fec = espressopp.integrator.FreeEnergyCompensation(system, center=[Lx/2, Ly/2, Lz/

$\to 2$])

fec.addForce(itype=3, filename="table_fec.tab", type=1)

integrator.addExtension(fec)
```

The FEC takes as arguments the system object as well as the center of the AT region. Then we add the actual force, which needs to be provided in a table (first column: resolution λ , second: energy, third: force). itype defines which type of interpolation should be used for values between the ones provided in the table. 1 corresponds to linear interpolation, 2 to akima splines, 3 to cubic splines. We suggest to use cubic splines. The FEC is applied on CG particles and distributed among the atoms belonging to the CG particle. type specifies the CG particle

type for which this correction should be applied. One can, for example, use different FECs for different molecules types.

Thermodynamic Force

When using force-based AdResS, or, alternatively, in addition to the FEC in H-AdResS, we can use the thermodynamic force. It can be set up in the following way, very similar to the FEC before:

```
# set up Thermodynamic Force
thdforce = espressopp.integrator.TDforce(system, verletlist)
thdforce.addForce(itype=3,filename="table_tf.tab",type=1)
integrator.addExtension(thdforce)
```

It works largely as for the FEC with the following differences: The table should not provide resolution values in the first column but actual distance values, this is, the distance from the (closest) AT region center. This allows to extend the application of the thermodynamic force slightly beyond the borders of the hybrid region where the resolution is constant. Furthermore, the Thermodynamic Force needs the verletlist as argument.

It is also possible to define a thermodynamic force, which is suited for an adaptive resolution setup with an AT region that is constructed via the overlap of several spherical regions. In this case, the extension needs more information:

It gets three more parameters, startdist, enddist and edgeweightmultiplier. startdist explicitely says at which distance from the center of the closest AT region defining particle the thermodynamic force starts to act and enddist says where it ends. Hence, these value should correspond to what is actually written in the table. edgeweightmultiplier is a parameter that speficies how precisely the thermodynamic force should be applied in the overlap regions of different spheres. For most applications, however, 20 should provide reasonable results (for details, see Kreis et al., J. Chem. Theory Comput. 12, 4067 (2016)). The 3 additional parameters are of course also present with some default values in the basic case, but they are ignored unless we have an AT region that is constructed via the overlap of several spherical regions.

3.7.3 Examples

We have provided several example scripts and setups that are available in the ESPResSo++ source code at examples/adress. Most of them are based on published papers.

The reader is strongly encouraged to play around with them and test what happens when the setups are modified. Possible questions to ask are provided at the end of the following subsections, which explain the individual examples in more detail.

Force-AdResS: Tetrahedral Liquid

Subfolder: fadress_tetraliquid. This example consists of the system that was used in the initial work introducing the force-based adaptive resolution method (see Praprotnik et al., J. Chem. Phys. 123, 224106 (2005) and Phys. Rev. E 73, 066701 (2006)). A liquid composed of artificial tetrahedral molecules, i.e. each molecule consists of 4 bonded atoms arranged in a tetrahedral geometry, is coupled to a CG model which describes the molecules as individual beads.

Questions: The geometry is set in such a way that the resolution changes along the x-axis of the box. Try changing the setup such that the AT region is of spherical shape. You can also try removing the thermostat. Does the system conserve energy? Also vary the size of the atomistic region and see what happens. Can you also make the system all-atomistic or all-CG? You can also try to compare computational times.

Force-AdResS: A Protein in Water

Subfolder: fadress_protein. This system is an aqueous solution of the regulatory protein ubiquitin. The atomistic protein and the atomistic water around it is coupled to a coarse-grained water model, which maps water molecules farther away from the protein to single beads. The CG water interaction was parametrized with iterative Boltzmann inversion (IBI). This system is similar to the setup which was used by Fogarty et al. (J. Chem. Phys. 142, 195101 (2015)) to study the structure and dynamics of a protein hydration shell.

Questions: The setup is significantly more complicated than the previous system. Try to understand the script. You can also have a look into the the actual source code and try to understand, for example, how the gromacs parser works. The example is set up as a fully atomistic simulation by setting the size of the atomistic region to a value larger than the simulation box. Try to change the script such that it is an actual adaptive setup. Do not forget the thermodynamic force! Furthermore, how is the high-resolution region positioned now?

Force-AdResS: Self-Adjusting Adaptive Resolution Simulations

Subfolder: fadress_selfadjusting. This setup demonstrates how force-based adaptive resolution simulations with self-adjusting high-resolution regions can be set up (Kreis et al., J. Chem. Theory Comput. 12, 4067 (2016)). The system is a polyalanine-9 molecule in aqueous solution. A spherical AT region is associated with each atom of the peptide such that the overall AT region formed by the overlap of all these spheres elegantly envelops the peptide. The peptide starts in an extended configuration and as it folds, the AT region surrounding it adjusts itself accordingly. At the outside, we use again a coarse-grained IBI single-bead model for the water molecules.

Questions: Can you change the system such that fewer atoms are associated with AT region, for example, only the heavy atoms? Can you change the update frequency of the shape of the AT region?

H-AdResS: Tetrahedral Liquid

Subfolder: hadress_tetraliquid. This is the system used by Potestio et al. in the paper that proposed the H-AdResS method (Phys. Rev. Lett. 110, 108301 (2013)). It is again a simple system composed of tetrahedral molecules that change their resolution and become individual beads in the CG region. The interpolation occurs along the x-axis. This example has three subfolders.

The first folder hadress_tetraliquid_plain runs a simple H-AdResS simulation without any free energy correction. Hence, the drift force strongly pushes molecules from one region to the other. The script contains analysis routines which measure both a density and a pressure profile along the direction of resolution change while the simulation is running. Gathering enough statistics takes a while, but we have also provided reference profiles which are obtained after a sufficiently long simulation. Have a look at them and try to interpret them.

The second folder hadress_tetraliquid_FEC contains the same setup but with a free energy correction. For this, two tables are provided, table_FEC_Helmholtz.dat and table_FEC_Gibbs.dat. They were derived via Kirkwood thermodynamic integration. The first one is based on the Helmholtz free energy difference per particle between the two subsystem, and the second one corresponds to the Gibbs free energy difference per particle. Two density and pressure profiles obtained while applying these correction are also shown. Try to interpret them.

The third folder hadress_tetraliquid_KTI contains a simple implementation of Kirkwood thermodynamic integration (KTI) which could in principle, when run for long enough, be used to derive the FEC. This is not an adaptive resolution simulation. Instead, we tell the AdResS integrator extension that we want to run KTI. Then, the extension does not modify the resolution values associated with the different molecules and we can change them by hand during the simulation. In this way, we can set up a simulation in which we change the resolution of all molecules in the system every few steps and slowly proceed from a complete CG system to an all-atom one. Have a look and try to understand what is going on.

There are many more interesting things you can try out: Are the H-AdResS simulations energy conserving? Add the commented Langevin thermostat and compare. Also vary the timestep. Additionally, you can change the size of the hybrid region. What happens if it becomes smaller or larger? Furthermore, what happens if you change the system from H-AdResS to force-based AdResS?

H-AdResS: Water

Subfolder: hadress_water. This is a slightly more advanced H-AdResS system in which an atomistic model is coupled to a coarse-grained one, mapping the three water atoms onto single beads.

Questions: Feel free to play around with the system. You could also try to figure out, how the gromacs parsers sets up the interactions and chooses the right H-AdResS interactions.

3.8 Thermodynamic integration

3.8.1 Theoretical explanation

Thermodynamic integration (TI) is a method used to calculate the free energy difference between two states A and B. For the theoretical background, see e.g. http://www.alchemistry.org. In this tutorial, we show how to perform TI calculations with ESPResSo++. We calculate the free energy of solvation of methanol in water. The complete python script is available in the ESPResSo+ source code under examples/thd_integration_solvation

To do TI, we define states A and B, with potentials U^A and U^B . We then construct a pathway of intermediate states between A and B by defining a parameter λ that takes values between 0 and 1 and writing the system potential U as a function of λ , U^A and U^B . The free energy difference between the states A and B is then given by

$$\Delta A = \int_0^1 \left\langle \frac{dU(\lambda)}{d\lambda} \right\rangle_{\lambda} d\lambda$$

In practise, we discretise λ and perform a series of MD simulations with different λ values between 0 and 1, sampling $\frac{dU(\lambda)}{d\lambda}$ in each simulation.

To calculate the solvation free energy of methanol in water, we use a box of water containing one methanol molecule. We simulate desolvation via two separate TI calculations. (Note that the procedure described here is decoupling, and solute-solute interactions will be treated differently if you're doing annihilation instead of decoupling, see Note 1.)

Step 1: free energy change for switching off the Coulombic interactions

State A: methanol has full non-bonded (Coulomb and Lennard Jones) interactions with the solvent

State B: methanol has only Lennard Jones interactions with the solvent

Step 2: free energy change for switching off the Lennard Jones interactions

State A: methanol has only Lennard Jones interactions with the solvent

State B: methanol has no interaction with the solvent

Step 1 can be done using a linear function of λ :

$$U(\lambda_C) = (1 - \lambda_C)U_C^A + U_{unaffected}$$

where U_C^A is the solute-solvent Coulombic interaction in state A. In ESPResSo++ the charges used for state A are the particle charges contained in the particle property charge. The charges in state B are zero, so $U_C^B(q)$ does not appear in the expression. (The case where A and B both have non-zero charges is not implemented in ESPResSo++). The term $U_{unaffected}$ is all other parts of the potential that don't change with λ_C including all bonded interactions, any solute-solute Coulombic interactions, solvent-solvent Coulombic interactions and all Lennard-Jones interactions. The parameter λ_C goes from 0 to 1 in Step 1.

Step 2 must be done using a softcore potential because of the singularity in the Lennard-Jones potential at $r_{ij} = 0$.

$$U(\lambda_L) = \sum_{i,j} U_L(r_{ij}, \lambda_L) + U_{unaffected}$$

$$U_L(r_{ij}, \lambda_L) = (1 - \lambda_L) U_H^A(r_A) + \lambda_L U_H^B(r_B)$$

$$r_A = (\alpha \sigma_A^6 \lambda^p + r_{ij}^6)^{1/6}$$

$$r_B = (\alpha \sigma_B^6 (1 - \lambda)^p + r_{ij}^6)^{1/6}$$

The terms $U_H^A(r_A)$ and $U_H^B(r_B)$ are the normal Lennard-Jones 12-6 hardcore potentials:

$$U_H^A(r_A) = 4.0\epsilon_A \left(\frac{\sigma_A}{r_A}^{12} - \frac{\sigma_A}{r_A}^{6}\right)$$

The sum $\sum_{i,j} U_L(r_{ij}, \lambda_L)$ is over all solute-solvent interactions. The term $U_{unaffected}$ is all other parts of the potential that don't change with λ_L including any solute-solute Lennard-Jones interactions and solvent-solvent Lennard-Jones interactions, which are treated using standard hardcore Lennard-Jones. (In this particular example of methanol, there are no solute-solute Lennard-Jones interactions). Finally α and p are adjustable parameters of the softcore potential.

The ESPResSo++ C++ code allows for different values of ϵ_A , ϵ_B , σ_A and σ_B for every pair of atomtypes interacting via this potential. In this example, we will set ϵ_B to 0 (we are switching off the Lennard-Jones interaction). The parameter λ_L goes from 0 to 1 in Step 2.

3.8.2 ESPResSo++ code

We must perform many separate simulations, each with a different λ value. It is convenient to define a list of λ values in the python script and use an index to access a different element of the list in each separate simulation. The script for the first simulation contains these lines:

```
# Parameters for Thermodynamic Integration
stateBIndices = [1,2,3,4,5,6] #indices of the methanol atoms
lambdaVectorCoul = [0.00, 0.05, 0.10, 0.15, 0.20, 0.25, 0.30, 0.35, 0.40, 0.45, 0.
 \hookrightarrow 50,
                                                          0.55, 0.60, 0.65, 0.70, 0.75, 0.80, 0.85, 0.90, 0.95, 1.00, 1.
 →000,
                                                         1.000, 1.000, 1.000, 1.000, 1.000, 1.000, 1.000, 1.000, 1.000,
                                                         1.000, 1.000, 1.000, 1.000, 1.000, 1.000, 1.000, 1.000,
                                                          1.000, 1.000, 1.000, 1.000, 1.000, 1.000, 1.000, 1.000,
                                                         1.000, 1.000, 1.000, 1.000, 1.000, 1.000, 1.000, 1.000,
                                                         1.000, 1.000, 1.000]
lambdaVectorVdwl = [0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00
 →00,
                                                         0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.00, 0.
  \hookrightarrow 025,
                                                          0.050, 0.075, 0.100, 0.125, 0.150, 0.175, 0.200, 0.225, 0.250,
                                                          0.275, 0.300, 0.325, 0.350, 0.375, 0.400, 0.425, 0.450, 0.475,
                                                          0.500, 0.525, 0.550, 0.575, 0.600, 0.625, 0.650, 0.675, 0.700,
                                                          0.725, 0.750, 0.775, 0.800, 0.825, 0.850, 0.875, 0.900, 0.925,
                                                         0.950, 0.975, 1.000]
lambdaIndex = 0
lambdaTICoul = lambdaVectorCoul[lambdaIndex]
lambdaTIVdwl = lambdaVectorVdwl[lambdaIndex]
```

The list lambdaVectorCoul contains the values of λ_C and the list lambdaVectorVdwl contains the values of λ_L . The total number of simulations to do Step 1 and Step 2 will be len(lambdaVectorCoul) or len(lambdaVectorVdwl). We must make a copy of the python script for each simulation, changing each time the value of lambdaIndex.

Next we set up the Coulombic interactions, assuming we already have created a system and a verletlist. The electrostatics method used is generalised reaction field.

Now we set up the softcore Lennard Jones interaction.

```
#atomtypeparameters - dictionary of format {atomtype: {'eps': epsilon, 'sig':..
⇒sigma}}
                      where atomtype is integer and epsilon and sigma are real
#defaults - dictionary containing a key 'combinationrule' with value 1 if the ...
⇔contents
           of atomtypeparameters need to be converted from c6,c12 format to
           epsilon, sigma format; can also be an empty dictionary if no conversion.
→needed
#sigmaSC, alphaSC, powerSC - parameters of the softcore potential
alphaSC = 0.5
powerSC = 1.0
epsilonB = 0.0
sigmaSC = 0.3
lj_adres_interaction = gromacs.setLennardJonesInteractionsTI(system, defaults,
                                       atomtypeparameters, verletlist, nbCutoff,
                                       epsilonB=epsilonB, sigmaSC=sigmaSC,
⇒alphaSC=alphaSC,
                                       powerSC=powerSC, lambdaTI=lambdaTIVdwl,
                                       pidlist=stateBIndices, annihilate=False,
                                       adress=True, ftpl=ftpl)
```

We open an output file. In the first line we write the values of λ_C and λ_L for this simulation.

During the MD run, every x number of MD steps, we return to the python level and calculate the derivatives of the energies with respect to λ .

```
dhdlCoul = qq_adres_interaction.computeEnergyDeriv()
dhdlVdwl = lj_adres_interaction.computeEnergyDeriv()
dhdlF.write(str(time)+" "+str(dhdlCoul)+" "+str(dhdlVdwl)+"\n")
```

After all simulations, we can now average $\frac{dU(\lambda)}{d\lambda}$ for each value of λ_C or λ_L , integrate over λ_C and λ_L , add the values ΔA_C and ΔA_L , and take the negative (because the procedure described here is desolvation and we want the free energy of solvation).

3.8.3 Some notes

1. This example given here uses decoupling (solute-solvent interactions are a function of λ , solute-solute interactions are not affected by changes in λ). In ESPResSo++ it is also possible to do annihilation, where both solute-solvent and solute-solute interactions are a function of λ , by setting annihilate=True when creating the non-bonded interactions.

- 2. The procedure described here is desolvation. To get the free energy of solvation, we take the negative of the value obtained after integration.
- 3. The example Python code snippets here the helper use gromacs.setLennardJonesInteractionsTI and gromacs.setCoulombInteractionsTI \$ESPRESSOHOME/src/tools/convert/gromacs.py, contained in but not necessary. You can do TI with ESPResSo++ without the Gromacs parser directly espresso.interaction.LennardJonesSoftcoreTI espresso.interaction.ReactionFieldGeneralizedTI. See the documentation of these two classes.

CHAPTER

FOUR

USER INTERFACE

4.1 Version - Object

Return version information of espressopp module

Example:

```
>>> version = espressopp.Version()
>>> print "Name = ", version.name
>>> print "Major version number = ", version.major
>>> print "Minor version number = ", version.minor
>>> print "Git revision = ", version.gitrevision
>>> print "boost version = ", version.boostversion
>>> print "Patchlevel = ", version.patchlevel
>>> print "Compilation date = ", version.date
>>> print "Compilation time = ", version.time
```

to print a full version info string:

```
>>> print version.info()
```

espressopp.Version()

4.2 PMI - Parallel Method Invocation

PMI allows users to write serial Python scripts that use functions and classes that are executed in parallel.

PMI is intended to be used in data-parallel environments, where several threads run in parallel and can communicate via MPI.

In PMI mode, a single thread of control (a python script that runs on the *controller*, i.e. the MPI root task) can invoke arbitrary functions on all other threads (the *workers*) in parallel via *call()*, *invoke()* and *reduce()*. When the function on the workers return, the control is returned to the controller.

This model is equivalent to the "Fork-Join execution model" used e.g. in OpenMP.

PMI also allows to create parallel instances of object classes via *create()*, i.e. instances that have a corresponding object instance on all workers. *call()*, *invoke()* and *reduce()* can be used to call arbitrary methods of these instances.

to execute arbitrary code on all workers, *exec_()* can be used, and to import python modules to all workers, use 'import_()'.

4.2.1 Main program

On the workers, the main program of a PMI script usually consists of a single call to the function *startWorker-Loop()*. On the workers, this will start an infinite loop on the workers that waits to receive the next PMI call,

while it will immediately return on the controller. On the workers, the loop ends only, when one of the commands *finalizeWorkers()* or *stopWorkerLoop()* is issued on the controller. A typical PMI main program looks like this:

```
>>> # compute 2*factorial(42) in parallel
>>> import pmi
>>>
>>> # start the worker loop
>>> # on the controller, this function returns immediately
>>> pmi.startWorkerLoop()
>>>
>>> # Do the parallel computation
>>> pmi.import_('math')
>>> pmi.reduce('lambda a,b: a+b', 'math.factorial', 42)
>>>
>>> # exit all workers
>>> pmi.finalizeWorkers()
```

Instead of using *finalizeWorkers()* at the end of the script, you can call *registerAtExit()* anywhere else, which will cause *finalizeWorkers()* to be called when the python interpreter exits.

Alternatively, it is possible to use PMI in an SPMD-like fashion, where each call to a PMI command on the controller must be accompanied by a corresponding call on the worker. This can be either a simple call to *receive()* that accepts any PMI command, or a call to the identical PMI command. In that case, the arguments of the call to the PMI command on the workers are ignored. In this way, it is possible to write SPMD scripts that profit from the PMI communication patterns.

```
>>> # compute 2*factorial(42) in parallel
>>> import pmi
>>>
>>> pmi.exec_('import math')
>>> pmi.reduce('lambda a,b: a+b', 'math.factorial', 42)
```

To start the worker loop, the command *startWorkerLoop()* can be issued on the workers. To stop the worker loop, *stopWorkerLoop()* can be issued on the controller, which will end the worker loop without exiting the workers.

4.2.2 Controller commands

These commands can be called in the controller script. When any of these commands is issued on a worker during the worker loop, a *UserError* is raised.

- call(), invoke(), reduce() to call functions and methods in parallel
- create() to create parallel object instances
- exec_() and import_() to execute arbitrary python code in parallel and to import classes and functions into the global namespace of pmi.
- sync() to make sure that all deleted PMI objects have been deleted.
- finalizeWorkers() to stop and exit all workers
- registerAtExit() to make sure that finalizeWorkers() is called when python exits on the controller
- stopWorkerLoop() to interrupt the worker loop an all workers and to return control to the single workers

4.2.3 Worker commands

These commands can be called on a worker.

- *startWorkerLoop()* to start the worker loop
- receive() to receive a single PMI command

• *call()*, *invoke()*, *reduce()*, *create()* and *exec_()* to receive a single corresponding PMI command. Note that these commands will ignore any arguments when called on a worker.

4.2.4 PMI Proxy metaclass

The Proxy metaclass can be used to easily generate front-end classes to distributed PMI classes. . . .

4.2.5 Useful constants and variables

The pmi module defines the following useful constants and variables:

- is Controller is True when used on the controller, False otherwise
- *isWorker* = not isController
- ID is the rank of the MPI task
- CONTROLLER is the rank of the Controller (normally the MPI root)
- workerStr is a string describing the thread ('Worker #' or 'Controller')
- in Worker Loop is True, if PMI currently executes the worker loop on the workers.

```
espressopp.pmi.exec_(*args)
```

Controller command that executes arbitrary python code on all (active) workers.

exec_() allows to execute arbitrary Python code on all workers. It can be used to define classes and functions on all workers. Modules should not be imported via exec_(), instead import_() should be used.

Each element of args should be string that is executed on all workers.

Example:

```
>>> pmi.exec_('import hello')
>>> hw = pmi.create('hello.HelloWorld')
```

```
espressopp.pmi.import_(*args)
```

Controller command that imports python modules on all (active) workers.

Each element of args should be a module name that is imported to all workers.

Example:

```
>>> pmi.import_('hello')
>>> hw = pmi.create('hello.HelloWorld')
```

```
espressopp.pmi.create(cls=None, *args, **kwds)
```

Controller command that creates an object on all workers.

cls describes the (new-style) class that should be instantiated. args are the arguments to the constructor of the class. Only classes that are known to PMI can be used, that is, classes that have been imported to pmi via <code>exec_()</code> or <code>import_()</code>.

Example:

```
>>> pmi.exec_('import hello')
>>> hw = pmi.create('hello.HelloWorld')
>>> print(hw)
MPI process #0: Hello World!
MPI process #1: Hello World!
...
```

Alternative: Note that in this case the class has to be imported to the calling module and via PMI.

```
>>> import hello
>>> pmi.exec_('import hello')
>>> hw = pmi.create(hello.HelloWorld)
>>> print(hw)
MPI process #0: Hello World!
MPI process #1: Hello World!
...
```

```
espressopp.pmi.call(*args, **kwds)
```

Call a function on all workers, returning only the return value on the controller.

function denotes the function that is to be called, args and kwds are the arguments to the function. If kwds contains keys that start with with the prefix '__pmictr_', they are stripped of the prefix and are passed only to the controller. If the function should return any results, it will be locally returned. Only functions that are known to PMI can be used, that is functions that have been imported to pmi via <code>exec_()</code> or <code>import_()</code>.

Example:

```
>>> pmi.exec_('import hello')
>>> hw = pmi.create('hello.HelloWorld')
>>> pmi.call(hw.hello)
>>> # equivalent:
>>> pmi.call('hello.HelloWorld', hw)
```

Note, that you can use only functions that are know to PMI when *call()* is called, i.e. functions in modules that have been imported via *exec_()*.

```
espressopp.pmi.invoke(*args, **kwds)
```

Invoke a function on all workers, gathering the return values into a list.

function denotes the function that is to be called, args and kwds are the arguments to the function. If kwds contains keys that start with with the prefix '__pmictr_', they are stripped of the prefix and are passed only to the controller.

On the controller, invoke() returns the results of the different workers as a list. On the workers, invoke returns None. Only functions that are known to PMI can be used, that is functions that have been imported to pmi via *exec_()* or *import_()*.

Example:

```
>>> pmi.exec_('import hello')
>>> hw = pmi.create('hello.HelloWorld')
>>> messages = pmi.invoke(hw.hello())
>>> # alternative:
>>> messages = pmi.invoke('hello.HelloWorld.hello', hw)
```

```
espressopp.pmi.reduce(*args, **kwds)
```

Invoke a function on all workers, reducing the return values to a single value.

reduceOp is the (associative) operator that is used to process the return values, function denotes the function that is to be called, args and kwds are the arguments to the function. If kwds contains keys that start with with the prefix '__pmictr_', they are stripped of the prefix and are passed only to the controller.

reduce() reduces the results of the different workers into a single value via the operation reduceOp. reduceOp is assumed to be associative. Both reduceOp and function have to be known to PMI, that is they must have been imported to pmi via *exec* () or *import* ().

Example:

```
>>> pmi.exec_('import hello')
>>> pmi.exec_('joinstr=lambda a,b: "\n".join(a,b)')
>>> hw = pmi.create('hello.HelloWorld')
>>> print(pmi.reduce('joinstr', hw.hello()))
>>> # equivalent:
```

```
espressopp.pmi.sync()
```

Controller command that deletes the PMI objects on the workers that have already been deleted on the controller.

```
espressopp.pmi.receive (expected=None)
```

Worker command that receives and handles the next PMI command.

This function waits to receive and handle a single PMI command. If expected is not None and the received command does not equal expected, raise a *UserError*.

```
espressopp.pmi.startWorkerLoop()
```

Worker command that starts the main worker loop.

This function starts a loop that expects to receive PMI commands until *stopWorkerLoop()* or *finalizeWorkers()* is called on the controller.

```
espressopp.pmi.finalizeWorkers()
```

Controller command that stops and exits all workers.

```
espressopp.pmi.stopWorkerLoop(doExit=False)
```

Controller command that stops all workers.

If doExit is set, the workers exit afterwards.

```
espressopp.pmi.registerAtExit()
```

Controller command that registers the function *finalizeWorkers()* via atexit.

```
class espressopp.pmi.Proxy (name, bases, dict)
```

A metaclass to be used to create frontend serial objects.

```
exception espressopp.pmi.UserError (msg)
```

Raised when PMI has encountered a user error.

4.3 System - Object

The main purpose of this class is to store pointers to some important other classes and thus make them available to C++. In a way the System class can be viewed as a container for system wide global variables. If you need to run more than one system at the same time you can combine several systems with the help of the Multisystem class.

4.3.1 In detail the System class holds pointers to:

- the *storage* (e.g. DomainDecomposition)
- the boundary conditions bc for the system (e.g. OrthorhombicBC)
- a random number generator rng which is for example used by a thermostat
- the skin which is needed for the Verlet lists and the cell grid
- a list of short range interactions that apply to the system these interactions are added with the *addInteraction()* method of the System

Example (not complete):

```
>>> LJSystem = espressopp.System()
>>> LJSystem.bc = espressopp.bc.OrthorhombicBC(rng, boxsize)
>>> LJSystem.rng
```

```
>>> LJSystem.skin = 0.4
>>> LJSystem.addInteraction(interLJ)
espressopp.System()
espressopp.System.addInteraction (interaction, name)
         Parameters
              • interaction -
              • name (string) – The optional name of the interaction.
         Return type bool
espressopp.System.getInteraction(number)
         Parameters number -
         Return type
espressopp.System.getNumberOfInteractions()
         Return type
espressopp.System.removeInteraction(number)
         Parameters number -
         Return type
espressopp.System.removeInteractionByName(self, name)
         Parameters name (str) – The name of the interaction to remove.
espressopp.System.scaleVolume(*args)
         Parameters *args -
         Return type
espressopp.System.setTrace(switch)
         Parameters switch -
```

4.4 BC - Boundary Condition Object

This is the abstract base class for all boundary condition objects. It cannot be used directly. All derived classes implement at least the following methods:

- getMinimumImageVector(pos1, pos2)
- getFoldedPosition(pos, imageBox)
- getUnfoldedPosition(pos, imageBox)
- getRandomPos()

pos, pos1 and pos2 are particle coordinates (type: (float, float, float)). imageBox (type: (int, int, int)) specifies the

```
espressopp.bc.BC.getFoldedPosition(pos, imageBox)
```

Parameters

- pos -
- imageBox (default: None)

Return type

4.5 OrthorhombicBC - Object

Return type

• imageBox -

Like all boundary condition objects, this class implements all the methods of the base class BC, which are described in detail in the documentation of the abstract class BC.

The OrthorhombicBC class is responsible for the orthorhombic boundary condition. Currently only periodic boundary conditions are supported.

Example:

```
>>> boxsize = (Lx, Ly, Lz)
>>> bc = espressopp.bc.OrthorhombicBC(rng, boxsize)
```

espressopp.bc.OrthorhombicBC(rng, boxL)

Parameters

- rng -
- **boxL** (real) (default: 1.0)

 $\verb|espressopp.bc.OrthorhombicBC.setBoxL| (boxL)$

Parameters boxL -

4.6 Storage - Storage Object

This is the base class for all storage objects. All derived classes implement at least the following methods:

• decompose()

Send all particles to their corresponding cell/cpu

• addParticle(pid, pos):

Add a particle to the storage

• removeParticle(pid):

Remove a particle with id number *pid* from the storage.

```
>>> system.storage.removeParticle(4)
```

There is an example in examples folder

• getParticle(pid):

Get a particle object. This can be used to get specific particle information:

```
>>> particle = system.storage.getParticle(15)
>>> print "Particle ID is : ", particle.id
>>> print "Particle position is : ", particle.pos
```

you cannot use this particle object to modify particle data. You have to use the modifyParticle command for that (see below).

• addAdrParticle(pid, pos, last_pos):

Add an AdResS Particle to the storage

- *setFixedTuplesAdress(fixed_tuple_list)*:
- addParticles(particle_list, *properties):

This routine adds particles with certain properties to the storage.

 $\boldsymbol{param\ particle List}\ \ list\ of\ particles\ (and\ properties)$ to be added

param properties property strings

Each particle in the list must be itself a list where each entry corresponds to the property specified in properties.

Example:

```
>>> addParticles([[id, pos, type, ...], 'id', 'pos', 'type', ...

→)
```

• modifyParticle(pid, property, value, decompose='yes')

This routine allows to modify any properties of an already existing particle.

Example:

```
>>> modifyParticle(pid, 'pos', Real3D(new_x, new_y, new_z))
```

• removeAllParticles():

This routine removes all particles from the storage.

• 'system':

The property 'system' returns the System object of the storage.

Examples:

espressopp.storage.Storage.addAdrATParticle(pid, *args)

Parameters

- pid-
- *args -

Return type

```
espressopp.storage.Storage.addParticle(pid, pos)
```

Parameters

```
• pid-
             • pos -
        Return type
espressopp.storage.Storage.addParticles(particleList, *properties)
        Parameters
             • particleList -
             • *properties -
        Return type
espressopp.storage.Storage.clearSavedPositions()
        Return type
espressopp.storage.Storage.getParticle(pid)
        Parameters pid-
        Return type
espressopp.storage.Storage.getRealParticleIDs()
        Return type
espressopp.storage.Storage.modifyParticle(pid, property, value)
        Parameters
             • pid-
             • property -
             • value -
        Return type
espressopp.storage.Storage.particleExists(pid)
        Parameters pid-
        Return type
espressopp.storage.Storage.printRealParticles()
        Return type
espressopp.storage.Storage.removeAllParticles()
        Return type
espressopp.storage.Storage.removeParticle(pid)
        Parameters pid-
        Return type
espressopp.storage.Storage.restorePositions()
        Return type
espressopp.storage.Storage.savePositions(idList)
        Parameters idList -
        Return type
espressopp.storage.Storage.setFixedTuplesAdress(fixedtuples)
        Parameters fixedtuples -
```

4.7 BerendsenBarostat - Berendsen barostat Object

This is the Berendsen barostat implementation according to the original paper [Berendsen84]. If Berendsen barostat is defined (as a property of integrator) then at the each run the system size and the particle coordinates will be scaled by scaling parameter μ according to the formula:

$$\mu = [1 - \Delta t / \tau (P_0 - P)]^{1/3}$$

where Δt - integration timestep, τ - time parameter (coupling parameter), P_0 - external pressure and P - instantaneous pressure.

Example:

```
>>> berendsenP = espressopp.integrator.BerendsenBarostat(system)
>>> berendsenP.tau = 0.1
>>> berendsenP.pressure = 1.0
>>> integrator.addExtension(berendsenP)
```

!IMPORTANT In order to run *npt* simulation one should separately define thermostat as well (e.g. Berendsen-Thermostat).

Definition:

In order to define the Berendsen barostat

```
>>> berendsenP = espressopp.integrator.BerendsenBarostat(system)
```

one should have the System defined.

Properties:

• berendsenP.tau

The property 'tau' defines the time parameter τ .

• berendsenP.pressure

The property 'pressure' defines the external pressure P_0 .

Setting the integration property:

```
>>> integrator.addExtension(berendsenP)
```

It will define Berendsen barostat as a property of integrator.

One more example:

```
>>> berendsen_barostat = espressopp.integrator.BerendsenBarostat(system)
>>> berendsen_barostat.tau = 10.0
>>> berendsen_barostat.pressure = 3.5
>>> integrator.addExtension(berendsen_barostat)
```

Canceling the barostat:

If one do not need the pressure regulation in system anymore or need to switch the ensamble or whatever:)

```
>>> # define barostat with parameters
>>> berendsen = espressopp.integrator.BerendsenBarostat(system)
>>> berendsen.tau = 0.8
>>> berendsen.pressure = 15.0
>>> integrator.addExtension(berendsen)
>>> ...
>>> # some runs
>>> ...
>>> # disconnect Berendsen barostat
```

```
>>> berendsen.disconnect()
>>> # the next runs will not include the system size and particle_
--coordinates scaling
```

Connecting the barostat back after the disconnection

```
>>> berendsen.connect()
```

References:

espressopp.integrator.BerendsenBarostat(system)

Parameters system -

4.8 BerendsenThermostat - Berendsen thermostat Object

This is the Berendsen thermostat implementation according to the original paper [Berendsen84]. If Berendsen thermostat is defined (as a property of integrator) then at the each run the system size and the particle coordinates will be scaled by scaling parameter λ according to the formula:

$$\lambda = [1 + \Delta t / \tau_T (T_0 / T - 1)]^{1/2}$$

where Δt - integration timestep, τ_T - time parameter (coupling parameter), T_0 - external temperature and T - instantaneous temperature.

Example:

```
>>> berendsenT = espressopp.integrator.BerendsenThermostat(system)
>>> berendsenT.tau = 1.0
>>> berendsenT.temperature = 1.0
>>> integrator.addExtension(berendsenT)
```

Definition:

In order to define the Berendsen thermostat

```
>>> berendsenT = espressopp.integrator.BerendsenThermostat(system)
```

one should have the System defined.

Properties:

• berendsenT.tau

The property 'tau' defines the time parameter τ_T .

• berendsenT.temperature

The property 'temperature' defines the external temperature T_0 .

Setting the integration property:

```
>>> integrator.addExtension(berendsenT)
```

It will define Berendsen thermostat as a property of integrator.

One more example:

```
>>> berendsen_thermostat = espressopp.integrator.BerendsenThermostat(system)
>>> berendsen_thermostat.tau = 0.1
>>> berendsen_thermostat.temperature = 3.2
>>> integrator.addExtension(berendsen_thermostat)
```

Canceling the thermostat:

```
>>> # define thermostat with parameters
>>> berendsen = espressopp.integrator.BerendsenThermostat(system)
>>> berendsen.tau = 2.0
>>> berendsen.temperature = 5.0
>>> integrator.addExtension(berendsen)
>>> ...
>>> # some runs
>>> ...
>>> # disconnect Berendsen thermostat
>>> berendsen.disconnect()
```

Connecting the thermostat back after the disconnection

```
>>> berendsen.connect()
```

espressopp.integrator.BerendsenThermostat(system)

Parameters system -

4.9 LangevinBarostat - Langevin-Hoover barostat Object

This is the barostat implementation to perform Langevin dynamics in a Hoover style extended system according to the paper [Quigley04]. It includes corrections of Hoover approach which were introduced by Martyna et al [Martyna94]. If LangevinBarostat is defined (as a property of integrator) the integration equations will be modified. The volume of system V is introduced as a dynamical variable:

$$\begin{split} \dot{\boldsymbol{r}}_i &= \frac{\boldsymbol{p}_i}{m_i} + \frac{p_\epsilon}{W} \boldsymbol{r}_i \\ \dot{\boldsymbol{p}}_i &= - \bigtriangledown_{\boldsymbol{r}_i} \Phi - (1 + \frac{n}{N_f}) \frac{p_\epsilon}{W} \boldsymbol{p}_i - \gamma \boldsymbol{p}_i + \boldsymbol{R}_i \\ \dot{V} &= dV p_\epsilon / W \\ \dot{p}_\epsilon &= nV (X - P_{ext}) + \frac{n}{N_f} \sum_{i=1}^N \frac{\boldsymbol{p}_i^2}{m_i} - \gamma_p p_\epsilon + R_p \end{split}$$

where volume has a fictitious mass W and associated momentum p_{ϵ} , γ_p - friction coefficient, P_{ext} - external pressure and X - instantaneous pressure without white noise contribution from thermostat, n - dimension, N_f - degrees of freedom (if there are no constrains and N is the number of particles in system $N_f = nN$). R_p - values which are drawn from Gaussian distribution of zero mean and unit variance scaled by

$$\sqrt{\frac{2k_BTW\gamma_p}{\Delta t}}$$

!IMPORTANT Terms $-\gamma p_i + R_i$ correspond to the termostat. They are not included here and will not be calculated if the Langevin Thermostat is not defined.

Example:

!IMPORTANT This barostat is supposed to be run in a couple with thermostat in order to simulate the npt ensamble, because the term R_p needs the temperature as a parameter.

Definition:

In order to define the Langevin-Hoover barostat

```
>>> langevinP = espressopp.integrator.LangevinBarostat(system, rng, _ desiredTemperature)
```

one should have the System and RNG defined and know the desired temperature.

Properties:

• langevinP.gammaP

The property 'gammaP' defines the friction coefficient γ_p .

• langevinP.pressure

The property 'pressure' defines the external pressure P_{ext} .

langevinP.mass

The property 'mass' defines the fictitious mass W.

Methods:

• setMassByFrequency(frequency)

Set the proper langevinP.mass using expression $W = dNk_bT/\omega_b^2$, where frequency, ω_b , is the frequency of required volume fluctuations. The value of ω_b should be less then the lowest frequency which appears in the NVT temperature spectrum [Quigley04] in order to match the canonical distribution. d - dimensions, N - number of particles, k_b - Boltzmann constant, T - desired temperature.

NOTE The *langevinP.mass* can be set both directly and using the (*setMassByFrequency*) (*frequency*))

Adding to the integration:

```
>>> integrator.addExtension(langevinP)
```

It will define Langevin-Hoover barostat as a property of integrator.

One more example:

Canceling the barostat:

If one do not need the pressure regulation in system anymore or need to switch the ensamble or whatever :)

Connecting the barostat back after the disconnection

```
>>> langevinBarostat.connect()
```

References:

espressopp.integrator.LangevinBarostat (system, rng, temperature)

Parameters

- system -
- rng -
- temperature -

4.10 CoulombRSpace - Coulomb potential and interaction Objects (*R* space part)

$$\sum_{i=1}^{N} \sum_{\substack{j>i\\r_{ij} < k_{max}}} \frac{q_i q_j}{r_{ij}} erfc(\alpha r_{ij}) - \frac{\alpha}{\sqrt{\pi}} \sum_{i=1}^{N} q_i^2$$

This is the *R* space part of potential of Coulomb long range interaction according to the Ewald summation technique. Good explanation of Ewald summation could be found here [Allen89], [Deserno98].

Example:

!IMPORTANT Coulomb interaction needs k-space part as well EwaldKSpace.

Definition:

It provides potential object CoulombRSpace and interaction object VerletListCoulombRSpace

The *potential* is based on parameters: Coulomb prefactor (coulomb_prefactor), Ewald parameter (alpha), and the cutoff in R space (rspacecutoff).

```
>>> coulombR_pot = espressopp.interaction.CoulombRSpace(coulomb_prefactor, alpha, rspacecutoff)
```

Potential Properties:

• coulombR_pot.prefactor

The property 'prefactor' defines the Coulomb prefactor.

 $\bullet \ \ coulomb R_pot.alpha$

The property 'alpha' defines the Ewald parameter alpha.

• coulombR_pot.cutoff

The property 'cutoff' defines the cutoff in R space.

The *interaction* is based on the Verlet list (*VerletList*)

```
>>> vl = espressopp.VerletList(system, rspacecutoff+skin)
>>> coulombR_int = espressopp.interaction.VerletListCoulombRSpace(vl)
```

It should include at least one potential

>>> coulombR_int.setPotential(type1=0, type2=0, potential = coulombR_pot)

Interaction Methods:

• setPotential(type1, type2, potential)

This method sets the *potential* for the particles of *type1* and *type2*. It could be a bunch of potentials for the different particle types.

• getVerletListLocal()

Access to the local Verlet list.

Adding the interaction to the system:

```
>>> system.addInteraction(coulombR_int)
```

espressopp.interaction.CoulombRSpace(prefactor, alpha, cutoff)

Parameters

- prefactor (real) (default: 1.0)
- alpha (real) (default: 1.0)
- cutoff (default: infinity)

espressopp.interaction.VerletListCoulombRSpace(vl)

Parameters v1 -

espressopp.interaction.VerletListCoulombRSpace.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListCoulombRSpace.getVerletList()

Return type A Python list of lists.

espressopp.interaction.VerletListCoulombRSpace.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

4.11 CoulombKSpaceEwald - Coulomb potential and interaction Objects (*K* space part)

$$\frac{1}{2\pi V} \sum_{\substack{m \in \mathbb{Z}^3 \\ 0 \le |m| \le k \max}} \frac{exp(-\frac{\pi^2}{\alpha^2}m'^2)}{m'^2} \left| \sum_{i=1}^N q_i \cdot exp(2\pi i r_i \cdot m') \right|^2$$

This is the *K* space part of potential of Coulomb long range interaction according to the Ewald summation technique. Good explanation of Ewald summation could be found here [Allen89], [Deserno98].

Example:

!IMPORTANT Coulomb interaction needs R space part as well CoulombRSpace.

Definition:

It provides potential object *CoulombKSpaceEwald* and interaction object *CellListCoulombKSpaceE-wald* based on all particles list.

The *potential* is based on the system information (System) and parameters: Coulomb prefactor (coulomb_prefactor), Ewald parameter (alpha), and the cutoff in K space (kspacecutoff).

```
>>> ewaldK_pot = espressopp.interaction.CoulombKSpaceEwald(system, _ 

--coulomb_prefactor, alpha, kspacecutoff)
```

Potential Properties:

• ewaldK_pot.prefactor

The property 'prefactor' defines the Coulomb prefactor.

• ewaldK_pot.alpha

The property 'alpha' defines the Ewald parameter alpha.

• ewaldK_pot.kmax

The property 'kmax' defines the cutoff in *K* space.

The *interaction* is based on the all particles list. It needs the information from Storage and *K* space part of potential.

Interaction Methods:

• getPotential()

Access to the local potential.

Adding the interaction to the system:

```
>>> system.addInteraction(ewaldK_int)
```

References:

espressopp.interaction.CoulombKSpaceEwald(system, prefactor, alpha, kmax)

Parameters

- system -
- prefactor -
- alpha -
- kmax-

espressopp.interaction.CellListCoulombKSpaceEwald(storage, potential)

Parameters

• storage -

• potential -

espressopp.interaction.CellListCoulombKSpaceEwald.getFixedPairList()

Return type A Python list of lists.

espressopp.interaction.CellListCoulombKSpaceEwald.getPotential()

Return type

4.12 decomp.py - Auxiliary python functions

• nodeGrid(n):

It determines how the processors are distributed and how the cells are arranged. n - number of processes

• *cellGrid(box_size, node_grid, rc, skin)*:

It returns an appropriate grid of cells.

• tuneSkin(system, integrator, minSkin=0.01, maxSkin=1.2, precision=0.001):

It tunes the skin size for the current system

• printTimeVsSkin(system, integrator, minSkin=0.01, maxSkin=1.5, skinStep = 0.01):

It prints time of running versus skin size in the range [minSkin, maxSkin] with the step skinStep

4.13 espressopp

4.13.1 espressopp.Exceptions

```
espressopp.Error(msg)

Parameters msg -

espressopp.ParticleDoesNotExistHere(msg)

Parameters msg -

espressopp.UnknownParticleProperty(msg)

Parameters msg -

espressopp.MissingFixedPairList(msg)
```

4.13.2 espressopp.FixedPairDistList

Parameters msg -

Return type

• pid2 -

espressopp.FixedPairDistList.addPairs(bondlist)

```
Parameters bondlist -
        Return type
espressopp.FixedPairDistList.getDist(pid1, pid2)
        Parameters
             • pid1 -
             • pid2 -
        Return type
espressopp.FixedPairDistList.getPairs()
        Return type
espressopp.FixedPairDistList.getPairsDist()
        Return type
espressopp.FixedPairDistList.size()
        Return type
4.13.3 espressopp.FixedPairList
espressopp.FixedPairList (storage)
        Parameters storage -
espressopp.FixedPairList.add(pid1, pid2)
        Parameters
             • pid1 -
             • pid2 -
        Return type
espressopp.FixedPairList.addBonds(bondlist)
        Parameters bondlist -
        Return type
espressopp.FixedPairList.getBonds()
        Return type
espressopp.FixedPairList.getLongtimeMaxBond()
        Return type
espressopp.FixedPairList.resetLongtimeMaxBond()
        Return type
espressopp.FixedPairList.size()
        Return type
```

4.13.4 FixedPairListAdress - Object

The FixedPairListAdress is the Fixed Pair List to be used for AdResS or H-AdResS simulations. When creating the FixedPairListAdress one has to provide the storage and the tuples. Afterwards the bonds can be added. In the example "bonds" is a python list of the form ((pid1, pid2), (pid3, pid4), ...) where each inner pair defines a bond between the particles with the given particle ids.

Example - creating the FixedPairListAdress and adding bonds:

```
>>> ftpl = espressopp.FixedTupleList(system.storage)
>>> fpl = espressopp.FixedPairListAdress(system.storage, ftpl)
>>> fpl.addBonds(bonds)
```

 $\verb|espressopp.FixedPairListAdress| (storage, fixed tuple List)|$

Parameters

- storage -
- fixedtupleList -

espressopp.FixedPairListAdress.add (pid1, pid2)

Parameters

- pid1 -
- pid2 -

Return type

espressopp.FixedPairListAdress.addBonds(bondlist)

Parameters bondlist -

Return type

espressopp.FixedPairListAdress.getBonds()

Return type

4.13.5 espressopp.FixedQuadrupleAngleList

espressopp.FixedQuadrupleAngleList (storage)

Parameters storage -

espressopp.FixedQuadrupleAngleList.add(pid1, pid2, pid3, pid4)

Parameters

- pid1 -
- pid2 -
- pid3 -
- pid4 -

Return type

espressopp.FixedQuadrupleAngleList.addQuadruples (quadruplelist)

Parameters quadruplelist -

Return type

 $\verb|espressopp.FixedQuadrupleAngleList.getAngle(|pid1|, pid2|, pid3|, pid4|)|$

Parameters

- pid1 -
- pid2 -
- pid3 -
- pid4 -

Return type

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```
espressopp.FixedQuadrupleAngleList.getQuadruples()
        Return type
espressopp.FixedQuadrupleAngleList.getQuadruplesAngles()
        Return type
espressopp.FixedQuadrupleAngleList.size()
        Return type
4.13.6 espressopp.FixedQuadrupleList
espressopp.FixedQuadrupleList(storage)
        Parameters storage -
espressopp.FixedQuadrupleList.add(pid1, pid2, pid3, pid4)
        Parameters
             • pid1 -
             • pid2 -
             • pid3 -
             • pid4 -
        Return type
espressopp.FixedQuadrupleList.addQuadruples (quadruplelist)
        Parameters quadruplelist -
        Return type
espressopp.FixedQuadrupleList.getQuadruples()
        Return type
espressopp.FixedQuadrupleList.size()
        Return type
4.13.7 espressopp.FixedSingleList
espressopp.FixedSingleList(storage)
        Parameters storage -
espressopp.FixedSingleList.add(pid1)
        Parameters pid1 -
        Return type
espressopp.FixedSingleList.addSingles (singlelist)
        Parameters singlelist -
        Return type
espressopp.FixedSingleList.getSingles()
        Return type
espressopp.FixedSingleList.size()
        Return type
```

4.13.8 espressopp.FixedTripleAngleList

```
espressopp.FixedTripleAngleList (storage)
        Parameters storage -
espressopp.FixedTripleAngleList.add(pid1, pid2, pid3)
        Parameters
             • pid1 -
             • pid2 -
              • pid3 -
        Return type
espressopp.FixedTripleAngleList.addTriples (triplelist)
        Parameters triplelist -
        Return type
espressopp.FixedTripleAngleList.getAngle(pid1, pid2, pid3)
        Parameters
             • pid1 -
             • pid2 -
             • pid3 -
        Return type
espressopp.FixedTripleAngleList.getTriples()
        Return type
espressopp.FixedTripleAngleList.getTriplesAngles()
        Return type
espressopp.FixedTripleAngleList.size()
        Return type
4.13.9 espressopp.FixedTripleList
espressopp.FixedTripleList (storage)
        Parameters storage -
espressopp.FixedTripleList.add(pid1, pid2, pid3)
        Parameters
             • pid1 -
             • pid2 -
             • pid3 -
        Return type
espressopp.FixedTripleList.addTriples(triplelist)
        Parameters triplelist -
        Return type
espressopp.FixedTripleList.getTriples()
```

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Return type

```
espressopp.FixedTripleList.size()
```

Return type

4.13.10 espressopp.FixedTripleListAdress

espressopp.FixedTripleListAdress(storage, fixedtupleList)

Parameters

- storage -
- fixedtupleList -

espressopp.FixedTripleListAdress.add (pid1, pid2)

Parameters

- pid1 -
- pid2 -

Return type

espressopp.FixedTripleListAdress.addTriples(triplelist)

Parameters triplelist -

Return type

4.13.11 espressopp.FixedTupleList

```
espressopp.FixedTupleList(storage)
```

Parameters storage -

espressopp.FixedTupleList.size()

Return type

4.13.12 FixedTupleListAdress - Object

The FixedTupleListAdress is important for AdResS and H-AdResS simulations. It is the connection between the atomistic and coarse-grained particles. It defines which atomistic particles belong to which coarse-grained particle. In the following example "tuples" is a python list of the form ((pid_CG1, pidAT11, pidAT12, pidAT13, ...), (pid_CG2, pidAT21, pidAT22, pidAT23, ...), ...). Each inner list (pid_CG1, pidAT11, pidAT12, pidAT13, ...) defines a tuple. The first number is the particle id of the coarse-grained particle while the following numbers are the particle ids of the corresponding atomistic particles.

Example - creating the FixedTupleListAdress:

```
>>> ftpl = espressopp.FixedTupleListAdress(system.storage)
>>> ftpl.addTuples(tuples)
>>> system.storage.setFixedTuples(ftpl)
```

espressopp.FixedTupleListAdress(storage)

Parameters storage -

espressopp.FixedTupleListAdress.addTuples(tuplelist)

Parameters tuplelist -

Return type

4.13.13 espressopp.Int3D

```
espressopp.___Int3D(*args)
          Parameters *args -
espressopp.__Int3D.\mathbf{x}(v, [0)
          Parameters
                • v -
                • [0 -
          Return type
espressopp.__Int3D.\mathbf{y}(v,[1)
          Parameters
                • [1-
          Return type
espressopp.__Int3D.\mathbf{z}(v, [2)
          Parameters
                • [2-
          Return type
espressopp.toInt3DFromVector(*args)
          Parameters *args -
espressopp.toInt3D(*args)
          Parameters *args -
espressopp.Int3D.toInt3D(*args)
     Try to convert the arguments to a Int3D, returns the argument, if it is already a Int3D.
espressopp.Int3D.toInt3DFromVector(*args)
     Try to convert the arguments to a Int3D.
     This function will only convert to a Int3D if x, y and z are specified.
```

4.13.14 espressopp.MultiSystem

4.13.15 espressopp.ParallelTempering

4.13.16 espressopp.Particle

```
espressopp.Particle(pid, storage)
Parameters
```

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```
• pid-
```

• storage -

class espressopp.Particle.ParticleLocal (pid, storage)

The local particle.

Throws an exception: * when the particle does not exists locally

TODO: Should throw an exception: * when a ghost particle is to be written * when data is to be read from a ghost that is not available

4.13.17 ParticleAccess - abstract base class for analysis/measurement/io

```
espressopp.ParticleAccess.perform_action()
```

Return type

4.13.18 espressopp.ParticleGroup

```
espressopp.ParticleGroup(storage)
```

Parameters storage -

espressopp.ParticleGroup.add(pid)

Parameters pid -

Return type

espressopp.ParticleGroup.has(pid)

Parameters pid-

Return type

espressopp.ParticleGroup.show()

Return type

espressopp.ParticleGroup.size()

Return type

4.13.19 espressopp.Real3D

```
espressopp.___Real3D(*args)
```

Parameters *args -

espressopp.__Real3D. $\mathbf{x}(v, [0)$

Parameters

- v -
- [O -

Return type

espressopp.__Real3D. $\mathbf{y}(v, [1)$

Parameters

- **v** -
- [1 -

Return type

```
espressopp.__Real3D.z (v, [2)

Parameters

v -

[2 -

Return type

espressopp.toReal3DFromVector(*args)

Parameters *args -

espressopp.toReal3D(*args)

Parameters *args -

espressopp.Real3D.toReal3D(*args)

Try to convert the arguments to a Real3D, returns the argument, if it is already a Real3D.

espressopp.Real3D.toReal3DFromVector(*args)

Try to convert the arguments to a Real3D.
```

This function will only convert to a Real3D if x, y and z are specified.

4.13.20 RealND -

This is the object which represents N-dimensional vector. It is an extended Real3D, basicly, it hase the same functionallity but in N-dimetions. First of all it is usefull for classes in 'espressopp.analysis'.

```
Description
```

```
espressopp.__RealND(*args)

Parameters *args -

espressopp.toRealNDFromVector(*args)

Parameters *args -

espressopp.toRealND(*args)

Parameters *args -

espressopp.RealND.toRealND(*args)

Try to convert the arguments to a RealND, returns the argument, if it is already a RealND.

espressopp.RealND.toRealNDFromVector(*args)

Try to convert the arguments to a RealND.
```

This function will only convert to a RealND if x, y and z are specified.

This function will only convert to a Tensor if x, y and z are specified.

4.13.21 espressopp.Tensor

```
espressopp. Tensor.toTensor(*args)
Try to convert the arguments to a Tensor, returns the argument, if it is already a Tensor.

espressopp. Tensor.toTensorFromVector(*args)
Try to convert the arguments to a Tensor.
```

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4.13.22 espressopp.VerletList

4.13.23 VerletListAdress - Object

The VerletListAdress is the Verlet List to be used for AdResS or H-AdResS simulations. When creating the VerletListAdress one has to provide the system and specify both cutoff for the CG interaction and adrcutoff for the atomistic interaction. Often, it is important to set the atomistic adrcutoff much bigger than the actual interaction's cutoff would be, since also the atomistic part of the VerletListAdress (adrPairs) is built based on the coarse-grained particle positions. For a much larger coarse-grained cutoff it is for example possible to also set the atomistic cutoff on the same value as the coarse-grained one.

Furthermore, the sizes of the explicit and hybrid region have to be provided (dEx and dHy in the example below) and the center of the atomistic region has to be set (adrCenter). Additionally, it can be chosen between a spherical and a slab-like geometry (sphereAdr).

The AdResS region can also be defined based on one or more particles. For a single particle, in this case a spherical region moves along with the particle. For many such region defining particles, the high-resolution/hybrid region corresponds to the overlap of the different spherical regions based on the individual particles (for details see Kreis et al., JCTC doi: 10.1021/acs.jctc.6b00440). Note that more region defining particles mean a higher computational overhead as these particles need to be communicated among all processors (also see explanations in AdResS.py). Also note that region defining particles should be normal/CG particles, not atomistic/AdResS ones.

Bascially the VerListAdress provides 4 lists:

- adrZone: A list which holds all particles in the atomistic and hybrid region
- cgZone: A list which holds all particles in the coarse-grained region
- adrPairs: A list which holds all pairs which have at least one particle in the adrZone, i.e. in the atomistic or hybrid region
- vlPairs: A list which holds all pairs which have both particles in the cgZone, i.e. in the coarse-grained region

Example - creating the VerletListAdress for a slab-type adress region fixed in space (only the x value of adrCenter is used):

```
>>> vl = espressopp.VerletListAdress(system, cutoff=rc, adrcut=rc, dEx=ex_

size, dHy=hy_size, adrCenter=[Lx/2, Ly/2, Lz/2])
```

or

```
>>> vl = espressopp.VerletListAdress(system, cutoff=rc, adrcut=rc, dEx=ex_

size, dHy=hy_size, adrCenter=[Lx/2, Ly/2, Lz/2], sphereAdr=False)
```

Example - creating the VerletListAdress for a spherical adress region centered on adrCenter and fixed in space:

```
>>> vl = espressopp.VerletListAdress(system, cutoff=rc, adrcut=rc, dEx=ex_

size, dHy=hy_size, adrCenter=[Lx/2, Ly/2, Lz/2], sphereAdr=True)
```

Example - creating the VerletListAdress for a spherical adress region centered on one particle and moving with the particle

```
>>> vl = espressopp.VerletListAdress(system, cutoff=rc, adrcut=rc, dEx=ex_

size, dHy=hy_size, pids=[adrCenterPID], sphereAdr=True)
```

Example - creating the VerletListAdress for a adress region based on the overlapping spherical regions by several particles

espressopp. VerletListAdress (system, cutoff, adrcut, dEx, dHy, adrCenter, pids, exclusionlist, sphereAdr)

Parameters

- system -
- cutoff -
- adrcut -
- dEx -
- dHy -
- adrCenter (default: [])
- pids (default: [])
- exclusionlist (default: [])
- **sphereAdr** (default: False)

espressopp.VerletListAdress.addAdrParticles(pids, rebuild)

Parameters

- pids -
- rebuild (default: True)

Return type

espressopp.VerletListAdress.exclude(exclusionlist)

Parameters exclusionlist -

Return type

espressopp.VerletListAdress.rebuild()

Return type

espressopp.VerletListAdress.totalSize()

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Return type

4.13.24 espressopp.VerletListTriple

espressopp.VerletListTriple.localSize()

Return type

espressopp.VerletListTriple.totalSize()

Return type

4.14 analysis

4.14.1 espressopp.analysis.AllParticlePos

```
\verb|espressopp.analysis.AllParticlePos.gatherAllPositions|()|
```

Return type

4.14.2 AnalysisBase - abstract base class for analysis/measurement

This abstract base class provides the interface and some basic functionality for classes that do analysis or observable measurements

It provides the following methods:

```
espressopp.analysis.AnalysisBase.compute()
Computes the instant value of the observable.
```

Return type a python list or a scalar

```
espressopp.analysis.AnalysisBase.getAverageValue()
```

Returns the average value for the observable and the standard deviation.

Return type a python list

```
espressopp.analysis.AnalysisBase.getNumberOfMeasurements()
```

counts the number of measurements that have been performed (standalone or in integrator) does _not_ include measurements that have been done using "compute()"

Return type

```
\verb|espressopp.analysis.AnalysisBase.performMeasurement|()|
```

Computes the observable and updates average and standard deviation

Return type

```
espressopp.analysis.AnalysisBase.reset()
Resets average and standard deviation
```

Return type

4.14.3 espressopp.analysis.Autocorrelation

```
espressopp.analysis.Autocorrelation(system)

Parameters system -

espressopp.analysis.Autocorrelation.clear()

Return type

espressopp.analysis.Autocorrelation.compute()

Return type

espressopp.analysis.Autocorrelation.gather(value)

Parameters value -

Return type
```

4.14.4 espressopp.analysis.CenterOfMass

```
espressopp.analysis.CenterOfMass(system)

Parameters system -
```

4.14.5 espressopp.analysis.ConfigsParticleDecomp

4.14.6 Configurations - Configurations Object

- *gather()* add configuration to trajectory
- clear() clear trajectory
- *back()* get last configuration of trajectory

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- *capacity* maximum number of configurations in trajectory further adding (*gather()*) configurations results in erasing oldest configuration before adding new one capacity=0 means: infinite capacity (until memory is full)
- size number of stored configurations

usage:

storing trajectory

```
>>> configurations = espressopp.Configurations(system)
>>> configurations.gather()
>>> for k in xrange(100):
>>> integrator.run(100)
>>> configurations.gather()
```

accessing trajectory data:

iterate over all stored configurations:

```
>>> for conf in configurations:
```

iterate over all particles stored in configuration:

```
>>> for pid in conf
>>> particle_coords = conf[pid]
>>> print pid, particle_coords
```

access particle with id <pid> of stored configuration <n>:

```
>>> print "particle coord: ",configurations[n][pid]
```

espressopp.analysis.Configurations(system)

Parameters system -

espressopp.analysis.Configurations.back()

Return type

espressopp.analysis.Configurations.clear()

Return type

espressopp.analysis.Configurations.gather()

Return type

4.14.7 ConfigurationsExt - ConfigurationsExt Object

- gather() add configuration to trajectory
- clear() clear trajectory
- back() get last configuration of trajectory
- *capacity* maximum number of configurations in trajectory further adding (*gather()*) configurations results in erasing oldest configuration before adding new one capacity=0 means: infinite capacity (until memory is full)
- size number of stored configurations

usage:

storing trajectory

```
>>> configurations = espressopp.ConfigurationsExt(system)
>>> configurations.gather()
>>> for k in xrange(100):
>>> integrator.run(100)
>>> configurations.gather()
```

accessing trajectory data:

iterate over all stored configurations:

```
>>> for conf in configurations:
```

iterate over all particles stored in configuration:

```
>>> for pid in conf
>>> particle_coords = conf[pid]
>>> print pid, particle_coords
```

access particle with id <pid> of stored configuration <n>:

```
>>> print "particle coord: ",configurations[n][pid]
```

espressopp.analysis.ConfigurationsExt(system)

Parameters system -

espressopp.analysis.ConfigurationsExt.back()

Return type

espressopp.analysis.ConfigurationsExt.clear()

Return type

espressopp.analysis.ConfigurationsExt.gather()

Return type

4.14.8 espressopp.analysis.Energy

```
espressopp.analysis.EnergyPot (system, per_atom)
```

Parameters

- system -
- per_atom (default: False)

espressopp.analysis.EnergyPot.compute()

Return type

espressopp.analysis.EnergyKin(system, per_atom)

Parameters

- system-
- per_atom (default: False)

espressopp.analysis.EnergyKin.compute()

Return type

espressopp.analysis.EnergyTot (system, per_atom)

Parameters

• system -

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4.14.9 espressopp.analysis.IntraChainDistSq

espressopp.analysis.IntraChainDistSq(system, fpl)

Parameters

- system -
- fpl -

espressopp.analysis.IntraChainDistSq.compute()

Return type

4.14.10 LBOutput - abstract base class for analysis / output in LB simulations

Abstract base class for arbitrary output from LB simulations. At the moment, the implemented realisations are:

- \bullet espressopp.analysis.LBOutputScreen to output local density rho and v_z component of the velocity as a function of the coordinate x.
- espressopp.analysis.LBOutputVzInTime to output velocity component v_z of a specific lattice site (the value used at the moment is $0.25 * N_i, 0, 0$) in time.
- espressopp.analysis.LBOutputVzOfX to output simulation progress and control flux conservation when using MD to LB coupling.

Note: Other types of output classes are possible. It is a subject of user requests.

4.14.11 LBOutputScreen - controls screen output in LB-simulations

Child class derived from the abstract class espressopp.analysis.LBOutput. It computes and outputs to the screen the simulation progress (finished step) and controls mass flux conservation when using MD-to-LB coupling. Ideally, the sum of mass fluxes should be zero, i.e. $j_{LB} + j_{MD} = 0$.

 $\verb|espressopp.analysis.LBOutputScreen| (\textit{system}, \textit{latticeboltzmann})$

Parameters

- **system** system object defined earlier in the python-script
- latticeboltzmann lattice boltzmann object defined earlier in the python-script

Note: this class should be called from external analysis class *espressopp.integrator.ExtAnalyze* with specified periodicity of invokation and after this added to the integrator. See an example for details.

Example to call the profiler:

```
>>> # initialise profiler (for example with the name outputScreen) with system and
>>> # lattice boltzmann objects as parameters:
>>> outputScreen = espressopp.analysis.LBOutputScreen(system,lb)
>>>
>>> # initialise external analysis object (for example extAnalysisNum1) with
>>> # previously created profiler and periodicity of invocation in steps:
>>> extAnalysisNum1=espressopp.integrator.ExtAnalyze(outputScreen,100)
```

```
>>>
>>> # add the external analysis object as an extension to the integrator
>>> integrator.addExtension(extAnalysisNum1)
```

espressopp.analysis.LBOutputScreen (system, latticeboltzmann)

Parameters

- system -
- latticeboltzmann -

4.14.12 LBOutputVzInTime - controls output of the velocity component on a site in time

Child class derived from the abstract class espressopp.analysis.LBOutput. It computes and outputs the velocity component v_z in time on a specific lattice site (the value used at the moment is $0.25 * N_i, 0, 0$).

espressopp.analysis.LBOutputVzInTime (system, latticeboltzmann)

Parameters

- **system** system object defined earlier in the python-script
- latticeboltzmann lattice boltzmann object defined earlier in the python-script

Note: this class should be called from external analysis class *espressopp.integrator.ExtAnalyze* with specified periodicity of invokation and after this added to the integrator. See an example for details.

Example to call the profiler:

espressopp.analysis.LBOutputVzInTime (system, latticeboltzmann)

Parameters

- system -
- latticeboltzmann -

4.14.13 LBOutputVzOfX - controls output of the velocity component profile

Child class derived from the abstract class <code>espressopp.analysis.LBOutput</code>. It computes and outputs simulation progress (finished step) and controls flux conservation when using MD to LB coupling.

 $\verb|espressopp.analysis.LBOutputVzOfX| (\textit{system}, \textit{latticeboltzmann})|$

Parameters

- **system** system object defined earlier in the python-script
- latticeboltzmann lattice boltzmann object defined earlier in the python-script

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Note: this class should be called from external analysis class *espressopp.integrator.ExtAnalyze* with specified periodicity of invokation and after this added to the integrator. See an example for details.

Example to call the profiler:

```
>>> # initialise profiler (for example with the name outputVzOfX) with system and
>>> # lattice boltzmann objects as parameters:
>>> outputVzOfX = espressopp.analysis.LBOutputVzOfX(system,lb)
>>>
>>> # initialise external analysis object (for example extAnalysisNum3) with
>>> # previously created profiler and periodicity of invocation in steps:
>>> extAnalysisNum3=espressopp.integrator.ExtAnalyze(outputVzOfX,100)
>>>
>>> # add the external analysis object as an extension to the integrator
>>> integrator.addExtension(extAnalysisNum3)
```

espressopp.analysis.LBOutputVzOfX(system, latticeboltzmann)

Parameters

- system -
- latticeboltzmann -

4.14.14 espressopp.analysis.MaxPID

```
espressopp.analysis.MaxPID(system)

Parameters system -
```

4.14.15 espressopp.analysis.MeanSquareDispl

```
espressopp.analysis.MeanSquareDispl(system, chainlength)
```

Parameters

- system -
- chainlength (default: None)

espressopp.analysis.MeanSquareDispl.computeG2()

Return type

```
espressopp.analysis.MeanSquareDispl.computeG3()
```

Return type

```
espressopp.analysis.MeanSquareDispl.strange()
```

Return type

4.14.16 espressopp.analysis.NPart

```
espressopp.analysis.NPart(system)
```

Parameters system -

4.14.17 espressopp.analysis.NeighborFluctuation

espressopp.analysis.NeighborFluctuation(system, radius)

Parameters

- system -
- radius -

4.14.18 espressopp.analysis.Observable

```
espressopp.analysis.Observable.compute()
```

Return type

4.14.19 espressopp.analysis.OrderParameter

```
espressopp.analysis.OrderParameter (system, cutoff, angular_momentum, do_cluster_analysis, include_surface_particles, ql\_low, ql\_high)
```

Parameters

- system -
- cutoff -
- angular_momentum(int)-(default: 6)
- do_cluster_analysis (default: False)
- include_surface_particles (default: False)
- **ql_low** (default: -1.0)
- **ql_high** (real) (default: 1.0)

4.14.20 espressopp.analysis.ParticleRadiusDistribution

```
espressopp.analysis.ParticleRadiusDistribution(system)
```

Parameters system -

4.14.21 espressopp.analysis.PotentialEnergy

The object that computes potential energy of different interactions.

```
espressopp.analysis.PotentialEnergy (system, potential, compute_method=None)
```

Parameters

- system (espressopp.System) The system object
- interaction (espressopp.interaction.Interaction) The interaction object.
- **compute_method** (str) If set to ALL (default) then compute total potential energies, if set to CG then compute only coarse-grained part (if feasible), if set to AT then compute only atomitic part of potential energy.

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4.14.22 espressopp.analysis.Pressure

```
espressopp.analysis.Pressure(system)

Parameters system -
```

4.14.23 PressureTensor - Analysis

This class computes the pressure tensor of the system. It can be used as standalone class in python as well as in combination with the integrator extension ExtAnalyze.

Standalone Usage:

```
>>> pt = espressopp.analysis.PressureTensor(system)
>>> print "pressure tensor of current configuration = ", pt.compute()
```

or

```
>>> pt = espressopp.analysis.PressureTensor(system)
>>> for k in xrange(100):
>>> integrator.run(100)
>>> pt.performMeasurement()
>>> print "average pressure tensor = ", pt.getAverageValue()
```

Usage in integrator with ExtAnalyze:

The following methods are supported:

- performMeasurement() computes the pressure tensor and updates average and standard deviation
- reset() resets average and standard deviation to 0
- compute() computes the instant pressure tensor, return value: [xx, yy, zz, xy, xz, yz]
- **getAverageValue()** returns the average pressure tensor and the standard deviation, return value: [xx, yy, zz, xy, xz, yz, +-xx, +-yy, +-zz, +-xy, +-xz, +-yz]
- **getNumberOfMeasurements()** counts the number of measurements that have been computed (standalone or in integrator) does _not_ include measurements that have been done using "compute()"

```
espressopp.analysis.PressureTensor(system)
```

Parameters system -

4.14.24 PressureTensorLayer - Analysis

This class computes the pressure tensor of the system in layer h0. It can be used as standalone class in python as well as in combination with the integrator extension ExtAnalyze.

Standalone Usage:

```
>>> pt = espressopp.analysis.PressureTensorLayer(system, h0, dh)
>>> print "pressure tensor of current configuration = ", pt.compute()
```

or

```
>>> pt = espressopp.analysis.PressureTensorLayer(system)
>>> for k in xrange(100):
>>> integrator.run(100)
>>> pt.performMeasurement()
>>> print "average pressure tensor = ", pt.getAverageValue()
```

Usage in integrator with ExtAnalyze:

The following methods are supported:

- performMeasurement() computes the pressure tensor and updates average and standard deviation
- reset() resets average and standard deviation to 0
- compute() computes the instant pressure tensor in layer h0, return value: [xx, yy, zz, xy, xz, yz]
- **getAverageValue()** returns the average pressure tensor and the standard deviation, return value: [xx, yy, zz, xy, xz, yz, +-xx, +-yy, +-zz, +-xy, +-xz, +-yz]
- **getNumberOfMeasurements**() counts the number of measurements that have been computed (standalone or in integrator) does _not_ include measurements that have been done using "compute()"

espressopp.analysis.**PressureTensorLayer** (system, h0, dh)

Parameters

- system -
- h0 -
- dh –

4.14.25 PressureTensorMultiLayer - Analysis

This class computes the pressure tensor of the system in n layers. Layers are perpendicular to Z direction and are equidistant(distance is Lz/n). It can be used as standalone class in python as well as in combination with the integrator extension ExtAnalyze.

Standalone Usage:

```
>>> pt = espressopp.analysis.PressureTensorMultiLayer(system, n, dh)
>>> for i in xrange(n):
>>> print "pressure tensor in layer %d: %s" % ( i, pt.compute())
```

or

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```
>>> pt = espressopp.analysis.PressureTensorMultiLayer(system, n, dh)
>>> for k in xrange(100):
>>> integrator.run(100)
>>> pt.performMeasurement()
>>> for i in xrange(n):
>>> print "average pressure tensor in layer %d: %s" % ( i, pt.compute())
```

Usage in integrator with ExtAnalyze:

The following methods are supported:

- performMeasurement() computes the pressure tensor and updates average and standard deviation
- reset() resets average and standard deviation to 0
- compute() computes the instant pressure tensor in n layers, return value: [xx, yy, zz, xy, xz, yz]
- **getAverageValue()** returns the average pressure tensor and the standard deviation, return value: [xx, yy, zz, xy, xz, yz, +-xx, +-yy, +-zz, +-xy, +-xz, +-yz]
- **getNumberOfMeasurements()** counts the number of measurements that have been computed (standalone or in integrator) does _not_ include measurements that have been done using "compute()"

espressopp.analysis.PressureTensorMultiLayer(system, n, dh)

Parameters

- system -
- n –
- dh -

4.14.26 espressopp.analysis.RDFatomistic

Class to compute radial distribution functions in adaptive resolution simulations in subregions of the box.

Option 1 (spanbased = True): the RDF can be calculated in a cuboid region in the center of the box (periodic in y,z, limited in x). In this case, particle pairs are considered for which at least one of them is in the defined cuboid region. This can be useful when the high resolution region has a slab geometry. No further normalization should be required.

Option 2 (spanbased = False): the routine can also calculate unnormalized RDFs using particle pairs with both particles being in the high resolution region (based on the resolution value lambda, the span parameter is not used then). This can be useful when atomistic region has complicated or spherical geometries.

In any case, only pairs of atomistic particles belonging to two different coarse-grained particles are considered. Furthermore, note that the routine uses L_y / half (L_y is the box length in y-direction) as the maximum distance for the RDF calculation, which is then binned according to rdfN during the computation. Hence, L_y should be the shortest box side (or, equally short as L_x and/or L_z).

Examples:

```
>>> rdf_0_1 = espressopp.analysis.RDFatomistic(system = system, type1 = 0, type2 = \( \to 1\), spanbased = True, span = 1.5)
>>> # creates the class for calculating the RDF between atomistic particles of \( \to type 1 \) and 0 between different molecules,
>>> # At least one of these particles has to be within plus/minus 1.5 from the \( \to center of the box in x-direction \)
```

```
>>> rdf_0_1.compute(100)
>>> # computes the rdf using 100 bins over a distance corresponding to L_y / 2.0
```

espressopp.analysis.RDFatomistic(system, type1, type2, spanbased, span)

Parameters

- **system** (shared_ptr<System>) system object
- type1 (int) type of atom 1
- **type2** (*int*) type of atom 2
- **spanbased** (bool) (default: True) If True, calculates RDFs in a cuboid region of radius span from the center (limited in x, periodic in y,z). If False, calculates RDFs with both particles being in the high resolution region (using lambda resolution values and ignoring span parameter).
- **span** (*real*) (default: 1.0) +/- distance from centre of box in x-direction of the cuboid region used for RDF calculation if spanbased == True. If spanbased == False, this parameter is not used.

```
espressopp.analysis.RDFatomistic.compute(rdfN)
```

Parameters rdfN (int) - number of bins

Return type list of reals

4.14.27 espressopp.analysis.RadialDistrF

```
espressopp.analysis.RadialDistrF(system)
Parameters system -
```

espressopp.analysis.RadialDistrF.compute(rdfN)

Parameters rdfN-

Return type

4.14.28 espressopp.analysis.StaticStructF

```
espressopp.analysis.StaticStructF(system)
```

Parameters system -

 $\verb|espressopp.analysis.StaticStructF.compute| (\textit{nqx}, \textit{nqy}, \textit{nqz}, \textit{bin_factor}, \textit{ofile})|$

Parameters

- nqx -
- nqy -
- nqz –
- bin_factor -
- ofile (default: None)

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Return type

espressopp.analysis.StaticStructF.computeSingleChain(nqx, nqy, nqz, bin_factor, chainlength, ofile)

Parameters

- nqx -
- nqy -
- nqz -
- bin factor -
- · chainlength -
- ofile (default: None)

Return type

4.14.29 espressopp.analysis.SystemMonitor

SystemMonitor prints and logs to file values obtained from Observables like temperature, pressure or potential energy.

espressopp.analysis.SystemMonitor(system, integrator, output)

Parameters

- system (espressopp.System) The system object.
- integrator (espressopp.integrator.MDIntegrator) The MD integrator
- output (espressopp.analysis.SystemMonitorOutputCSV) The output object.

espressopp.analysis.SystemMonitor.add_observable(name, observable, is_visible)
The function adds new observable to SystemMonitor.

Parameters

- name (str) The name of observable
- observable The observable, eg. espressopp.analysis.PotentialEnergy
- **is_visible** (bool) If set to True then values will be print on console.

espressopp.analysis.SystemMonitor.info()

The method print out on console the values of observables.

CSV Output

The output of SystemMonitor to CSV files.

 $\verb|espressopp.analysis.SystemMonitorOutputCSV| (file_name, delimiter)$

Parameters

- file name (str) The name of CSV file.
- **delimiter** (*str*) The field delimiter, by default it is tabulator.

Example

4.14.30 espressopp.analysis.PotentialEnergy

The object that computes potential energy of different interactions.

espressopp.analysis.PotentialEnergy (system, potential, compute_method=None)

Parameters

- system (espressopp.System) The system object
- interaction (espressopp.interaction.Interaction) The interaction object.
- **compute_method** (str) If set to ALL (default) then compute total potential energies, if set to CG then compute only coarse-grained part (if feasible), if set to AT then compute only atomitic part of potential energy.

4.14.31 espressopp.analysis.Temperature

```
espressopp.analysis.Temperature(system)
Parameters system -
```

4.14.32 espressopp.analysis.Test

```
espressopp.analysis.Test(system)

Parameters system –
```

4.14.33 espressopp.analysis.TotalVelocity

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Examples

Reset the velocity

```
>>> total_velocity = espressopp.analysis.TotalVelocity(system)
>>> total_velocity.reset()
```

Extension to integrator

This extension can also be attached to integrator and run *reset()* every *n-th* steps.

```
>>> total_velocity = espressopp.analysis.TotalVelocity(system)
>>> ext_remove_com = espressopp.analysis.ExtAnalyze(total_velocity, 10)
>>> integrator.addExtension(ext_remove_com)
```

4.14.34 espressopp.analysis. Velocities

4.14.35 espressopp.analysis.VelocityAutocorrelation

```
espressopp.analysis.VelocityAutocorrelation(system)
```

Parameters system-

4.14.36 espressopp.analysis.Viscosity

```
espressopp.analysis.Viscosity(system)

Parameters system -
espressopp.analysis.Viscosity.compute(t0, dt, T)
```

Parameters

- t0 -
- dt -
- T -

Return type

```
espressopp.analysis.Viscosity.gather()
```

Return type

4.14.37 espressopp.analysis.XDensity

```
espressopp.analysis.XDensity(system)

Parameters system -
espressopp.analysis.XDensity.compute(rdfN)

Parameters rdfN -
Return type
```

4.14.38 espressopp.analysis.XPressure

```
espressopp.analysis.XPressure(system)
    Parameters system -
espressopp.analysis.XPressure.compute(N)
    Parameters N -
    Return type
```

4.15 bc

4.16 check

4.16.1 espressopp.check.System

4.17 esutil

4.17.1 espressopp.esutil.GammaVariate

```
espressopp.esutil.GammaVariate (alpha, beta)
Parameters
• alpha -
```

- u-p---
- beta -

4.17.2 espressopp.esutil.Grid

4.17.3 espressopp.esutil.NormalVariate

```
\verb|espressopp.esutil.NormalVariate| (\textit{mean}, \textit{sigma})|
```

Parameters

- **mean** (real) (default: 0.0)
- **sigma** (real) (default: 1.0)

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4.17.4 espressopp.esutil.RNG

4.17.5 espressopp.esutil.UniformOnSphere

4.17.6 espressopp.esutil.collectives

espressopp.esutil.locateItem(here)

Parameters here -

espressopp.esutil.collectives.locateItem(here)

locate the node with here=True (e.g. indicating that data of a distributed storage is on the local node). This is a collective SPMD function.

here is a boolean value, which should be True on at most one node. Returns on the controller the number of the node with here=True, or an KeyError exception if no node had the item, i.e. had here=True.

4.18 external

Homogeneous Transformation Matrices and Quaternions.

A library for calculating 4x4 matrices for translating, rotating, reflecting, scaling, shearing, projecting, orthogonalizing, and superimposing arrays of 3D homogeneous coordinates as well as for converting between rotation matrices, Euler angles, and quaternions. Also includes an Arcball control object and functions to decompose transformation matrices.

Authors Christoph Gohlke, Laboratory for Fluorescence Dynamics, University of California, Irvine

Version 2011.01.25

4.18.1 Requirements

- Python 2.6 or 3.1
- Numpy 1.5
- transformations.c 2010.04.10 (optional implementation of some functions in C)

4.18.2 Notes

The API is not stable yet and is expected to change between revisions.

This Python code is not optimized for speed. Refer to the transformations.c module for a faster implementation of some functions.

Documentation in HTML format can be generated with epydoc.

Matrices (M) can be inverted using numpy.linalg.inv(M), concatenated using numpy.dot(M0, M1), or used to transform homogeneous coordinates (v) using numpy.dot(M, v) for shape (4, *) "point of arrays", respectively numpy.dot(v, M.T) for shape (*, 4) "array of points".

Use the transpose of transformation matrices for OpenGL glMultMatrixd().

Calculations are carried out with numpy.float64 precision.

Vector, point, quaternion, and matrix function arguments are expected to be "array like", i.e. tuple, list, or numpy arrays.

Return types are numpy arrays unless specified otherwise.

Angles are in radians unless specified otherwise.

Quaternions w+ix+jy+kz are represented as [w, x, y, z].

A triple of Euler angles can be applied/interpreted in 24 ways, which can be specified using a 4 character string or encoded 4-tuple:

Axes 4-string: e.g. 'sxyz' or 'ryxy'

- first character: rotations are applied to 's'tatic or 'r'otating frame
- remaining characters : successive rotation axis 'x', 'y', or 'z'

Axes 4-tuple: e.g. (0, 0, 0, 0) or (1, 1, 1, 1)

- inner axis: code of axis ('x':0, 'y':1, 'z':2) of rightmost matrix.
- parity: even (0) if inner axis 'x' is followed by 'y', 'y' is followed by 'z', or 'z' is followed by 'x'. Otherwise odd (1).
- repetition: first and last axis are same (1) or different (0).
- frame : rotations are applied to static (0) or rotating (1) frame.

4.18.3 References

- 1. Matrices and transformations. Ronald Goldman. In "Graphics Gems I", pp 472-475. Morgan Kaufmann, 1990.
- 2. More matrices and transformations: shear and pseudo-perspective. Ronald Goldman. In "Graphics Gems II", pp 320-323. Morgan Kaufmann, 1991.
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4.18.4 Examples

```
>>> alpha, beta, gamma = 0.123, -1.234, 2.345
>>> origin, xaxis, yaxis, zaxis = (0, 0, 0), (1, 0, 0), (0, 1, 0), (0, 0, 1)
>>> I = identity_matrix()
>>> Rx = rotation_matrix(alpha, xaxis)
>>> Ry = rotation_matrix(beta, yaxis)
>>> Rz = rotation_matrix(gamma, zaxis)
>>> R = concatenate_matrices(Rx, Ry, Rz)
```

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```
>>> euler = euler_from_matrix(R, 'rxyz')
>>> numpy.allclose([alpha, beta, gamma], euler)
>>> Re = euler_matrix(alpha, beta, gamma, 'rxyz')
>>> is_same_transform(R, Re)
True
>>> al, be, ga = euler_from_matrix(Re, 'rxyz')
>>> is_same_transform(Re, euler_matrix(al, be, ga, 'rxyz'))
True
>>> qx = quaternion_about_axis(alpha, xaxis)
>>> qy = quaternion_about_axis(beta, yaxis)
>>> qz = quaternion_about_axis(gamma, zaxis)
>>> q = quaternion_multiply(qx, qy)
>>> q = quaternion_multiply(q, qz)
>>> Rq = quaternion_matrix(q)
>>> is_same_transform(R, Rq)
True
>>> S = scale_matrix(1.23, origin)
>>> T = translation_matrix((1, 2, 3))
>>> Z = shear_matrix(beta, xaxis, origin, zaxis)
>>> R = random_rotation_matrix(numpy.random.rand(3))
>>> M = concatenate_matrices(T, R, Z, S)
>>> scale, shear, angles, trans, persp = decompose_matrix(M)
>>> numpy.allclose(scale, 1.23)
True
>>> numpy.allclose(trans, (1, 2, 3))
True
>>> numpy.allclose(shear, (0, math.tan(beta), 0))
>>> is_same_transform(R, euler_matrix(axes='sxyz', *angles))
True
>>> M1 = compose_matrix(scale, shear, angles, trans, persp)
>>> is_same_transform(M, M1)
>>> v0, v1 = random_vector(3), random_vector(3)
>>> M = rotation_matrix(angle_between_vectors(v0, v1), vector_product(v0, v1))
>>> v2 = numpy.dot(v0, M[:3,:3].T)
>>> numpy.allclose(unit_vector(v1), unit_vector(v2))
True
```

class espressopp.external.transformations.Arcball(initial=None) Virtual Trackball Control.

```
>>> ball = Arcball()
>>> ball = Arcball(initial=numpy.identity(4))
>>> ball.place([320, 320], 320)
>>> ball.down([500, 250])
>>> ball.drag([475, 275])
>>> R = ball.matrix()
>>> numpy.allclose(numpy.sum(R), 3.90583455)
True
>>> ball = Arcball(initial=[1, 0, 0, 0])
>>> ball.place([320, 320], 320)
>>> ball.setaxes([1,1,0], [-1, 1, 0])
>>> ball.setconstrain(True)
>>> ball.down([400, 200])
>>> ball.drag([200, 400])
>>> R = ball.matrix()
>>> numpy.allclose(numpy.sum(R), 0.2055924)
True
>>> ball.next()
```

```
down (point)
          Set initial cursor window coordinates and pick constrain-axis.
          Update current cursor window coordinates.
     getconstrain()
          Return state of constrain to axis mode.
     matrix()
          Return homogeneous rotation matrix.
     next (acceleration=0.0)
          Continue rotation in direction of last drag.
     place (center, radius)
          Place Arcball, e.g. when window size changes.
          center [sequence[2]] Window coordinates of trackball center.
          radius [float] Radius of trackball in window coordinates.
     setaxes(*axes)
          Set axes to constrain rotations.
     setconstrain (constrain)
          Set state of constrain to axis mode.
espressopp.external.transformations.angle_between_vectors (v\theta,
                                                                                  v1,
                                                                                          di-
                                                                           rected=True,
                                                                           axis=0)
     Return angle between vectors.
     If directed is False, the input vectors are interpreted as undirected axes, i.e. the maximum angle is pi/2.
     >>> a = angle_between_vectors([1, -2, 3], [-1, 2, -3])
     >>> numpy.allclose(a, math.pi)
     \rightarrow \rightarrow a = angle_between_vectors([1, -2, 3], [-1, 2, -3], directed=False)
     >>> numpy.allclose(a, 0)
     True
     >>> v0 = [[2, 0, 0, 2], [0, 2, 0, 2], [0, 0, 2, 2]]
     >>> v1 = [[3], [0], [0]]
     >>> a = angle_between_vectors(v0, v1)
     >>> numpy.allclose(a, [0., 1.5708, 1.5708, 0.95532])
     True
     >>> v0 = [[2, 0, 0], [2, 0, 0], [0, 2, 0], [2, 0, 0]]
     >>> v1 = [[0, 3, 0], [0, 0, 3], [0, 0, 3], [3, 3, 3]]
     >>> a = angle_between_vectors(v0, v1, axis=1)
     >>> numpy.allclose(a, [1.5708, 1.5708, 1.5708, 0.95532])
     True
espressopp.external.transformations.arcball_constrain_to_axis(point, axis)
     Return sphere point perpendicular to axis.
espressopp.external.transformations.arcball_map_to_sphere(point, center, ra-
                                                                           dius)
     Return unit sphere coordinates from window coordinates.
espressopp.external.transformations.arcball_nearest_axis (point, axes)
     Return axis, which arc is nearest to point.
espressopp.external.transformations.clip_matrix(left, right, bottom, top, near, far,
                                                              perspective=False)
     Return matrix to obtain normalized device coordinates from frustrum.
     The frustrum bounds are axis-aligned along x (left, right), y (bottom, top) and z (near, far).
```

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Normalized device coordinates are in range [-1, 1] if coordinates are inside the frustrum.

If perspective is True the frustrum is a truncated pyramid with the perspective point at origin and direction along z axis, otherwise an orthographic canonical view volume (a box).

Homogeneous coordinates transformed by the perspective clip matrix need to be dehomogenized (devided by w coordinate).

```
>>> frustrum = numpy.random.rand(6)
>>> frustrum[1] += frustrum[0]
>>> frustrum[3] += frustrum[2]
>>> frustrum[5] += frustrum[4]
>>> M = clip_matrix(perspective=False, *frustrum)
>>> numpy.dot(M, [frustrum[0], frustrum[2], frustrum[4], 1.0])
array([-1., -1., -1., 1.])
>>> numpy.dot(M, [frustrum[1], frustrum[3], frustrum[5], 1.0])
array([ 1., 1., 1., 1.])
>>> M = clip_matrix(perspective=True, *frustrum)
>>> v = numpy.dot(M, [frustrum[0], frustrum[2], frustrum[4], 1.0])
>>> v / v[3]
array([-1., -1., -1., 1.])
>>> v = numpy.dot(M, [frustrum[1], frustrum[3], frustrum[4], 1.0])
>>> v / v[3]
array([ 1., 1., -1., 1.])
```

espressopp.external.transformations.compose_matrix (scale=None, shear=None, angles=None, translate=None, perspective=None)

Return transformation matrix from sequence of transformations.

This is the inverse of the decompose_matrix function.

Sequence of transformations: scale: vector of 3 scaling factors shear: list of shear factors for x-y, x-z, y-z axes angles: list of Euler angles about static x, y, z axes translate: translation vector along x, y, z axes perspective: perspective partition of matrix

```
>>> scale = numpy.random.random(3) - 0.5
>>> shear = numpy.random.random(3) - 0.5
>>> angles = (numpy.random.random(3) - 0.5) * (2*math.pi)
>>> trans = numpy.random.random(3) - 0.5
>>> persp = numpy.random.random(4) - 0.5
>>> M0 = compose_matrix(scale, shear, angles, trans, persp)
>>> result = decompose_matrix(M0)
>>> M1 = compose_matrix(*result)
>>> is_same_transform(M0, M1)
True
```

espressopp.external.transformations.concatenate_matrices(*matrices)

Return concatenation of series of transformation matrices.

```
>>> M = numpy.random.rand(16).reshape((4, 4)) - 0.5
>>> numpy.allclose(M, concatenate_matrices(M))
True
>>> numpy.allclose(numpy.dot(M, M.T), concatenate_matrices(M, M.T))
True
```

espressopp.external.transformations.decompose_matrix(matrix)

Return sequence of transformations from transformation matrix.

matrix [array_like] Non-degenerative homogeneous transformation matrix

Return tuple of: scale: vector of 3 scaling factors shear: list of shear factors for x-y, x-z, y-z axes angles: list of Euler angles about static x, y, z axes translate: translation vector along x, y, z axes perspective: perspective partition of matrix

Raise ValueError if matrix is of wrong type or degenerative.

```
>>> T0 = translation_matrix((1, 2, 3))
>>> scale, shear, angles, trans, persp = decompose_matrix(T0)
>>> T1 = translation_matrix(trans)
>>> numpy.allclose(T0, T1)
True
>>> S = scale_matrix(0.123)
>>> scale, shear, angles, trans, persp = decompose_matrix(S)
>>> scale[0]
0.123
>>> R0 = euler_matrix(1, 2, 3)
>>> scale, shear, angles, trans, persp = decompose_matrix(R0)
>>> R1 = euler_matrix(*angles)
>>> numpy.allclose(R0, R1)
True
```

espressopp.external.transformations.euler_from_matrix(matrix, axes='sxyz')

Return Euler angles from rotation matrix for specified axis sequence.

axes: One of 24 axis sequences as string or encoded tuple

Note that many Euler angle triplets can describe one matrix.

```
>>> R0 = euler_matrix(1, 2, 3, 'syxz')
>>> al, be, ga = euler_from_matrix(R0, 'syxz')
>>> R1 = euler_matrix(al, be, ga, 'syxz')
>>> numpy.allclose(R0, R1)
True
>>> angles = (4.0*math.pi) * (numpy.random.random(3) - 0.5)
>>> for axes in _AXES2TUPLE.keys():
... R0 = euler_matrix(axes=axes, *angles)
... R1 = euler_matrix(axes=axes, *euler_from_matrix(R0, axes))
... if not numpy.allclose(R0, R1): print(axes, "failed")
```

espressopp.external.transformations.euler_from_quaternion(quaternion, axes='sxyz')

Return Euler angles from quaternion for specified axis sequence.

```
>>> angles = euler_from_quaternion([0.99810947, 0.06146124, 0, 0])
>>> numpy.allclose(angles, [0.123, 0, 0])
True
```

espressopp.external.transformations.euler_matrix(ai, aj, ak, axes='sxyz')

Return homogeneous rotation matrix from Euler angles and axis sequence.

ai, aj, ak: Euler's roll, pitch and yaw angles axes: One of 24 axis sequences as string or encoded tuple

espressopp.external.transformations.identity_matrix()

Return 4x4 identity/unit matrix.

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```
>>> I = identity_matrix()
>>> numpy.allclose(I, numpy.dot(I, I))
True
>>> numpy.sum(I), numpy.trace(I)
(4.0, 4.0)
>>> numpy.allclose(I, numpy.identity(4, dtype=numpy.float64))
True
```

espressopp.external.transformations.inverse_matrix(matrix)

Return inverse of square transformation matrix.

espressopp.external.transformations.is_same_transform(matrix0, matrix1)
Return True if two matrices perform same transformation.

```
>>> is_same_transform(numpy.identity(4), numpy.identity(4))
True
>>> is_same_transform(numpy.identity(4), random_rotation_matrix())
False
```

espressopp.external.transformations.orthogonalization_matrix(lengths, angles)

Return orthogonalization matrix for crystallographic cell coordinates.

Angles are expected in degrees.

The de-orthogonalization matrix is the inverse.

```
>>> O = orthogonalization_matrix((10., 10., 10.), (90., 90., 90.))
>>> numpy.allclose(O[:3, :3], numpy.identity(3, float) * 10)
True
>>> O = orthogonalization_matrix([9.8, 12.0, 15.5], [87.2, 80.7, 69.7])
>>> numpy.allclose(numpy.sum(0), 43.063229)
True
```

espressopp.external.transformations.projection_from_matrix(matrix, pseudo=False)

Return projection plane and perspective point from projection matrix.

Return values are same as arguments for projection_matrix function: point, normal, direction, perspective, and pseudo.

```
>>> point = numpy.random.random(3) - 0.5
>>> normal = numpy.random.random(3) - 0.5
>>> direct = numpy.random.random(3) - 0.5
>>> persp = numpy.random.random(3) - 0.5
>>> P0 = projection_matrix(point, normal)
>>> result = projection_from_matrix(P0)
>>> P1 = projection_matrix(*result)
>>> is_same_transform(P0, P1)
True
>>> P0 = projection_matrix(point, normal, direct)
>>> result = projection_from_matrix(P0)
>>> P1 = projection_matrix(*result)
>>> result = projection_matrix(*result)
```

```
True
>>> P0 = projection_matrix(point, normal, perspective=persp, pseudo=False)
>>> result = projection_from_matrix(P0, pseudo=False)
>>> P1 = projection_matrix(*result)
>>> is_same_transform(P0, P1)
True
>>> P0 = projection_matrix(point, normal, perspective=persp, pseudo=True)
>>> result = projection_from_matrix(P0, pseudo=True)
>>> P1 = projection_matrix(*result)
>>> is_same_transform(P0, P1)
True
```

```
espressopp.external.transformations.projection_matrix(point, normal, direction=None, perspective=None, pseudo=False)
```

Return matrix to project onto plane defined by point and normal.

Using either perspective point, projection direction, or none of both.

If pseudo is True, perspective projections will preserve relative depth such that Perspective = dot(Orthogonal, PseudoPerspective).

```
>>> P = projection_matrix((0, 0, 0), (1, 0, 0))
>>> numpy.allclose(P[1:, 1:], numpy.identity(4)[1:, 1:])
True
>>> point = numpy.random.random(3) - 0.5
>>> normal = numpy.random.random(3) - 0.5
>>> direct = numpy.random.random(3) - 0.5
>>> persp = numpy.random.random(3) - 0.5
>>> P0 = projection_matrix(point, normal)
>>> P1 = projection_matrix(point, normal, direction=direct)
>>> P2 = projection_matrix(point, normal, perspective=persp)
>>> P3 = projection_matrix(point, normal, perspective=persp, pseudo=True)
>>> is_same_transform(P2, numpy.dot(P0, P3))
True
>>> P = projection_matrix((3, 0, 0), (1, 1, 0), (1, 0, 0))
>>> v0 = (numpy.random.rand(4, 5) - 0.5) * 20.0
>>> v0[3] = 1.0
>>> v1 = numpy.dot(P, v0)
>>> numpy.allclose(v1[1], v0[1])
True
>>> numpy.allclose(v1[0], 3.0-v1[1])
```

espressopp.external.transformations.quaternion_about_axis (angle, axis)
Return quaternion for rotation about axis.

```
>>> q = quaternion_about_axis(0.123, (1, 0, 0))
>>> numpy.allclose(q, [0.99810947, 0.06146124, 0, 0])
True
```

espressopp.external.transformations.quaternion_conjugate(quaternion)

Return conjugate of quaternion.

```
>>> q0 = random_quaternion()
>>> q1 = quaternion_conjugate(q0)
>>> q1[0] == q0[0] and all(q1[1:] == -q0[1:])
True
```

```
espressopp.external.transformations.quaternion_from_euler(ai, aj, ak, axes='sxyz')
```

Return quaternion from Euler angles and axis sequence.

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ai, aj, ak: Euler's roll, pitch and yaw angles axes: One of 24 axis sequences as string or encoded tuple

```
>>> q = quaternion_from_euler(1, 2, 3, 'ryxz')
>>> numpy.allclose(q, [0.435953, 0.310622, -0.718287, 0.444435])
True
```

espressopp.external.transformations.quaternion_from_matrix(matrix, ispre-cise=False)

Return quaternion from rotation matrix.

If isprecise=True, the input matrix is assumed to be a precise rotation matrix and a faster algorithm is used.

```
>>> q = quaternion_from_matrix(identity_matrix(), True)
>>> numpy.allclose(q, [1., 0., 0., 0.])
\rightarrow \rightarrow q = quaternion_from_matrix(numpy.diag([1., -1., -1., 1.]))
>>> numpy.allclose(q, [0, 1, 0, 0]) or numpy.allclose(q, [0, -1, 0, 0])
>>> R = rotation_matrix(0.123, (1, 2, 3))
>>> q = quaternion_from_matrix(R, True)
>>> numpy.allclose(q, [0.9981095, 0.0164262, 0.0328524, 0.0492786])
>>> R = [[-0.545, 0.797, 0.260, 0], [0.733, 0.603, -0.313, 0],
        [-0.407, 0.021, -0.913, 0], [0, 0, 0, 1]]
>>> q = quaternion_from_matrix(R)
>>> numpy.allclose(q, [0.19069, 0.43736, 0.87485, -0.083611])
>>> R = [[0.395, 0.362, 0.843, 0], [-0.626, 0.796, -0.056, 0],
        [-0.677, -0.498, 0.529, 0], [0, 0, 0, 1]]
>>> q = quaternion_from_matrix(R)
>>> numpy.allclose(q, [0.82336615, -0.13610694, 0.46344705, -0.29792603])
True
>>> R = random_rotation_matrix()
>>> q = quaternion_from_matrix(R)
>>> is_same_transform(R, quaternion_matrix(q))
True
```

 $\verb|espressopp.external.transformations.quaternion_imag(|quaternion|)|$

Return imaginary part of quaternion.

```
>>> quaternion_imag([3.0, 0.0, 1.0, 2.0])
[0.0, 1.0, 2.0]
```

espressopp.external.transformations.quaternion_inverse(quaternion)
Return inverse of quaternion.

```
>>> q0 = random_quaternion()
>>> q1 = quaternion_inverse(q0)
>>> numpy.allclose(quaternion_multiply(q0, q1), [1, 0, 0, 0])
True
```

espressopp.external.transformations.quaternion_matrix(quaternion)
Return homogeneous rotation matrix from quaternion.

```
>>> M = quaternion_matrix([0.99810947, 0.06146124, 0, 0])
>>> numpy.allclose(M, rotation_matrix(0.123, (1, 0, 0)))
True
>>> M = quaternion_matrix([1, 0, 0, 0])
>>> numpy.allclose(M, identity_matrix())
True
>>> M = quaternion_matrix([0, 1, 0, 0])
>>> numpy.allclose(M, numpy.diag([1, -1, -1, 1]))
True
```

espressopp.external.transformations.quaternion_multiply(quaternion1, quaternion0)

Return multiplication of two quaternions.

```
>>> q = quaternion_multiply([4, 1, -2, 3], [8, -5, 6, 7])
>>> numpy.allclose(q, [28, -44, -14, 48])
True
```

espressopp.external.transformations.quaternion_real(quaternion)
Return real part of quaternion.

```
>>> quaternion_real([3.0, 0.0, 1.0, 2.0])
3.0
```

espressopp.external.transformations.quaternion_slerp(quat0, quat1, fraction, spin=0, shortest-path=True)

Return spherical linear interpolation between two quaternions.

espressopp.external.transformations.random_quaternion(rand=None)
Return uniform random unit quaternion.

rand: array like or None Three independent random variables that are uniformly distributed between 0 and 1

```
>>> q = random_quaternion()
>>> numpy.allclose(1.0, vector_norm(q))
True
>>> q = random_quaternion(numpy.random.random(3))
>>> len(q.shape), q.shape[0]==4
(1, True)
```

espressopp.external.transformations.random_rotation_matrix(rand=None)
Return uniform random rotation matrix.

rnd: array like Three independent random variables that are uniformly distributed between 0 and 1 for each returned quaternion.

```
>>> R = random_rotation_matrix()
>>> numpy.allclose(numpy.dot(R.T, R), numpy.identity(4))
True
```

espressopp.external.transformations.random_vector(size)

Return array of random doubles in the half-open interval [0.0, 1.0).

```
>>> v = random_vector(10000)
>>> numpy.all(v >= 0.0) and numpy.all(v < 1.0)
True
>>> v0 = random_vector(10)
```

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```
>>> v1 = random_vector(10)
>>> numpy.any(v0 == v1)
False
```

espressopp.external.transformations.reflection_from_matrix(matrix)

Return mirror plane point and normal vector from reflection matrix.

```
>>> v0 = numpy.random.random(3) - 0.5
>>> v1 = numpy.random.random(3) - 0.5
>>> M0 = reflection_matrix(v0, v1)
>>> point, normal = reflection_from_matrix(M0)
>>> M1 = reflection_matrix(point, normal)
>>> is_same_transform(M0, M1)
True
```

espressopp.external.transformations.reflection_matrix(point, normal)

Return matrix to mirror at plane defined by point and normal vector.

```
>>> v0 = numpy.random.random(4) - 0.5
>>> v0[3] = 1.0
>>> v1 = numpy.random.random(3) - 0.5
>>> R = reflection_matrix(v0, v1)
>>> numpy.allclose(2., numpy.trace(R))
True
>>> numpy.allclose(v0, numpy.dot(R, v0))
True
>>> v2 = v0.copy()
>>> v2[:3] += v1
>>> v3 = v0.copy()
>>> v2[:3] -= v1
>>> numpy.allclose(v2, numpy.dot(R, v3))
True
```

 $\verb|espressopp.external.transformations.rotation_from_matrix| (\textit{matrix})$

Return rotation angle and axis from rotation matrix.

```
>>> angle = (random.random() - 0.5) * (2*math.pi)
>>> direc = numpy.random.random(3) - 0.5
>>> point = numpy.random.random(3) - 0.5
>>> R0 = rotation_matrix(angle, direc, point)
>>> angle, direc, point = rotation_from_matrix(R0)
>>> R1 = rotation_matrix(angle, direc, point)
>>> is_same_transform(R0, R1)
True
```

espressopp.external.transformations.rotation_matrix(angle, direction, point=None)

Return matrix to rotate about axis defined by point and direction.

```
>>> R = rotation_matrix(math.pi/2.0, [0, 0, 1], [1, 0, 0])
>>> numpy.allclose(numpy.dot(R, [0, 0, 0, 1]), [1., -1., 0., 1.])
True
>>> angle = (random.random() - 0.5) * (2*math.pi)
>>> direc = numpy.random.random(3) - 0.5
>>> point = numpy.random.random(3) - 0.5
>>> R0 = rotation_matrix(angle, direc, point)
>>> R1 = rotation_matrix(angle-2*math.pi, direc, point)
True
>>> R0 = rotation_matrix(angle, direc, point)
>>> R1 = rotation_matrix(angle, direc, point)
>>> R1 = rotation_matrix(angle, direc, point)
>>> R1 = rotation_matrix(-angle, -direc, point)
>>> is_same_transform(R0, R1)
```

```
True
>>> I = numpy.identity(4, numpy.float64)
>>> numpy.allclose(I, rotation_matrix(math.pi*2, direc))
True
>>> numpy.allclose(2., numpy.trace(rotation_matrix(math.pi/2,
... direc, point)))
True
```

espressopp.external.transformations.scale_from_matrix(matrix)

Return scaling factor, origin and direction from scaling matrix.

```
>>> factor = random.random() * 10 - 5
>>> origin = numpy.random.random(3) - 0.5
>>> direct = numpy.random.random(3) - 0.5
>>> S0 = scale_matrix(factor, origin)
>>> factor, origin, direction = scale_from_matrix(S0)
>>> S1 = scale_matrix(factor, origin, direction)
>>> is_same_transform(S0, S1)
True
>>> S0 = scale_matrix(factor, origin, direct)
>>> factor, origin, direction = scale_from_matrix(S0)
>>> S1 = scale_matrix(factor, origin, direction)
>>> is_same_transform(S0, S1)
True
```

espressopp.external.transformations.scale_matrix(factor, origin=None, direction=None)

Return matrix to scale by factor around origin in direction.

Use factor -1 for point symmetry.

```
>>> v = (numpy.random.rand(4, 5) - 0.5) * 20.0
>>> v[3] = 1.0
>>> S = scale_matrix(-1.234)
>>> numpy.allclose(numpy.dot(S, v)[:3], -1.234*v[:3])
True
>>> factor = random.random() * 10 - 5
>>> origin = numpy.random.random(3) - 0.5
>>> direct = numpy.random.random(3) - 0.5
>>> S = scale_matrix(factor, origin)
>>> S = scale_matrix(factor, origin, direct)
```

espressopp.external.transformations.shear_from_matrix(matrix)

Return shear angle, direction and plane from shear matrix.

```
>>> angle = (random.random() - 0.5) * 4*math.pi
>>> direct = numpy.random.random(3) - 0.5
>>> point = numpy.random.random(3) - 0.5
>>> normal = numpy.cross(direct, numpy.random.random(3))
>>> S0 = shear_matrix(angle, direct, point, normal)
>>> angle, direct, point, normal = shear_from_matrix(S0)
>>> S1 = shear_matrix(angle, direct, point, normal)
>>> is_same_transform(S0, S1)
True
```

espressopp.external.transformations.**shear_matrix** (*angle*, *direction*, *point*, *normal*) Return matrix to shear by angle along direction vector on shear plane.

The shear plane is defined by a point and normal vector. The direction vector must be orthogonal to the plane's normal vector.

A point P is transformed by the shear matrix into P" such that the vector P-P" is parallel to the direction vector and its extent is given by the angle of P-P'-P", where P' is the orthogonal projection of P onto the

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shear plane.

```
>>> angle = (random.random() - 0.5) * 4*math.pi
>>> direct = numpy.random.random(3) - 0.5
>>> point = numpy.random.random(3) - 0.5
>>> normal = numpy.cross(direct, numpy.random.random(3))
>>> S = shear_matrix(angle, direct, point, normal)
>>> numpy.allclose(1.0, numpy.linalg.det(S))
True
```

```
espressopp.external.transformations.superimposition_matrix (v0, v1, scaling=False, us-esvd=True)
```

Return matrix to transform given vector set into second vector set.

v0 and v1 are shape (3, *) or (4, *) arrays of at least 3 vectors.

If usesvd is True, the weighted sum of squared deviations (RMSD) is minimized according to the algorithm by W. Kabsch [8]. Otherwise the quaternion based algorithm by B. Horn [9] is used (slower when using this Python implementation).

The returned matrix performs rotation, translation and uniform scaling (if specified).

```
>>> v0 = numpy.random.rand(3, 10)
>>> M = superimposition_matrix(v0, v0)
>>> numpy.allclose(M, numpy.identity(4))
>>> R = random_rotation_matrix(numpy.random.random(3))
>>> v0 = ((1,0,0), (0,1,0), (0,0,1), (1,1,1))
>>> v1 = numpy.dot(R, v0)
>>> M = superimposition_matrix(v0, v1)
>>> numpy.allclose(v1, numpy.dot(M, v0))
>>> v0 = (numpy.random.rand(4, 100) - 0.5) * 20.0
>>> v0[3] = 1.0
>>> v1 = numpy.dot(R, v0)
>>> M = superimposition_matrix(v0, v1)
>>> numpy.allclose(v1, numpy.dot(M, v0))
>>> S = scale_matrix(random.random())
>>> T = translation_matrix(numpy.random.random(3)-0.5)
>>> M = concatenate_matrices(T, R, S)
>>> v1 = numpy.dot(M, v0)
>>> v0[:3] += numpy.random.normal(0.0, 1e-9, 300).reshape(3, -1)
>>> M = superimposition_matrix(v0, v1, scaling=True)
>>> numpy.allclose(v1, numpy.dot(M, v0))
>>> M = superimposition_matrix(v0, v1, scaling=True, usesvd=False)
>>> numpy.allclose(v1, numpy.dot(M, v0))
True
>>> v = numpy.empty((4, 100, 3), dtype=numpy.float64)
>>> v[:, :, 0] = v0
>>> M = superimposition_matrix(v0, v1, scaling=True, usesvd=False)
>>> numpy.allclose(v1, numpy.dot(M, v[:, :, 0]))
```

espressopp.external.transformations.translation_from_matrix(matrix)

Return translation vector from translation matrix.

```
>>> v0 = numpy.random.random(3) - 0.5
>>> v1 = translation_from_matrix(translation_matrix(v0))
>>> numpy.allclose(v0, v1)
True
```

espressopp.external.transformations.translation_matrix (direction) Return matrix to translate by direction vector.

```
>>> v = numpy.random.random(3) - 0.5
>>> numpy.allclose(v, translation_matrix(v)[:3, 3])
True
```

espressopp.external.transformations.unit_vector(data, axis=None, out=None)
Return ndarray normalized by length, i.e. eucledian norm, along axis.

```
>>> v0 = numpy.random.random(3)
>>> v1 = unit_vector(v0)
>>> numpy.allclose(v1, v0 / numpy.linalg.norm(v0))
\rightarrow > v0 = numpy.random.rand(5, 4, 3)
>>> v1 = unit_vector(v0, axis=-1)
>>> v2 = v0 / numpy.expand_dims(numpy.sqrt(numpy.sum(v0*v0, axis=2)), 2)
>>> numpy.allclose(v1, v2)
True
>>> v1 = unit_vector(v0, axis=1)
>>> v2 = v0 / numpy.expand_dims(numpy.sqrt(numpy.sum(v0*v0, axis=1)), 1)
>>> numpy.allclose(v1, v2)
True
>>> v1 = numpy.empty((5, 4, 3), dtype=numpy.float64)
>>> unit_vector(v0, axis=1, out=v1)
>>> numpy.allclose(v1, v2)
True
>>> list(unit_vector([]))
>>> list(unit_vector([1.0]))
[1.0]
```

espressopp.external.transformations.vector_norm(data, axis=None, out=None)
Return length, i.e. eucledian norm, of ndarray along axis.

```
>>> v = numpy.random.random(3)
>>> n = vector_norm(v)
>>> numpy.allclose(n, numpy.linalg.norm(v))
>>> v = numpy.random.rand(6, 5, 3)
>>> n = vector_norm(v, axis=-1)
>>> numpy.allclose(n, numpy.sqrt(numpy.sum(v*v, axis=2)))
>>> n = vector_norm(v, axis=1)
>>> numpy.allclose(n, numpy.sqrt(numpy.sum(v*v, axis=1)))
>>> v = numpy.random.rand(5, 4, 3)
>>> n = numpy.empty((5, 3), dtype=numpy.float64)
>>> vector_norm(v, axis=1, out=n)
>>> numpy.allclose(n, numpy.sqrt(numpy.sum(v*v, axis=1)))
True
>>> vector_norm([])
0.0
>>> vector_norm([1.0])
1.0
```

 $\verb|espressopp.external.transformations.vector_product| (v0, v1, axis=0)$

Return vector perpendicular to vectors.

```
>>> v = vector_product([2, 0, 0], [0, 3, 0])
>>> numpy.allclose(v, [0, 0, 6])
True
>>> v0 = [[2, 0, 0, 2], [0, 2, 0, 2], [0, 0, 2, 2]]
```

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```
>>> v1 = [[3], [0], [0]]
>>> v = vector_product(v0, v1)
>>> numpy.allclose(v, [[0, 0, 0, 0], [0, 0, 6, 6], [0, -6, 0, -6]])
True
>>> v0 = [[2, 0, 0], [2, 0, 0], [0, 2, 0], [2, 0, 0]]
>>> v1 = [[0, 3, 0], [0, 0, 3], [0, 0, 3], [3, 3, 3]]
>>> v = vector_product(v0, v1, axis=1)
>>> numpy.allclose(v, [[0, 0, 6], [0, -6, 0], [6, 0, 0], [0, -6, 6]])
True
```

4.19 integrator

4.19.1 AdResS - Object

The AdResS object is an extension to the integrator. It makes sure that the integrator also processes the atomistic particles and not only the CG particles. Hence, this object is of course only used when performing AdResS or H-AdResS simulations.

In detail the AdResS extension makes sure:

- that also the forces on the atomistic particles are initialized and set to by Adress::initForces
- that also the atomistic particles are integrated and propagated by Adress::integrate1 and Adress::integrate2

Example - how to turn on the AdResS integrator extension:

```
>>> adress = espressopp.integrator.Adress(system, verletlist, fixedtuplelist)
>>> integrator.addExtension(adress)
```

If KTI is set to True, then the resolution parameters are not updated. This can be used for example for Kirkwood thermodynamic integration, during which one manually sets the whole system on different resolution parameters. KTI = True then prevents overwriting these manually set values. Furthermore, when having moving AdResS regions based on particles, regionupdates specifies the update frequency of the AdResS region in number of steps (or, to be more precise, calls of communicateAdrPositions()). Note that there is a tradeoff: The more frequently the AdResS region is updated, the more gradually and accurately the AdResS region changes and adapts it shape. This could allow for a smaller overall AdResS region and possibly a smoother simulation. However, when having many AdResS region defining particles, these frequent updates can become computationally significant and cost additional simulation time. The optimum is highly system and application dependent.

espressopp.integrator.Adress(_system,_verletlist,_fixedtuplelist, KTI, regionupdates)

Parameters

- _system (shared_ptr<System>) system object
- _verletlist (shared_ptr<VerletListAdress>) verletlist object
- _fixedtuplelist (shared_ptr<FixedTupleListAdress>) fixedtuplelist object
- KTI (bool) (default: False) update resolution parameter? (Yes: set False, No: set True)
- **regionupdates** (*int*) (default: 1) after how many steps does the AdResS region needs to be updated?

4.19.2 BerendsenBarostatAnisotropic - Berendsen barostat Object

#TODO fix these comments This is the Berendsen barostat implementation according to the original paper [Berendsen84]. If Berendsen barostat is defined (as a property of integrator) then at the each run the system size and the particle coordinates will be scaled by scaling parameter μ according to the formula:

$$\mu = [1 - \Delta t / \tau (P_0 - P)]^{1/3}$$

where Δt - integration timestep, τ - time parameter (coupling parameter), P_0 - external pressure and P - instantaneous pressure.

Example:

```
>>> berendsenP = espressopp.integrator.BerendsenBarostatAnisotropic(system)
>>> berendsenP.tau = 0.1
>>> berendsenP.pressure = 1.0
>>> integrator.addExtension(berendsenP)
```

!IMPORTANT In order to run *npt* simulation one should separately define thermostat as well (e.g. Berendsen-Thermostat).

Definition:

In order to define the Berendsen barostat

```
>>> berendsenP = espressopp.integrator.

GrandsenBarostatAnisotropic(system)
```

one should have the System defined.

Properties:

• berendsenP.tau

The property 'tau' defines the time parameter τ .

• berendsenP.pressure

The property 'pressure' defines the external pressure P_0 .

Setting the integration property:

```
>>> integrator.addExtension(berendsenP)
```

It will define Berendsen barostat as a property of integrator.

One more example:

```
>>> berendsen_barostat = espressopp.integrator.BerendsenBarostatAnisotropic(system)
>>> berendsen_barostat.tau = 10.0
>>> berendsen_barostat.pressure = 3.5
>>> integrator.addExtension(berendsen_barostat)
```

Canceling the barostat:

If one do not need the pressure regulation in system anymore or need to switch the ensamble or whatever:)

```
>>> # define barostat with parameters
>>> berendsen = espressopp.integrator.

BerendsenBarostatAnisotropic(system)
>>> berendsen.tau = 0.8
>>> berendsen.pressure = 15.0
>>> integrator.addExtension(berendsen)
>>> ...
>>> # some runs
```

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Connecting the barostat back after the disconnection

```
>>> berendsen.connect()
```

espressopp.integrator.BerendsenBarostatAnisotropic(system)

Parameters system -

4.19.3 CapForce - Integrator Extension

This class can be used to forcecap all particles or a group of particles. Force capping means that the force vector of a particle is rescaled so that the length of the force vector is <= capforce

Example Usage:

```
>>> capforce = espressopp.integrator.CapForce(system, 1000.0)
>>> integrator.addExtension(capForce)
```

CapForce can also be used to forcecap only a group of particles:

```
>>> particle_group = [45, 67, 89, 103]
>>> capforce = espressopp.integrator.CapForce(system, 1000.0, particle_group)
>>> integrator.addExtension(capForce)
```

espressopp.integrator.CapForce (system, capForce, particleGroup)

Parameters

- system -
- capForce -
- particleGroup (default: None)

4.19.4 espressopp.integrator.DPDThermostat

espressopp.integrator.**DPDThermostat** (system, vl)

Parameters

- system -
- v1 -

4.19.5 ExtAnalyze - Integrator Extension

This class can be used to execute nearly all analysis objects within the main integration loop which allows to automatically accumulate time averages (with standard deviation error bars).

Example Usage:

espressopp.integrator.ExtAnalyze(action_obj, interval)

Parameters

- action_obj -
- interval (int) (default: 1)

4.19.6 espressopp.integrator.ExtForce

espressopp.integrator.ExtForce (system, extForce, particleGroup)

Parameters

- system -
- extForce -
- particleGroup (default: None)

4.19.7 espressopp.integrator.Extension

Return type

4.19.8 espressopp.integrator.FixPositions

espressopp.integrator.FixPositions (system, particleGroup, fixMask)

Parameters

- system -
- particleGroup -
- fixMask -

4.19.9 espressopp.integrator.FreeEnergyCompensation

Free Energy Compensation used in Hamiltonian Adaptive Resolution Simulations (H-AdResS). This works for spherical or slab adaptive resolution geometries. However, it only works for fixed, non-moving atomistic region (otherwise, H-AdResS is not properly defined).

Example:

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```
>>> fec = espressopp.integrator.FreeEnergyCompensation(system, center=[Lx/2, Ly/2, Lz/2])
>>> # set up the fec module with the center in the center of the box
>>> fec.addForce(itype=3,filename="tablefec.xvg",type=typeCG)
>>> # set up the actual force
>>> integrator.addExtension(fec)
>>> # add to previously defined integrator
```

espressopp.integrator.FreeEnergyCompensation(system, center, sphereAdr)

Parameters

- **system** (shared_ptr<System>) system object
- **center** (list of reals) (default: [], corresponds to (0.0, 0.0, 0.0) position) center of high resolution region
- **sphereAdr** (bool) (default: False) Spherical AdResS region (True) vs. slab geometry with resolution change in x-direction (False)

espressopp.integrator.FreeEnergyCompensation.addForce(itype, filename, type)

Parameters

- itype (int) interpolation type 1: linear, 2: Akima, 3: Cubic
- **filename** (*string*) filename for TD force file
- type (int) particle type on which the TD force needs to be applied

espressopp.integrator.FreeEnergyCompensation.computeCompEnergy()

Return type real

4.19.10 espressopp.integrator.lsokinetic

```
espressopp.integrator. Isokinetic (system)
```

Parameters system -

4.19.11 LatticeBoltzmann - class for lattice Boltzmann methods

The LatticeBoltzmann (LB) class is an extension to the integrator class of ESPResSo++. The main purpose of the LB-fluid in our simulation package is NOT in fluid dynamics applications or investigation of fluid-solid interfacial phenomena. We aim at complex soft matter systems, where the LB-fluid is a bulk solvent and therefore one has rather use some MD particles as solutes. Examples of such systems range from colloids (point-like MD-particles) to polymer chains (point-like MD-particles connected into chains) dissolved in some solvent (LB-fluid) with specific static and dynamic properties.

It is therefore done ON PURPOSE that the user specifies parameters for LB-fluid in Lennard-Jones (LJ) units. In the kernel of the C++ code we transform these into LB-units, if neccessary. Such strategy helps users coming from MD-background to think of the LB-fluid as if it has particle-based structure: to mimic the solvent one only has to specify such parameters as liquid density, rho, temperature, T, and viscosity, η . For a standard LJ-fluid one has: $rho \sim 1[\sigma^{-3}]$, $T \sim 1[\epsilon]$, and $\eta \sim 5[units]$.

Note: Experienced LB-users may find our approach unusual. However, we kindly ask them for a feedback, as for us it is also quite novel. Particularly, we are interested in suggestions on expansion of the LB-possibilities and would like at first get an overview of "what do the people need?". Being it either BGK-scheme, implementation of boundary conditions or something else.

It creates a simulation box with specified dimensions and allocates necessary memory for a lattice Boltzmann simulation. By default we use D3Q19 lattice model (in three dimensions and with 19-velocities on the node model).

LatticeBoltzmann constructor expects 5 parameters (and a system pointer). These are: lattice size in 3D Ni, lattice spacing a, lattice timestep tau, number of dimensions and number of velocity vectors on a lattice node. The lattice size, Ni, is an obligatory parameter and must be set at the beginning of the simulation.

The default lattice model is D3Q19 (numDims = 3, numVels = 19) and both lattice spacing and timestep are set to 1.

Note that at the present stage of development we aim at D3Q19 model. If you want to use something else, please, feel free to modify the code.

Originally, we had planned this module to operate in 3D only, so if you need a 2D version, there is a bit more tuning involved. On the other hand, adding different 3D lattice models (such as D3Q15 or D3Q27) is rather straightforward.

Example

```
>>> lb = espressopp.integrator.LatticeBoltzmann(system, Ni=Int3D(20, 20, 20))
>>> # creates a cubic box of 20^3 nodes with default spacing parameters in D3Q19_

-model.
```

Example

```
>>> lb = espressopp.integrator.LatticeBoltzmann(system, Ni=Int3D(30, 20, 20), a = 0.5, tau = 0.5)
>>> # creates a box of 30*20*20 nodes with lattice spacing of 0.5 and timestep of 0.5.
>>> # The model of the lattice is D3Q19.
```

After initialization of the Lattice Boltzmann module, one has a possibility to set several properties of the system: gamma_b and gamma_s are bulk and shear gammas (default values are 0.);

gamma odd and gamma even are (hey-hey, surprise!) odd and even gammas (defaults 0.);

(if you are unsure what these gammas are, please refer to any lattice Boltzmann review. In short, they control correspondent viscosities of the liquid.)

lbTemp is the temperature in lb units for setting up fluctuations (default is 0.);

Example

```
>>> lb = espressopp.integrator.LatticeBoltzmann(system, Ni=Int3D(20, 20, 20))
>>> lb.lbTemp = 0.0000005
>>> # creates a box of 20^3 nodes with lattice spacing of 1. and timestep of 1.

\[
\to D3Q19 model.
\]
>>> # then the fluctuations with the temperature of 0.0000005 are initialized.
```

Example

```
>>> lb = espressopp.integrator.LatticeBoltzmann(system, Ni=Int3D(20, 20, 20))
>>> lb.gamma_b = 0.5
>>> lb.gamma_s = 0.5
>>> # creates a box of 20^3 nodes with lattice spacing of 1. and timestep of 1.

\[ \to D3Q19 model. \]
>>> # then the bulk and shear gammas are set to 0.5
```

espressopp.integrator.LatticeBoltzmann(system, nodeGrid, Ni, a, tau, numDims, numVels)

Parameters

• system -

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```
nodeGrid –
Ni –
a – (default: 1.)
tau – (default: 1.)
numDims (int) – (default: 3)
numVels (int) – (default: 19)
```

4.19.12 LBInit - abstract class for LatticeBoltzmann initialization and external force management

This abstract class provides the interface to (re-)initialize populations and handle external forces.

```
espressopp.integrator.LBInit.createDenVel (rho0, u0) to set initial density and velocity of the LB-fluid.
```

Parameters

- **rho0** density
- **u0** velocity

At the moment we support the following options for LB-fluid initialization:

- •espressopp.integrator.LBInitPopUniform for uniformly distributed density and velocity, i.e. on every lattice site the density is rho0 and velocity is u0;
- •espressopp.integrator.LBInitPopWave for uniform density at every lattice site, but harmonic velocity $v_z(x)$ with the period of lattice sites in x-direction;

```
espressopp.integrator.LBInit.setForce(value) to set an external force onto LB-fluid.
```

```
Parameters value (Real3D) – value of the force
```

```
espressopp.integrator.LBInit.addForce (value) to add a new external force to the existing one.
```

Parameters value (Real3D) – value of the force

Two main external force types are implemented:

- espressopp.integrator.LBInitConstForce to manage constant (gravity-like) force acting on every lattice site and
- espressopp.integrator.LBInitPeriodicForce to manage harmonic (position-dependent) force

```
espressopp.integrator.LBInit.addForce(force)
```

```
Parameters force -
```

Return type

```
espressopp.integrator.LBInit.createDenVel(rho0, u0)
```

Parameters

- rho0 -
- u0 –

Return type

```
espressopp.integrator.LBInit.setForce (force)
```

Parameters force -

4.19.13 LBInitConstForce - handles constant (gravity-like) external force

This class sets or adds a constant (gravity-like) external force to the LB-fluid. At first, one has to create an instance. Only after it one may set or add this force to the system.

Example to set the extend force to (0., 0., 0.0005):

```
>>> lbforce1 = espressopp.integrator.LBInitConstForce(system,lb)
>>> lbforce1.setForce(Real3D(0.,0.,0.0005))
>>> # a vector sets the external body force directly in lb-units
```

Example to add an extenal force of (0.0001, 0., 0.) to the existing forces:

```
>>> lbforce2 = espressopp.integrator.LBInitConstForce(system,lb)
>>> lbforce2.addForce(Real3D(0.0001,0.,0.))
>>> # a vector sets the external body force directly in lb-units
```

espressopp.integrator.LBInitConstForce (system, latticeboltzmann)

Parameters

- system -
- latticeboltzmann -

4.19.14 LBInitPeriodicForce - handles external periodic forces

This class sets or adds an external periodic forces to the LB-fluid. At first, one has to create an instance. Only after it one may set or add this force to the system.

Note: Please note, that you have to specify the amplitude of the force. Its particular values at every lattice site will be calculated automatically.

Example to set an external force:

```
>>> lbforce1 = espressopp.integrator.LBInitPeriodicForce(system,lb)
>>> lbforce1.setForce(Real3D(0.,0.,0.0005))
>>> # a vector sets the external body force amplitude
```

Example to add an external force with the amplitude (0.0001, 0., 0.):

```
>>> lbforce2 = espressopp.integrator.LBInitPeriodicForce(system,lb)
>>> lbforce2.addForce(Real3D(0.0001,0.,0.))
>>> # a vector adds the external body force with a Real3D amplitude
```

espressopp.integrator.LBInitPeriodicForce (system, latticeboltzmann)

Parameters

- system -
- latticeboltzmann -

4.19.15 LBInitPopUniform - creates initial populations with uniform density and velocity

This class creates LB-fluid with uniform density rho0 and velocity u0. You have only to specify the corresponding parameters.

Example:

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```
>>> initPop = espressopp.integrator.LBInitPopUniform(system,lb)
>>> initPop.createDenVel(1.0, Real3D(0.,0.,0.0))
>>> # first number is the density, second number is a vector of velocity
```

espressopp.integrator.LBInitPopUniform(system, latticeboltzmann)

Parameters

- system -
- latticeboltzmann -

4.19.16 LBInitPopWave - creates initial populations with uniform density and harmonic velocity

This class creates LB-fluid with uniform density and harmonic velocity: $v_x = 0$, $v_y = 0$, $v_z(i) = A * sin(2 * \pi * i/N_x)$,

where A is the amplitude of the velocity wave, N_x is the number of lattice nodes in x-direction and i is the node index that the velocity is calculated for.

This may be used to test the system: total moment is zero and the liquid tends to equilibrium, i.e. relaxes to a uniform zero velocity.

Example:

```
>>> initPop = espressopp.integrator.LBInitPopWave(system,lb)
>>> initPop.createDenVel(1.0, Real3D(0.,0.,0.0005))
>>> # the Real3D vector in this case includes amplitudes of the velocities
```

espressopp.integrator.LBInitPopWave (system, latticeboltzmann)

Parameters

- system-
- latticeboltzmann -

4.19.17 espressopp.integrator.LangevinThermostat

Langevin Thermostat

Example:

```
>>> langevin = espressopp.integrator.LangevinThermostat(system)
>>> # set up the thermostat
>>> langevin.gamma = gamma
>>> # set friction coefficient gamma
>>> langevin.temperature = temp
>>> # set temperature
>>> langevin.adress = True
>>> # set adress (default is False)
>>> integrator.addExtension(langevin)
>>> # add extensions to a previously defined integrator
```

espressopp.integrator.LangevinThermostat(system)

```
Parameters system(shared_ptr<System>) - system object
```

```
espressopp.integrator.LangevinThermostat.addExclusions(pidlist)
```

Parameters pidlist (list of ints) – list of particle ids to be excluded from thermostating. In adaptive (AdResS) simulations, add ids of atomistic particles to be excluded

(thermostats acts in this case on atomistic level). For normal simulations, add normal or coarse-grained particle ids.

4.19.18 espressopp.integrator.LangevinThermostat1D

espressopp.integrator.LangevinThermostat1D(system)

Parameters system -

```
4.19.19 espressopp.integrator.MDIntegrator

espressopp.integrator.MDIntegrator.addExtension(extension)

Parameters extension -

Return type

espressopp.integrator.MDIntegrator.getExtension(k)

Parameters k -

Return type

espressopp.integrator.MDIntegrator.getNumberOfExtensions()
```

Return type

espressopp.integrator.MDIntegrator.run(niter)

4.19.20 espressopp.integrator.Settle

Parameters niter -

```
espressopp.integrator.Settle(system, fixedtuplelist, mO, mH, distHH, distOH)
```

Parameters

- system-
- fixedtuplelist -
- mO(real) (default: 16.0)
- **mH** (real) (default: 1.0)
- **distHH** (real) (default: 1.58)
- **distOH** (real) (default: 1.0)

 $\verb|espressopp.integrator.Settle.addMolecules| (\textit{moleculelist})$

Parameters moleculelist -

Return type

4.19.21 espressopp.integrator.StochasticVelocityRescaling

```
espressopp.integrator.StochasticVelocityRescaling(system)
Parameters system -
```

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4.19.22 espressopp.integrator.TDforce

Thermodynamic force.

Example - how to turn on thermodynamic force (except for multiple moving spherical regions)

Example - how to turn on thermodynamic force for multiple moving spherical regions

espressopp.integrator.**TDforce** (system, verletlist, startdist, enddist, edgeweightmultiplier)

Parameters

- **system** (shared_ptr<System>) system object
- verletlist (shared ptr<VerletListAdress>) verletlist object
- **startdist** (real) (default: 0.0) starting distance from center at which the TD force is actually applied. Needs to be altered when using several moving spherical regions (not used for static or single moving region)
- **enddist** (real) (default: 0.0) end distance from center up to which the TD force is actually applied. Needs to be altered when using several moving spherical regions (not used for static or single moving region)
- edgeweightmultiplier (int) (default: 20) interpolation parameter for multiple overlapping spherical regions (see Kreis et al., JCTC doi: 10.1021/acs.jctc.6b00440), the default should be fine for most applications (not used for static or single moving region)

espressopp.integrator.TDforce.addForce(itype, filename, type)

Parameters

- itype (int) interpolation type 1: linear, 2: Akima, 3: Cubic
- **filename** (*string*) filename for TD force file
- type (int) particle type on which the TD force needs to be applied

4.19.23 espressopp.integrator.VelocityVerlet

```
espressopp.integrator.VelocityVerlet(system)
```

Parameters system -

4.19.24 espressopp.integrator.VelocityVerletOnGroup

espressopp.integrator.VelocityVerletOnGroup(system, group)

Parameters

- system -
- group -

4.19.25 espressopp.integrator.VelocityVerletOnRadius

espressopp.integrator.VelocityVerletOnRadius (system, dampingmass)

Parameters

- system -
- · dampingmass -

4.19.26 espressopp.integrator.MinimizeEnergy

This is a very simple approach to perform energy minimization of the system. The module uses a steepest descent method. The position of particles is updated following the equation:

$$p_{i+1} = p_i + min(\gamma F_i, d_{max})$$

where p_{i+} is a new position, p_i is a position at current step with corresponding force F_i . The parameters γ and d_{max} are set by user and control the relaxation of the energy and the maximum update of the coordinates per step.

Additionally, a variable γ step is also implemented. In this case, the position of particles is updated following the equation:

$$p_{i+1} = p_i + d_{max} / f_{max} F_i$$

where f_{max} is a maximum force in a single step of steepest descent method. $\gamma = d_{max}/f_{max}$ is automatically adjusted to a force magnitude.

In both cases, the routine runs until the maximum force is bigger than f_{max} or for at most n steps.

Please note This module does not support any integrator extensions.

Example1

```
>>> em = espressopp.integrator.MinimizeEnergy(system, gamma=0.001, ftol=0.01, max_

displacement=0.0001)

>>> em.run(10000)
```

Example2

```
>>> em = espressopp.integrator.MinimizeEnergy(system, gamma=0.01, ftol=0.01, max_

displacement=0.01, variable_step_flag=True)

>>> em.run(10000)
```

API

espressopp.integrator.MinimizeEnergy (system, gamma, ftol, max_displacement, variable_step_flag)

Parameters

• system (espressopp.System) - The espressopp system object.

4.19. integrator 99

- gamma (float) The gamma value.
- **ftol** (float) The force tolerance
- max_displacement (float) The maximum displacement.
- **variable_step_flag** (bool) The flag of adjusting gamma to the force strength.

espressopp.integrator.MinimizeEnergy.run (max_steps, verbose)

Parameters

- max_steps (int) The maximum number of steps to run.
- **verbose** (bool) If set to True then display information about maximum force during the iterations.

Returns The true if the maximum force in the system is lower than ftol otherwise false.

Return type bool

espressopp.integrator.MinimizeEnergy.f_max
The maximum force in the system.

espressopp.integrator.MinimizeEnergy.displacement

The maximum displacement used during the run of MinimizeEnergy

espressopp.integrator.MinimizeEnergy.step

The current iteration step.

4.20 interaction

4.20.1 espressopp.interaction.AngularCosineSquared

Calculates the Angular Cosine Squared interaction

$$U = K(\cos(\theta) - \cos(\theta_0))^2$$

espressopp.interaction.AngularCosineSquared(K, theta0)

Parameters

- **K** (real) (default: 1.0)
- theta0 (real) (default: 0.0)

espressopp.interaction.FixedTripleListAngularCosineSquared(system, vl, potential)

Parameters

- system -
- **v1** -
- potential -

espressopp.interaction.FixedTripleListAngularCosineSquared.getFixedTripleList()

Return type A Python list of lists.

 $\verb|espressopp.interaction.FixedTripleListAngularCosineSquared.setPotential| (type l, the context of the contex$

type2,

po-

ten-

tial)

Parameters

- type1 -
- type2 -
- potential -

class espressopp.interaction.AngularCosineSquared.AngularCosineSquared
 The AngularCosineSquared potential.

4.20.2 espressopp.interaction.AngularHarmonic

Calculates the Angular Harmonic interaction

$$U = K(\theta - \theta_0)^2$$

espressopp.interaction.AngularHarmonic(K, theta0)

Parameters

- **K** (real) (default: 1.0)
- theta0 (real) (default: 0.0)

espressopp.interaction.FixedTripleListAngularHarmonic(system, vl, potential)

Parameters

- system -
- v1 -
- potential -

Parameters

- type1 -
- type2 -
- potential -

class espressopp.interaction.AngularHarmonic.AngularHarmonic
 The AngularHarmonic potential.

4.20.3 espressopp.interaction.AngularPotential

This is an abstract class, only needed to be inherited from.

espressopp.interaction.AngularPotential.computeEnergy(*args)

Parameters *args -

Return type

espressopp.interaction.AngularPotential.computeForce(*args)

Parameters *args -

Return type

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4.20.4 espressopp.interaction.AngularUniqueCosineSquared

Calculates the angular unique cosine squared interaction.

$$U = K(\cos(\theta) - \cos(\theta_0))^2$$

 $\verb|espressopp.interaction.AngularUniqueCosineSquared|(K)$

Parameters K (real) – (default: 1.0)

espressopp.interaction.FixedTripleAngleListAngularUniqueCosineSquared(system,

ftcl,

po-

ten-

tial)

Parameters

- system -
- ftcl -
- potential -

 $\verb|espressopp.interaction.FixedTripleAngleListAngularUniqueCosineSquared.getFixedTripleCosineSquared.getFixedTripleCosin$

Return type A Python list of lists.

Parameters potential -

class espressopp.interaction.AngularUniqueCosineSquared.AngularUniqueCosineSquared
 The AngularUniqueCosineSquared potential.

4.20.5 espressopp.interaction.AngularUniqueHarmonic

Calculates the Angular Unique Harmonic interaction

$$U = K(\theta - \theta_0)^2$$

espressopp.interaction.AngularUniqueHarmonic(K)

Parameters K (real) – (default: 1.0)

espressopp.interaction.FixedTripleAngleListAngularUniqueHarmonic(system,

ftal, potential)

Parameters

- system -
- ftal -
- potential -

espressopp.interaction.FixedTripleAngleListAngularUniqueHarmonic.setPotential(potential)

Parameters potential -

4.20.6 espressopp.interaction.AngularUniquePotential

```
This is an abstract class, only needed to be inherited from.
```

```
espressopp.interaction.AngularUniquePotential.computeEnergy (*args)
```

Parameters *args -

Return type

espressopp.interaction.AngularUniquePotential.computeForce(*args)

Parameters *args -

Return type

4.20.7 espressopp.interaction.Cosine

Calculates the Cosine Interaction

$$U = K(1 + \cos(\theta - \theta_0))$$

espressopp.interaction.Cosine(K, theta0)

Parameters

- **K** (real) (default: 1.0)
- **theta0** (real) (default: 0.0)

espressopp.interaction.FixedTripleListCosine(system, vl, potential)

Parameters

- system -
- v1 -
- potential -

espressopp.interaction.FixedTripleListCosine.getFixedTripleList()

Return type A Python list of lists.

espressopp.interaction.FixedTripleListCosine.setPotential(potential)

Parameters potential -

class espressopp.interaction.Cosine.Cosine
The Cosine potential.

4.20.8 CoulombKSpaceP3M - Coulomb potential and interaction Objects (*K* space part)

This is the *K* space part of potential of Coulomb long range interaction according to the P3M summation technique. Good explanation of P3M summation could be found here [Allen89], [Deserno98].

Example:

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!IMPORTANT Coulomb interaction needs R space part as well CoulombRSpace.

Definition:

It provides potential object *CoulombKSpaceP3M* and interaction object *CellListCoulombKSpaceP3M* based on all particles list.

The *potential* is based on the system information (System) and parameters: Coulomb prefactor (coulomb_prefactor), P3M parameter (alpha), and the cutoff in K space (kspacecutoff).

```
>>> ewaldK_pot = espressopp.interaction.CoulombKSpaceP3M(system, coulomb_ 

->prefactor, alpha, kspacecutoff)
```

Potential Properties:

• ewaldK_pot.prefactor

The property 'prefactor' defines the Coulomb prefactor.

ewaldK_pot.alpha

The property 'alpha' defines the P3M parameter *alpha*.

• ewaldK_pot.kmax

The property 'kmax' defines the cutoff in *K* space.

The *interaction* is based on the all particles list. It needs the information from Storage and K space part of potential.

```
>>> ewaldK_int = espressopp.interaction.CellListCoulombKSpaceP3M(system. 

storage, ewaldK_pot)
```

Interaction Methods:

• getPotential()

Access to the local potential.

Adding the interaction to the system:

```
>>> system.addInteraction(ewaldK_int)
```

espressopp.interaction.CoulombKSpaceP3M(system, C_pref, alpha, M, P, rcut, interpolation)

Parameters

- system -
- C_pref -
- alpha -
- M –
- P -
- rcut -
- interpolation (int) (default: 200192)

 $\verb|espressopp.interaction.CellListCoulombKSpaceP3M| (\textit{storage}, \textit{potential})$

Parameters

- storage -
- potential -

 $\verb|espressopp.interaction.CellListCoulombKSpaceP3M.getPotential()|\\$

Return type

4.20.9 espressopp.interaction.CoulombTruncated

$$U = k \frac{q_i q_j}{d_{ij}}$$

where k is the user-supplied prefactor, q_i is the charge of particle i, and d_{ij} is interparticle distance

In this interaction potential, a different charge can be associated with each particle. For a truncated Coulomb interaction potential where only one q_iq_j value is specified for all interactions, see CoulombTruncatedUniqueCharge.

espressopppp.interaction.CoulombTruncated(prefactor, cutoff)

Parameters

- **prefactor** (real) (default: 1.0) user-supplied prefactor k
- **cutoff** (real) (default: infinity) user-supplied interaction cutoff

espressopppp.interaction.VerletListCoulombTruncated(vl)

Parameters v1 (VerletList) – verlet list object defined earlier in python script

espressopppp.interaction.VerletListCoulombTruncated.getPotential(type1, type2)

Parameters

- type1 (integer) type of first atom in pair
- type2 (integer) type of second atom in pair

Parameters

- type1 (integer) type of first atom in pair
- type2 (integer) type of second atom in pair
- potential (CoulombTruncated potential) potential object defined earlier in python script

 $\verb|espressoppp.interaction.FixedPairListTypesCoulombTruncated| (\textit{system}, \textit{vl}) \\$

Parameters

- system (System) system object defined earlier in the python script
- v1 (FixedPairList) fixedpairlist object defined earlier in the python script

Parameters

- type1 (integer) type of first atom in pair
- type2 (integer) type of second atom in pair
- **potential** (CoulombTruncated potential) **potential** object defined earlier in python script

#Example:

class espressopp.interaction.CoulombTruncated.CoulombTruncated
 The CoulombTruncated potential.

4.20.10 espressopp.interaction.CoulombTruncatedUniqueCharge

$$U = \frac{Q}{d}$$

where Q is the product of the charges of the two particles and d is their distance from each other.

In this interaction potential, a unique $Q = q_i q_j$ value is specified per potential. For a more flexible truncated Coulomb interaction potential where each individual particle has its own charge q_i , see CoulombTruncated.

espressopp.interaction.CoulombTruncatedUniqueCharge(qq, cutoff, shift)

Parameters

- **qq** (real) (default: 1.0)
- cutoff (default: infinity)
- **shift** (default: "auto")

 $\verb|espressopp.interaction.VerletListCoulombTruncatedUniqueCharge| (vl) \\$

Parameters v1 -

espressopp.interaction.VerletListCoulombTruncatedUniqueCharge.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListCoulombTruncatedUniqueCharge.setPotential(type1, type2,

po-

ten-

tial)

Parameters

- type1 -
- type2 -
- potential -

 $\verb|espressopp.interaction.CellListCoulombTruncatedUniqueCharge| (|stor)|$

Parameters stor -

 $\verb|espressopp.interaction.CellListCoulombTruncatedUniqueCharge.setPotential| (type I, the context of the conte$

type2,

po-

ten-

tial)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.FixedPairListCoulombTruncatedUniqueCharge(system,

vl, potential)

Parameters

- system -
- v1 -
- potential -

espressopp.interaction.FixedPairListCoulombTruncatedUniqueCharge.setPotential(potential)

Parameters potential -

 ${\bf class}\ {\tt espressopp.interaction.} Coulomb {\tt TruncatedUniqueCharge.Coulomb {\tt TruncatedUniqueCharge}}\ . {\tt Coulomb {\tt TruncatedUniqueCharge}}\ {\tt Decoulomb {\tt TruncatedUniqueCharge}}\ {\tt potential.}$

4.20.11 espressopp.interaction.DihedralHarmonicCos

$$U = K(\cos(\phi) - \cos(\phi_0))^2$$

 $\verb|espressopp.interaction.DihedralHarmonicCos| (\textit{K}, \textit{phi0})|$

Parameters

- \mathbf{K} (real) (default: 0.0)
- **phi0** (real) (default: 0.0)

espressopp.interaction.FixedQuadrupleListDihedralHarmonicCos(system, fql, potential)

Parameters

- system -
- fql -
- potential -

espressopp.interaction.FixedQuadrupleListDihedralHarmonicCos.getFixedQuadrupleList()

Return type A Python list of lists.

espressopp.interaction.FixedQuadrupleListDihedralHarmonicCos.setPotential(potential)

Parameters potential -

class espressopp.interaction.DihedralHarmonicCos.DihedralHarmonicCos
 The DihedralHarmonicCos potential.

4.20.12 espressopp.interaction.DihedralHarmonicNCos

The dihedral harmonic potential

$$U(\phi_{ijkl}) = K[1 + \cos(N \cdot \phi_{ijkl} - \phi_0)]$$

where the K is a constant, the angles should be provided in radians. The N is a multiplicity.

Reference: http://www.uark.edu/ua/fengwang/DLPOLY2/node49.html

espressopp.interaction.DihedralHarmonicNCos(K, phi0, multiplicity)

Parameters

- **K** (real) (default: 0.0)
- **phi0** (real) (default: 0.0)
- multiplicity (int) (default: 1)

espressopp.interaction.FixedQuadrupleListDihedralHarmonicNCos(system, fql potential)

Parameters

- system -
- fql -
- potential -

espressopp.interaction.FixedQuadrupleListDihedralHarmonicNCos.getFixedQuadrupleList()

Return type A Python list of lists.

Parameters potential -

 ${\bf class} \ {\tt espressopp.interaction.Dihedral Harmonic NCos.Dihedral Harmonic NCos} \\ The \ Dihedral Harmonic NCos \ potential.$

class espressopp.interaction.DihedralHarmonicNCos.FixedQuadrupleListDihedralHarmonicNCosLo

The (local) DihedralHarmonicNCos interaction using FixedQuadruple lists.

4.20.13 espressopp.interaction.DihedralHarmonicUniqueCos

$$U = K(\cos(\phi) - \cos(\phi_0))^2$$

espressopp.interaction. $\mathbf{DihedralHarmonicUniqueCos}(K)$

Parameters K (real) – (default: 0.0)

 $\verb|espressopp.interaction.FixedQuadrupleAngleListDihedralHarmonicUniqueCos| (system, in the context of the con$

fqal,

po-

ten-

tial)

Parameters

- system-
- fqal -

• potential -

 $\verb|espressopp.interaction.FixedQuadrupleAngleListDihedralHarmonicUniqueCos.getFixedQuadrupleAngleCos.getFixedQuadrupleAng$

Return type A Python list of lists.

 ${\tt espressopp.interaction.FixedQuadrupleAngleListDihedralHarmonicUniqueCos.} \textbf{setPotential} \ (\textit{potential} \ \texttt{Cos.}) \ (\textit{po$

Parameters potential -

 ${\bf class}\ {\tt espressopp.interaction.Dihedral Harmonic Unique Cos.Dihedral Harmonic Unique Cos} \\ {\bf The\ Dihedral Harmonic Unique Cos\ potential.}$

4.20.14 espressopp.interaction.DihedralPotential

This is an abstract class, only needed to be inherited from.

espressopp.interaction.DihedralPotential.computeEnergy(*args)

Parameters *args -

Return type

espressopp.interaction.DihedralPotential.computeForce(*args)

Parameters *args -

Return type

4.20.15 espressopp.interaction.DihedralUniquePotential

This is an abstract class, only needed to be inherited from.

espressopp.interaction.DihedralUniquePotential.computeEnergy (*args)

Parameters *args -

Return type

espressopp.interaction.DihedralUniquePotential.computeForce(*args)

Parameters *args -

Return type

4.20.16 espressopp.interaction.DihedralRB

The proper dihedral with Ryckaert-Bellemans form.

$$U_{rb}(\phi_{ijkl}) = \sum_{n=0}^{5} K_n(cos(\theta))^n$$

where the $\theta = \phi - 180^{\circ}$ and $K_{0...5}$ are the coefficients.

By default the IUPAC convention is used, where ϕ is the angle between planes ijk and jkl. The 0° corresponds to the cis configuration.

Reference: http://www.gromacs.org/Documentation/Manual

espressopp.interaction.DihedralRB(K0, K1, K2, K3, K4, K5, iupac)

Parameters

- **KO** (real) (default: 0.0)
- **K1** (real) (default: 0.0)
- **K2** (real) (default: 0.0)

- **K3** (real) (default: 0.0)
- **K4** (real) (default: 0.0)
- **K5** (real) (default: 0.0)
- iupac (default: True)

espressopp.interaction.FixedQuadrupleListDihedralRB(system, vl, potential)

Parameters

- system -
- v1 -
- potential -

espressopp.interaction.FixedQuadrupleListDihedralRB.getFixedQuadrupleList()

Return type A Python list of lists.

Parameters

- type1 -
- type2 -
- potential -

4.20.17 FENE interaction

Implementation of the Finitely Extensible Non-linear Elastic potential:

$$U(r) = -\frac{1}{2}r_{\text{max}}^2 K \log \left[1 - \left(\frac{r - r_0}{r_{\text{max}}}\right)^2\right]$$

espressopp.interaction.**FENE** (*K*, *r0*, *rMax*, *cutoff*, *shift*)

Parameters

- **K** (real) (default: 1.0)
- **r0** (real) (default: 0.0)
- **rMax** (real) (default: 1.0)
- cutoff (real) (default: infinity)
- **shift** (real) (default: 0.0)

 $\verb|espressopp.interaction.FixedPairListFENE| (\textit{system}, \textit{pair_list}, \textit{potential})|$

Parameters

- **system** (object) your system espressopp.System()
- pair_list (object) list of bonds espressopp.FixedPairList()
- potential (object) espressopp.interaction.FENE()

espressopp.interaction.FixedPairListFENE.getFixedPairList()

Return type A Python list of lists.

espressopp.interaction.FixedPairListFENE.getPotential()

Return type object

espressopp.interaction.FixedPairListFENE.setFixedPairList (pair_list)

Parameters pair_list (fixedpairlist) -

espressopp.interaction.FixedPairListFENE.setPotential(potential)

Parameters potential -

Example of usage

4.20.18 espressopp.interaction.FENECapped

$$U = -\frac{1}{2}r_{max}^2 K \cdot \log\left(1 - \frac{D - r_0^2}{r_{max}}\right)$$

where D = dist if

 $cap_{rad}^2 > dist$

and $D = cap_{rad}$ else.

espressopp.interaction.**FENECapped**(*K*, *r*0, *rMax*, *cutoff*, *caprad*, *shift*)

Parameters

```
• K (real) – (default: 1.0)
```

• ${\tt r0}$ (real) – (default: 0.0)

• **rMax** (real) - (default: 1.0)

• **cutoff** – (default: infinity)

• **caprad** (real) - (default: 1.0)

• **shift** (real) – (default: 0.0)

 $\verb|espressopp.interaction.FixedPairListFENECapped| (\textit{system}, \textit{vl}, \textit{potential})$

Parameters

- \bullet system -
- **v1** -
- potential -

espressopp.interaction.FixedPairListFENECapped.getFixedPairList()

Return type A Python list of lists.

espressopp.interaction.FixedPairListFENECapped.getPotential()

Return type

espressopp.interaction.FixedPairListFENECapped.setFixedPairList (fixedpairlist)

Parameters fixedpairlist -

espressopp.interaction.FixedPairListFENECapped.setPotential(potential)

Parameters potential -

class espressopp.interaction.FENECapped.FENECapped
 The FENECapped potential.

4.20.19 GravityTruncated

This is an implementation of a truncated (cutoff) Gravity Potential

$$U = P \cdot \frac{m_1 \cdot m_2}{|p_1 - p_2|}$$

where m_i is the mass of the *i* th particle, p_i its position and *P* a prefactor.

espressopp.interaction.GravityTruncated(prefactor, cutoff)

Parameters

- prefactor (real) (default: 1.0)
- **cutoff** (default: infinity)

espressopp.interaction. VerletListGravityTruncated(vl)

Parameters v1 -

espressopp.interaction.VerletListGravityTruncated.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListGravityTruncated.getVerletList()

Return type A Python list of lists.

Parameters

- type1 -
- type2 -
- potential -

4.20.20 espressopp.interaction.Harmonic

$$U = K(d - r_0)^2$$

espressopp.interaction.Harmonic(K, r0, cutoff, shift)

Parameters

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```
• K (real) – (default: 1.0)
              • r0 (real) – (default: 0.0)
              • cutoff – (default: infinity)
              • shift (real) – (default: 0.0)
espressopp.interaction.FixedPairListHarmonic (system, vl, potential)
        Parameters
              • system -
              • v1 -
              • potential -
espressopp.interaction.FixedPairListHarmonic.getFixedPairList()
         Return type A Python list of lists.
espressopp.interaction.FixedPairListHarmonic.setFixedPairList(fixedpairlist)
        Parameters fixedpairlist -
espressopp.interaction.FixedPairListHarmonic.setPotential(potential)
        Parameters potential -
espressopp.interaction.FixedPairListTypesHarmonic(system, vl)
        Parameters
              • system -
espressopp.interaction.FixedPairListTypesHarmonic.getFixedPairList()
         Return type A Python list of lists.
espressopp.interaction.FixedPairListTypesHarmonic.setFixedPairList(fixedpairlist)
        Parameters fixedpairlist -
espressopp.interaction.FixedPairListTypesHarmonic.setPotential(typeI)
                                                                        type2,
                                                                        potential)
         Parameters
              • type1 -
              • type2 -
              • potential -
espressopp.interaction.FixedPairListTypesHarmonic.getPotential(typeI,
                                                                        type2)
        Parameters
              • type1 -
              • type2 -
         Return type
class espressopp.interaction.Harmonic.Harmonic
    The Harmonic potential.
```

4.20.21 espressopp.interaction.HarmonicUnique

$$U = K(d - d_{cur})^2;$$

espressopp.interaction. HarmonicUnique(K)

Parameters K (real) – (default: 1.0)

espressopp.interaction.FixedPairDistListHarmonicUnique (system, fpl, potential)

Parameters

- system -
- fpl -
- potential -

espressopp.interaction.FixedPairDistListHarmonicUnique.getFixedPairList()

Return type A Python list of lists.

espressopp.interaction.FixedPairDistListHarmonicUnique.setFixedPairList (fixedpairlist)

Parameters fixedpairlist -

espressopp.interaction.FixedPairDistListHarmonicUnique.setPotential(potential)

Parameters potential -

class espressopp.interaction.HarmonicUnique.HarmonicUnique
The HarmonicUnique potential.

4.20.22 espressopp.interaction.Interaction

This is an abstract class, only needed to be inherited from.

espressopp.interaction.Interaction.bondType()

Return type int

espressopp.interaction.Interaction.computeEnergy()

Return type real

espressopp.interaction.Interaction.computeEnergyAA()

Return type real

espressopp.interaction.Interaction.computeEnergyDeriv()

Return type real

espressopp.interaction.Interaction.computeEnergyCG()

Return type real

espressopp.interaction.Interaction.computeVirial()

Return type real

4.20.23 espressopp.interaction.LJcos

if $r^2 \leq border_{pot}$, then:

$$U = 4\left(\frac{1}{r^{12}} - \frac{1}{r^6}\right) + 1 - \phi$$

else:

$$U = \frac{1}{2}\phi(\cos(\alpha r^2 + \beta) - 1)$$

espressopp.interaction.LJcos(phi)

Parameters phi (real) - (default: 1.0)

espressopp.interaction.VerletListLJcos(vl)

Parameters v1 -

espressopp.interaction.VerletListLJcos.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListLJcos.getVerletList()

Return type A Python list of lists.

espressopp.interaction.VerletListLJcos.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

 $\verb|espressopp.interaction.VerletListAdressLJcos| (vl, \textit{fixedtupleList})$

Parameters

- **v1** -
- fixedtupleList -

espressopp.interaction.VerletListAdressLJcos.setPotentialAT(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListAdressLJcos.setPotentialCG(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListHadressLJcos(vl, fixedtupleList)

Parameters

- v1 -
- fixedtupleList -

espressopp.interaction.VerletListHadressLJcos.setPotentialAT(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.CellListLJcos(stor)

Parameters stor -

espressopp.interaction.CellListLJcos.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.FixedPairListLJcos (system, vl, potential)

Parameters

- system-
- v1 -
- potential -

espressopp.interaction.FixedPairListLJcos.getFixedPairList()

Return type A Python list of lists.

espressopp.interaction.FixedPairListLJcos.setFixedPairList(fixedpairlist)

Parameters fixedpairlist -

espressopp.interaction.FixedPairListLJcos.setPotential(potential)

Parameters potential -

class espressopp.interaction.LJcos.LJcos
 The Lennard-Jones potential.

4.20.24 espressopp.interaction.LennardJones

$$V(r) = 4\varepsilon \left[\left(\frac{\sigma}{r} \right)^{12} - \left(\frac{\sigma}{r} \right)^{6} \right]$$

espressopp.interaction.LennardJones (epsilon, sigma, cutoff, shift)

Parameters

• **epsilon** (real) – (default: 1.0)

```
• sigma (real) - (default: 1.0)
              • cutoff – (default: infinity)
              • shift – (default: "auto")
espressopp.interaction.VerletListLennardJones(vl)
         Parameters v1 -
espressopp.interaction.VerletListLennardJones.getPotential(type1, type2)
         Parameters
              • type1 -
              • type2 -
         Return type
espressopp.interaction.VerletListLennardJones.getVerletList()
         Return type A Python list of lists.
espressopp.interaction.VerletListLennardJones.setPotential(type1, type2, po-
         Parameters
              • type1 -
              • type2 -
              • potential -
espressopp.interaction.VerletListAdressLennardJones(vl, fixedtupleList)
         Parameters
              • v1 -
              • fixedtupleList -
\verb|espressopp.interaction.VerletListAdressLennardJones.setPotentialAT| ( \textit{type1},
                                                                               type2,
                                                                              poten-
                                                                              tial)
         Parameters
              • type1 -
              • type2 -
              • potential -
espressopp.interaction.VerletListAdressLennardJones.setPotentialCG(type1,
                                                                               type2,
                                                                              poten-
                                                                              tial)
         Parameters
              • type1 -
              • type2 -
              • potential -
espressopp.interaction.VerletListAdressLennardJones2(vl, fixedtupleList)
         Parameters
              • v1 -
```

• fixedtupleList -

```
\verb|espressopp.interaction.VerletListAdressLennardJones2.setPotentialAT| (type I, to be a substitution of the property of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     type2,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ten-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tial)
                                                     Parameters
                                                                                    • type1 -
                                                                                    • type2 -
                                                                                    • potential -
\verb|espressopp.interaction.VerletListAdressLennardJones2.setPotentialCG| (type I, the interaction of the int
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     po-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ten-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tial)
                                                     Parameters
                                                                                    • type1 -
                                                                                    • type2 -
                                                                                    • potential -
espressopp.interaction.VerletListHadressLennardJones(vl, fixedtupleList)
                                                     Parameters
                                                                                    • v1 -
                                                                                    • fixedtupleList -
\verb|espressopp.interaction.VerletListHadressLennardJones.setPotentialAT| (type I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    po-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ten-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tial)
                                                     Parameters
                                                                                    • type1 -
                                                                                    • type2 -
                                                                                    • potential -
espressopp.interaction.VerletListHadressLennardJones.setPotentialCG(type1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    po-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ten-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tial)
                                                     Parameters
                                                                                     • type1 -
                                                                                    • type2 -
                                                                                    • potential -
espressopp.interaction.VerletListHadressLennardJones2(vl, fixedtupleList)
                                                     Parameters
                                                                                    • v1 -
                                                                                    • fixedtupleList -
```

```
espressopp.interaction.VerletListHadressLennardJones2.setPotentialAT (type I)
                                                                              type2,
                                                                              po-
                                                                              ten-
                                                                              tial)
        Parameters
             • type1 -
             • type2 -
             • potential -
\verb|espressopp.interaction.VerletListHadressLennardJones2.setPotentialCG| (type I,
                                                                              type2,
                                                                              po-
                                                                              ten-
                                                                              tial)
        Parameters
             • type1 -
             • type2 -
             • potential -
espressopp.interaction.CellListLennardJones(stor)
        Parameters stor -
espressopp.interaction.CellListLennardJones.setPotential(type1, type2, poten-
        Parameters
             • type1 -
             • type2 -
             • potential -
espressopp.interaction.FixedPairListLennardJones (system, vl, potential)
        Parameters
             • system -
             • v1 -
             • potential -
espressopp.interaction.FixedPairListLennardJones.getFixedPairList()
        Return type A Python list of lists.
espressopp.interaction.FixedPairListLennardJones.getPotential()
        Return type
espressopp.interaction.FixedPairListLennardJones.setFixedPairList (fixedpairlist)
        Parameters fixedpairlist -
espressopp.interaction.FixedPairListLennardJones.setPotential(potential)
        Parameters potential -
class espressopp.interaction.LennardJones.LennardJones
    The Lennard-Jones potential.
```

4.20.25 espressopp.interaction.LennardJonesAutoBonds

$$V(r) = 4\varepsilon \left[\left(\frac{\sigma}{r}\right)^{12} - \left(\frac{\sigma}{r}\right)^{6} \right]$$

espressopp.interaction.LennardJonesAutoBonds (epsilon, sigma, cutoff, bondlist, max-crosslinks)

Parameters

- epsilon (real) (default: 1.0)
- **sigma** (real) (default: 1.0)
- cutoff (default: infinity)
- bondlist (default: None)
- maxcrosslinks (int) (default: 2)

espressopp.interaction.VerletListLennardJonesAutoBonds (vl)

Parameters v1 -

espressopp.interaction.VerletListLennardJonesAutoBonds.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListLennardJonesAutoBonds.getVerletList()

Return type A Python list of lists.

espressopp.interaction.VerletListLennardJonesAutoBonds.setPotential(type1, type2,

po-

ten-

4: -1

tial)

Parameters

- type1 -
- type2 -
- potential -

 ${\tt espressopp.interaction.} \textbf{VerletListAdressLennardJonesAutoBonds} (\textit{vl}, & \textit{fixedtu-pleList})$

Parameters

- v1 -
- fixedtupleList -

 $\verb|espressopp.interaction.VerletListAdressLennardJonesAutoBonds.setPotential| (|typeI|, |text|) | |text| | |te$

type2, po-

ten-

tial)

Parameters

• type1 -

- type2 -
- potential -

espressopp.interaction.VerletListHadressLennardJonesAutoBonds(vl, fixedtu-pleList)

Parameters

- v1 -
- fixedtupleList -

espressopp.interaction.VerletListHadressLennardJonesAutoBonds.setPotential(type1, type2,

iype

po-

tential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.CellListLennardJonesAutoBonds(stor)

Parameters stor -

 $\verb|espressopp.interaction.CellListLennardJonesAutoBonds.setPotential| (type I, the property of the property o$

type2,

potential)

Parameters

- type1 -
- type2 -
- potential -

 ${\tt espressopp.interaction.} \textbf{FixedPairListLennardJonesAutoBonds} (\textit{system}, \textit{vl}, \textit{potential})$

Parameters

- \bullet system -
- **v1** -
- potential -

espressopp.interaction.FixedPairListLennardJonesAutoBonds.setPotential(potential)

Parameters potential -

class espressopp.interaction.LennardJonesAutoBonds.LennardJonesAutoBonds
 The Lennard-Jones auto bonds potential.

4.20.26 espressopp.interaction.LennardJonesCapped

$$V(r) = 4\varepsilon \left[\left(\frac{\sigma}{r} \right)^{12} - \left(\frac{\sigma}{r} \right)^{6} \right]$$

where r is either the distance or the capped distance, depending on which is greater.

espressopp.interaction.LennardJonesCapped(epsilon, sigma, cutoff, caprad, shift)

Parameters

```
• epsilon (real) – (default: 1.0)
              • sigma (real) - (default: 1.0)
              • cutoff – (default: infinity)
              • caprad (real) - (default: 0.0)
              • shift – (default: "auto")
espressopp.interaction.VerletListLennardJonesCapped(vl)
         Parameters v1 -
espressopp.interaction.VerletListLennardJonesCapped.getPotential(typeI,
         Parameters
              • type1 -
              • type2 -
         Return type
espressopp.interaction.VerletListLennardJonesCapped.setPotential(typeI,
                                                                             type2,
                                                                             poten-
                                                                             tial)
         Parameters
              • type1 -
              • type2 -
              • potential -
espressopp.interaction.VerletListAdressLennardJonesCapped(vl, fixedtupleList)
         Parameters
              • v1 -
              • fixedtupleList -
\verb|espressopp.interaction.VerletListAdressLennardJonesCapped.getPotentialAT| (type I,
                                                                                      type2)
         Parameters
              • type1 -
              • type2 -
         Return type
\verb|espressopp.interaction.VerletListAdressLennardJonesCapped.getPotentialCG| (type I,
                                                                                      type2)
         Parameters
              • type1 -
              • type2 -
         Return type
\verb|espressopp.interaction.VerletListAdressLennardJonesCapped.setPotentialAT| (type I,
                                                                                      type2,
                                                                                      po-
                                                                                      ten-
                                                                                      tial)
```

Parameters

```
• type1 -
                                                                                                                 • type2 -
                                                                                                                 • potential -
\verb|espressopp.interaction.VerletListAdressLennardJonesCapped.setPotentialCG| (type I, the context of the conte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 po-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ten-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tial)
                                                                       Parameters
                                                                                                                 • type1 -
                                                                                                                 • type2 -
                                                                                                                 • potential -
\verb|espressopp.interaction.VerletListHadressLennardJonesCapped| (vl, fixed tuple List)|
                                                                       Parameters
                                                                                                                 • v1 -
                                                                                                                 • fixedtupleList -
espressopp.interaction.VerletListHadressLennardJonesCapped.getPotentialAT (type1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          type2)
                                                                       Parameters
                                                                                                                 • type1 -
                                                                                                                 • type2 -
                                                                        Return type
\verb|espressopp.interaction.VerletListHadressLennardJonesCapped.getPotentialCG| (type I, the context of the cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          type2)
                                                                       Parameters
                                                                                                                 • type1 -
                                                                                                                 • type2 -
                                                                       Return type
\verb|espressopp.interaction.VerletListHadressLennardJonesCapped.setPotentialAT| (type I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          po-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ten-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tial)
                                                                        Parameters
                                                                                                                 • type1 -
                                                                                                                 • type2 -
                                                                                                                 • potential -
\verb|espressopp.interaction.VerletListHadressLennardJonesCapped.setPotentialCG| (type I, the context of the cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          po-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tial)
                                                                       Parameters
                                                                                                                 • type1 -
```

- type2 -
- potential -

 $\verb|espressopp.interaction.CellListLennardJonesCapped| (stor)$

Parameters stor -

espressopp.interaction.CellListLennardJonesCapped.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.CellListLennardJonesCapped.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.FixedPairListLennardJonesCapped(system, vl, potential)

Parameters

- system -
- v1 -
- potential -

espressopp.interaction.FixedPairListLennardJonesCapped.getPotential()

Return type

espressopp.interaction.FixedPairListLennardJonesCapped.setPotential(potential)

Parameters potential -

class espressopp.interaction.LennardJonesCapped.LennardJonesCapped
 The Lennard-Jones potential.

4.20.27 espressopp.interaction.LennardJonesEnergyCapped

$$V(r) = 4\varepsilon \left[\left(\frac{\sigma}{r} \right)^{12} - \left(\frac{\sigma}{r} \right)^{6} \right]$$

where r is either the distance or the capped distance, depending on which is greater.

espressopp.interaction.LennardJonesEnergyCapped(epsilon, sigma, cutoff, caprad, shift)

Parameters

- epsilon (real) (default: 1.0)
- **sigma** (real) (default: 1.0)
- **cutoff** (default: infinity)
- caprad (real) (default: 0.0)
- **shift** (default: "auto")

```
espressopp.interaction.VerletListLennardJonesEnergyCapped(vl)
                                                                  Parameters v1 -
\verb|espressopp.interaction.VerletListLennardJonesEnergyCapped.getPotential| (type I, to be a substitution of the property of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  type2)
                                                                  Parameters
                                                                                                         • type1 -
                                                                                                         • type2 -
                                                                   Return type
espressopp.interaction.VerletListLennardJonesEnergyCapped.setPotential(type1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 po-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ten-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tial)
                                                                  Parameters
                                                                                                         • type1 -
                                                                                                         • type2 -
                                                                                                         • potential -
espressopp.interaction.VerletListAdressLennardJonesEnergyCapped(vl, fixedtu-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pleList)
                                                                  Parameters
                                                                                                        • v1 -
                                                                                                         • fixedtupleList -
\verb|espressopp.interaction.VerletListAdressLennardJonesEnergyCapped.getPotentialAT| (type 1, the context of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      type2)
                                                                  Parameters
                                                                                                         • type1 -
                                                                                                         • type2 -
                                                                  Return type
\verb|espressopp.interaction.VerletListAdressLennardJonesEnergyCapped.getPotentialCG| (type I, the property of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      type2)
                                                                  Parameters
                                                                                                         • type1 -
                                                                                                         • type2 -
                                                                   Return type
espressopp.interaction.VerletListAdressLennardJonesEnergyCapped.setPotentialAT (type1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     po-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ten-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tial)
                                                                  Parameters
                                                                                                          • type1 -
                                                                                                          • type2 -
```

• potential -

```
\verb|espressopp.interaction.VerletListAdressLennardJonesEnergyCapped.setPotentialCG| (type I, the context of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           po-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ten-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tial)
                                                                                           Parameters
                                                                                                                                               • type1 -
                                                                                                                                               • type2 -
                                                                                                                                               • potential -
\verb|espressopp.interaction.VerletListHadressLennardJonesEnergyCapped| (vl,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fixedtu-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pleList)
                                                                                           Parameters
                                                                                                                                               • v1 -
                                                                                                                                               • fixedtupleList -
\verb|espressopp.interaction.VerletListHadressLennardJonesEnergyCapped.getPotentialAT| (type I, the context of th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        type2)
                                                                                           Parameters
                                                                                                                                               • type1 -
                                                                                                                                               • type2 -
                                                                                           Return type
espressopp.interaction.VerletListHadressLennardJonesEnergyCapped.getPotentialCG (type I),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        type2)
                                                                                           Parameters
                                                                                                                                               • type1 -
                                                                                                                                               • type2 -
                                                                                            Return type
\verb|espressopp.interaction.VerletListHadressLennardJonesEnergyCapped.setPotentialAT| (type I, the property of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       po-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ten-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tial)
                                                                                           Parameters
                                                                                                                                               • type1 -
                                                                                                                                               • type2 -
                                                                                                                                               • potential -
\verb|espressopp.interaction.VerletListHadressLennardJonesEnergyCapped.setPotentialCG| (type I, the standard of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       po-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ten-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tial)
                                                                                           Parameters
                                                                                                                                                 • type1 -
```

• type2 -

• potential -

espressopp.interaction.CellListLennardJonesEnergyCapped(stor)

Parameters stor -

espressopp.interaction.CellListLennardJonesEnergyCapped.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

 $\verb|espressopp.interaction.CellListLennardJonesEnergyCapped.setPotential| (type I, to be a constant of the con$

type2,

po-

ten-

tial)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.FixedPairListLennardJonesEnergyCapped(system, vl, potential)

Parameters

- system -
- v1 -
- potential -

espressopp.interaction.FixedPairListLennardJonesEnergyCapped.getPotential()

Return type

espressopp.interaction.FixedPairListLennardJonesEnergyCapped.setPotential(potential)

Parameters potential -

 ${\bf class}\ {\tt espressopp.interaction.LennardJonesEnergyCapped.LennardJonesEnergyCapped}\ The\ Lennard-Jones\ potential.$

4.20.28 espressopp.interaction.LennardJonesExpand

$$V(r) = 4\varepsilon \left[\left(\frac{\sigma}{r} \right)^{12} - \left(\frac{\sigma}{r} \right)^{6} \right]$$

espressopp.interaction.LennardJonesExpand (epsilon, sigma, delta, cutoff, shift)

Parameters

- **epsilon** (*real*) (default: 1.0)
- **sigma** (real) (default: 1.0)
- **delta** (real) (default: 0.0)
- **cutoff** (default: infinity)
- shift (default: "auto")

espressopp.interaction.VerletListLennardJonesExpand(vl)

Parameters v1 -

espressopp.interaction.VerletListLennardJonesExpand.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.CellListLennardJonesExpand(stor)

Parameters stor -

espressopp.interaction.CellListLennardJonesExpand.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.FixedPairListLennardJonesExpand(system, vl, potential)

Parameters

- system-
- v1 -
- potential -

espressopp.interaction.FixedPairListLennardJonesExpand.setPotential (potential)

${\bf Parameters\ potential} -$

class espressopp.interaction.LennardJonesExpand.LennardJonesExpand
 The LennardJonesExpand potential.

4.20.29 espressopp.interaction.LennardJonesGromacs

 $\text{if } d^2 > r_1^2$

$$U = 4\varepsilon(\frac{sigma^{12}}{d^{12}} - \frac{sigma^{6}}{d^{6}}) + (d - r_{1})^{3}(ljsw3 + ljsw4(d - r_{1}) + ljsw5)$$

else

$$U = 4\varepsilon (\frac{\sigma^{12}}{d^{12}} - \frac{\sigma^6}{d^6})$$

```
espressopp.interaction.LennardJonesGromacs (epsilon, sigma, r1, cutoff, shift)
         Parameters
              • epsilon (real) – (default: 1.0)
              • sigma (real) - (default: 1.0)
              • r1 (real) – (default: 0.0)
              • cutoff – (default: infinity)
              • shift – (default: "auto")
espressopp.interaction.VerletListLennardJonesGromacs(vl)
         Parameters v1 -
espressopp.interaction.VerletListLennardJonesGromacs.getPotential(typeI,
                                                                            type2)
         Parameters
              • type1 -
              • type2 -
         Return type
espressopp.interaction.VerletListLennardJonesGromacs.setPotential(typeI,
                                                                            type2,
                                                                            poten-
                                                                            tial)
         Parameters
              • type1 -
              • type2 -
              • potential -
espressopp.interaction.CellListLennardJonesGromacs(stor)
         Parameters stor -
espressopp.interaction.CellListLennardJonesGromacs.setPotential(typeI,
                                                                          type2,
                                                                          potential)
         Parameters
              • type1 -
              • type2 -
              • potential -
espressopp.interaction.FixedPairListLennardJonesGromacs(system, vl, potential)
         Parameters
              • system -
              • v1 -
              • potential -
espressopp.interaction.FixedPairListLennardJonesGromacs.setPotential(potential)
         Parameters potential -
class espressopp.interaction.LennardJonesGromacs.LennardJonesGromacs
    The LennardJonesGromacs potential.
```

4.20.30 espressopp.interaction.Morse

This class provides methods to compute forces and energies of the Morse potential.

$$U = \varepsilon \left(e^{-2\alpha(r - r_{min})} - 2e^{-\alpha(r - r_{min})} \right)$$

espressopp.interaction.Morse(epsilon, alpha, rMin, cutoff, shift)

Parameters

- epsilon (real) (default: 1.0)
- **alpha** (real) (default: 1.0)
- **rMin** (real) (default: 0.0)
- **cutoff** (default: infinity)
- **shift** (default: "auto")

espressopp.interaction.VerletListMorse (vl)

Parameters v1 -

espressopp.interaction.VerletListMorse.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListMorse.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListAdressMorse(vl, fixedtupleList)

Parameters

- **v1** -
- fixedtupleList -

espressopp.interaction.VerletListAdressMorse.setPotentialAT(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListAdressMorse.setPotentialCG(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListHadressMorse(vl, fixedtupleList)

Parameters

- v1 -
- fixedtupleList -

Parameters

- type1 -
- type2 -
- potential -

Parameters

- type1 -
- type2 -
- potential -

 $\verb|espressopp.interaction.CellListMorse| (stor)$

Parameters stor -

espressopp.interaction.CellListMorse.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.FixedPairListMorse(system, vl, potential)

Parameters

- system -
- v1 -
- potential -

espressopp.interaction.FixedPairListMorse.setPotential(potential)

Parameters potential -

class espressopp.interaction.Morse.Morse
 The Morse potential.

4.20.31 espressopp.interaction.OPLS

This class provides methods to compute forces and energies of the OPLS dihedral potential. To create a new dihedral potential.

$$U = \sum_{j=1}^{4} K_j (1 + \cos(j\phi))$$

espressopp.interaction.**OPLS** (K1, K2, K3, K4)

Parameters

```
• K1 (real) – (default: 1.0)
```

• **K2** (real) – (default: 0.0)

• **K3** (real) – (default: 0.0)

• **K4** (real) – (default: 0.0)

espressopp.interaction.FixedQuadrupleListOPLS(system, vl, potential)

Parameters

- system -
- v1 -
- potential -

espressopp.interaction.FixedQuadrupleListOPLS.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

class espressopp.interaction.OPLS.OPLS
 The OPLS potential.

4.20.32 espressopp.interaction.Potential

This is an abstract class, only needed to be inherited from.

```
espressopp.interaction.Potential.computeEnergy(*args)
```

Parameters *args -

Return type

espressopp.interaction.Potential.computeForce(*args)

Parameters *args -

Return type

4.20.33 espressopp.interaction.PotentialUniqueDist

This is an abstract class, only needed to be inherited from.

```
espressopp.interaction.PotentialUniqueDist.computeEnergy(*args)
```

Parameters *args -

Return type

espressopp.interaction.PotentialUniqueDist.computeForce(*args)

Parameters *args -

Return type

4.20.34 espressopp.interaction.PotentialVSpherePair

This is an abstract class, only needed to be inherited from.

espressopp.interaction.PotentialVSpherePair.computeEnergy(*args)

Parameters *args -

Return type

espressopp.interaction.PotentialVSpherePair.computeForce(*args)

Parameters *args -

Return type

4.20.35 espressopp.interaction.Quartic

This class provides methods to compute forces and energies of the Quartic potential.

$$U = \frac{K}{4} \left(d^2 - r_0^2 \right)^2$$

espressopp.interaction.Quartic(K, r0, cutoff, shift)

Parameters

- **K** (real) (default: 1.0)
- **r0** (real) (default: 0.0)
- cutoff (default: infinity)
- **shift** (real) (default: 0.0)

espressopp.interaction.FixedPairListQuartic(system, vl, potential)

Parameters

- system -
- v1 -
- potential -

espressopp.interaction.FixedPairListQuartic.getFixedPairList()

Return type A Python list of lists.

 $\verb|espressopp.interaction.FixedPairListQuartic.setFixedPairList(|fixedpairlist|)|$

Parameters fixedpairlist -

espressopp.interaction.FixedPairListQuartic.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

class espressopp.interaction.Quartic.Quartic

The Quartic potential.

4.20.36 espressopp.interaction.ReactionFieldGeneralized

This class provides methods to compute forces and energies of the generalized reaction field.

$$U = PQ \left(\frac{1}{d} - \frac{\left(1 + \frac{(\varepsilon_1 - 4\varepsilon_2)(1 + \kappa r_c) - 2\varepsilon_2 \kappa r_c^2}{(\varepsilon_1 + 2\varepsilon_2)(1 + \kappa r_c) + \varepsilon_2 \kappa r_c^2} \right)}{r_c^3 2} \cdot d^2 - \frac{3\varepsilon_2}{r_c(2\varepsilon_2 + 1)} \right)$$

where P is a prefactor, Q is the product of the charges of the two particles, d is their distance from each other, and r_c the cutoff-radius.

espressopp.interaction.ReactionFieldGeneralized(prefactor, kappa, epsilon1, ep*silon2*, *cutoff*, *shift*)

Parameters

- prefactor (real) (default: 1.0)
- **kappa** (*real*) (default: 0.0)
- **epsilon1** (real) (default: 1.0)
- epsilon2 (real) (default: 80.0)
- **cutoff** (default: infinity)
- **shift** (default: "auto")

espressopp.interaction. VerletListReactionFieldGeneralized (vl)

Parameters v1 -

 $\verb|espressopp.interaction.VerletListReactionFieldGeneralized.getPotential| (\textit{type1}, in the latest open and the latest open$ type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListReactionFieldGeneralized.setPotential(type1, type2,

po-

ten-

tial)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListAdressReactionFieldGeneralized(vl, fixedtupleList)

Parameters

- v1 -
- fixedtupleList -

espressopp.interaction.VerletListAdressReactionFieldGeneralized.setPotentialAT (type1,

type2,

po-

ten-

tial)

Parameters

```
• type1 -
                                                                                                                   • type2 -
                                                                                                                   • potential -
\verb|espressopp.interaction.VerletListAdressReactionFieldGeneralized.setPotentialCG| (type I, the property of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       po-
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                                                                         Parameters
                                                                                                                   • type1 -
                                                                                                                   • type2 -
                                                                                                                   • potential -
\verb|espressopp.interaction.VerletListHadressReactionFieldGeneralized| (vl, in the context of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 fixedtu-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pleList)
                                                                         Parameters
                                                                                                                   • v1 -
                                                                                                                   • fixedtupleList -
\verb|espressopp.interaction.VerletListHadressReactionFieldGeneralized.setPotentialAT| (type I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 type2,
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                                                                         Parameters
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                                                                                                                   • type2 -
                                                                                                                   • potential -
\verb|espressopp.interaction.VerletListHadressReactionFieldGeneralized.setPotentialCG| (type I, the context of th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 type2,
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                                                                         Parameters
                                                                                                                   • type1 -
                                                                                                                    • type2 -
                                                                                                                   • potential -
espressopp.interaction.CellListReactionFieldGeneralized(stor)
                                                                         Parameters stor -
espressopp.interaction.CellListReactionFieldGeneralized.setPotential(type1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         type2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         po-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tial)
                                                                         Parameters
```

type1 –type2 –

• potential -

4.20.37 espressopp.interaction.SoftCosine

This class provides methods to compute forces and energies ofthe SoftCosine potential.

$$V(r) = A \left[1.0 + \cos \left(\frac{\pi r}{r_c} \right) \right]$$

espressopp.interaction.SoftCosine(A, cutoff, shift)

Parameters

- **A**(real) (default: 1.0)
- cutoff (default: infinity)
- **shift** (default: "auto")

espressopp.interaction.VerletListSoftCosine(stor)

Parameters stor -

espressopp.interaction.VerletListSoftCosine.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.CellListSoftCosine(stor)

Parameters stor -

espressopp.interaction.CellListSoftCosine.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.FixedPairListSoftCosine(system, vl, potential)

Parameters

- system -
- v1 -
- potential -

espressopp.interaction.FixedPairListSoftCosine.setPotential(potential)

Parameters potential -

 ${\bf class} \ {\bf espressopp.interaction.SoftCosine.SoftCosine} \\ {\bf The \ SoftCosine \ potential.}$

4.20.38 espressopp.interaction.StillingerWeberPairTerm

This class provides methods to compute forces and energies of 2 body term of Stillinger-Weber potential.

$$U = \varepsilon A \left[\frac{d}{\sigma}^{-p} (B - 1) \right] exp \left(\frac{1}{\frac{d}{\sigma} - r_c} \right)$$

where r_c is the cutoff-radius.

espressopp.interaction. StillingerWeberPairTerm(A, B, p, q, epsilon, sigma, cutoff)

Parameters

- A -
- B -
- p -
- epsilon (real) (default: 1.0)
- **sigma** (real) (default: 1.0)
- **cutoff** (default: infinity)

espressopp.interaction.VerletListStillingerWeberPairTerm(vl)

Parameters v1 -

 $\verb|espressopp.interaction.VerletListStillingerWeberPairTerm.getPotential| (typeI,$ type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListStillingerWeberPairTerm.getVerletList()

Return type A Python list of lists.

 $\verb|espressopp.interaction.VerletListStillingerWeberPairTerm.setPotential| (type I, the stillinger weberPairTerm) | the stillinger weberPairTerm | the stil$ type2,

po-

ten-

tial)

Parameters

- type1 -
- type2 -
- potential -

 ${\tt espressopp.interaction.} \textbf{VerletListAdressStillingerWeberPairTerm} \ (vl. \ \textit{fixedtu-}$ pleList)

Parameters

- v1 -
- fixedtupleList -

espressopp.interaction.VerletListAdressStillingerWeberPairTerm.setPotentialAT(type1, type2,

poten-

tial)

Parameters

- type1 -
- type2 -
- potential -

 ${\tt espressopp.interaction.VerletListAdressStillingerWeberPairTerm.} {\tt setPotentialCG} \ ({\it typeI},$ type2,

po-

tential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListHadressStillingerWeberPairTerm(vl, fixedtupleList)

Parameters

- v1 -
- fixedtupleList -

 ${\tt espressopp.interaction.} Verlet {\tt ListHadressStillingerWeberPairTerm.} {\tt setPotentialAT} (\textit{type } l, \textit{type } l, \textit{typ$

type2,

po-

ten-

tial)

Parameters

- type1 -
- type2 -
- potential -

 ${\tt espressopp.interaction.} Verlet {\tt ListHadressStillingerWeberPairTerm.} \textbf{setPotentialCG} (\textit{type1}, \textit{type1}, \textit{type1}, \textit{type2}, \textit{type3}, \textit{type3},$

type2, po-

ten-

tial)

Parameters

- type1 -
- type2 -
- potential -

 $\verb|espressopp.interaction.CellListStillingerWeberPairTerm| (stor)$

Parameters stor -

espressopp.interaction.CellListStillingerWeberPairTerm.setPotential(type1,

type2,

po-

ten-

tial)

Parameters

• type1 -

- type2 -
- potential -

espressopp.interaction.FixedPairListStillingerWeberPairTerm(system, vl, potential)

Parameters

- system -
- v1 -
- potential -

espressopp.interaction.FixedPairListStillingerWeberPairTerm.setPotential(potential)

Parameters potential -

class espressopp.interaction.StillingerWeberPairTerm.StillingerWeberPairTerm
The Lennard-Jones potential.

4.20.39 espressopp.interaction.StillingerWeberPairTermCapped

This class provides methods to compute forces and energies of 2 body term of Stillinger-Weber potential.

If the distance is smaller than the cap-radius:

$$U = A[d_{12}^{-p}(B-1)]e^{\frac{1}{d_{12}-r_c}}$$

where r_c is the cutoff-radius.

espressopp.interaction. StillingerWeberPairTermCapped (A, B, p, q, epsilon, sigma, cutoff, caprad)

Parameters

- A -
- B –
- p -
- q-
- epsilon (real) (default: 1.0)
- **sigma** (real) (default: 1.0)
- **cutoff** (default: infinity)
- **caprad** (real) (default: 0.0)

espressopp.interaction.VerletListStillingerWeberPairTermCapped(vl)

Parameters v1 -

espressopp.interaction.VerletListStillingerWeberPairTermCapped.getCaprad()

Return type

espressopp.interaction. VerletListStillingerWeberPairTermCapped. getPotential (typeI, type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListStillingerWeberPairTermCapped.getVerletList()

Return type A Python list of lists.

```
espressopp.interaction. VerletListStillingerWeberPairTermCapped. setPotential (typeI, type2, potential)
```

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction. VerletListAdressStillingerWeberPairTermCapped (vl, fixed-tu-pleList)

Parameters

- v1 -
- fixedtupleList -

espressopp.interaction.VerletListAdressStillingerWeberPairTermCapped.setPotentialAT (type1 type2

poten-

tial)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListAdressStillingerWeberPairTermCapped.setPotentialCG(typeItype2

poten-

tial)

Parameters

- type1 -
- type2 -
- potential -

fixed-

tupleList)

Parameters

- v1 -
- fixedtupleList -

 $\verb|espressopp.interaction.VerletListHadressStillingerWeberPairTermCapped.setPotentialAT| (type the context of the context of$

type po-

tential

Parameters

- type1 -
- type2 -
- potential -

 $\verb|espressopp.interaction.VerletListHadressStillingerWeberPairTermCapped.\textbf{setPotentialCG}| (\textit{type} algorithms) | \textit{type} algorithms | \textit{type} algorithms$

poten-

tial

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.CellListStillingerWeberPairTermCapped(stor)

Parameters stor -

 $\verb|espressopp.interaction.CellListStillingerWeberPairTermCapped.setPotential| (type I, the stilling of the st$

type2,

po-

ten-

tial)

Parameters

- type1 -
- type2 -
- potential -

 $\verb|espressopp.interaction.FixedPairListStillingerWeberPairTermCapped| (\textit{system}, \\$

vl, po-

tential)

Parameters

- system -
- vl -
- potential -

espressopp.interaction.FixedPairListStillingerWeberPairTermCapped.setPotential(potential)

Parameters potential -

class espressopp.interaction.StillingerWeberPairTermCapped.StillingerWeberPairTermCapped
 The Lennard-Jones potential.

4.20.40 espressopp.interaction.StillingerWeberTripleTerm

This class provides methods to compute forces and energies of the Stillinger Weber Triple Term potential.

if
$$d_{12} >= r_{c_1}$$
 or $d_{32} >= r_{c_2}$

$$U = 0.0$$

else

$$U = \varepsilon \lambda e^{\frac{\sigma \gamma_1}{|r_{12}| - \sigma r_{c_1}}} + \frac{\sigma \gamma_2}{|r_{32}| - \sigma r_{c_2}} \left(\frac{r_{12} r_{32}}{|r_{12}| \cdot |r_{32}|} - \cos(\theta_0) \right)^2$$

```
espressopp.interaction.StillingerWeberTripleTerm(gamma, theta0, lmbd, epsilon,
                                                                                                                                                                      sigma, cutoff)
                          Parameters
                                         • gamma (real) - (default: 0.0)
                                         • theta0 (real) - (default: 0.0)
                                          • lmbd (real) – (default: 0.0)
                                         • epsilon (real) – (default: 1.0)
                                          • sigma (real) - (default: 1.0)
                                         • cutoff – (default: infinity)
espressopp.interaction.VerletListStillingerWeberTripleTerm(system, vl3)
                          Parameters
                                         • system -
                                          • v13 -
\verb|espressopp.interaction.VerletListStillingerWeberTripleTerm.getPotential| (type I, to be a substitution of the context of t
                                                                                                                                                                                                                                                    type2,
                                                                                                                                                                                                                                                    type3)
                          Parameters
                                         • type1 -
                                         • type2 -
                                          • type3 -
                          Return type
\verb|espressopp.interaction.VerletListStillingerWeberTripleTerm.getVerletListTriple()|
                          Return type A Python list of lists.
espressopp.interaction.VerletListStillingerWeberTripleTerm.setPotential(type1,
                                                                                                                                                                                                                                                    type2,
                                                                                                                                                                                                                                                    type3,
                                                                                                                                                                                                                                                    po-
                                                                                                                                                                                                                                                    ten-
                                                                                                                                                                                                                                                    tial)
                          Parameters
                                         • type1 -
                                          • type2 -
                                         • type3 -
                                         • potential -
espressopp.interaction.FixedTripleListStillingerWeberTripleTerm (system, ftl,
                                                                                                                                                                                                                        potential)
                          Parameters
                                         • system -
                                         • ftl -
                                         • potential -
espressopp.interaction.FixedTripleListStillingerWeberTripleTerm.getFixedTripleList()
```

tential)

Return type A Python list of lists.

```
\verb|espressopp.interaction.FixedTripleListStillingerWeberTripleTerm.setPotential| (typeI,
                                                                                          type2,
                                                                                          type3,
                                                                                          po-
```

Parameters

- type1 -
- type2 -
- type3 -
- potential -

class espressopp.interaction.StillingerWeberTripleTerm.StillingerWeberTripleTerm The StillingerWeberTripleTerm potential.

4.20.41 espressopp.interaction.Tabulated

espressopp.interaction.**Tabulated**(*itype*, *filename*, *cutoff*)

Parameters

- itype -
- filename -
- **cutoff** (default: infinity)

espressopp.interaction.VerletListAdressTabulated(vl, fixedtupleList)

Parameters

- v1 -
- fixedtupleList -

espressopp.interaction.VerletListAdressTabulated.setPotentialAT(typeI, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

 $\verb|espressopp.interaction.VerletListAdressTabulated.setPotentialCG| (type I,$

type2,

potential)

Parameters

- type1 -
- type2 -
- potential -

 $\verb|espressopp.interaction.VerletListHadressTabulated| (vl, fixed tuple List)|$

Parameters

• v1 -

• fixedtupleList -

espressopp.interaction.VerletListHadressTabulated.setPotentialAT(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction. VerletListHadressTabulated. setPotentialCG (typeI, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListTabulated(vl)

Parameters v1 -

espressopp.interaction.VerletListTabulated.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListTabulated.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.CellListTabulated(stor)

Parameters stor -

espressopp.interaction.CellListTabulated.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

 $\verb|espressopp.interaction.FixedPairListTabulated| (\textit{system}, \textit{vl}, \textit{potential})$

Parameters

- system -
- v1 -
- potential -

```
espressopp.interaction.FixedPairListTabulated.setPotential(potential)
         Parameters potential -
espressopp.interaction.FixedPairListTypesTabulated(system, ftl)
         Parameters
              • system (espressopp.System) - The Espresso++ system object.
              • ftl (espressopp.FixedPairList) - The FixedPair list.
espressopp.interaction.FixedPairListTypesTabulated.setPotential(typeI,
                                                                            type2,
                                                                            potential)
    Defines bond potential for interaction between particles of types type1-type2-type3.
         Parameters
              • type1 (int) – Type of particle 1.
              • type2 (int) – Type of particle 2.
              • potential (espressopp.interaction.Potential) - The potential to set
class espressopp.interaction.Tabulated.Tabulated
    The Tabulated potential.
4.20.42 espressopp.interaction.TabulatedAngular
espressopp.interaction.TabulatedAngular(itype, filename)
         Parameters
              • itype (int) – The interpolation type: 1 - linear, 2 - akima spline, 3 - cubic spline
              • filename (str) – The tabulated potential filename.
espressopp.interaction.FixedTripleListTabulatedAngular(system, ftl, potential)
         Parameters
              • system (espressopp.System) - The Espresso++ system object.
              • ftl (espressopp.FixedTripleList) - The FixedTripleList.
              • potential (espressopp.interaction.Potential) - The potential.
espressopp.interaction.FixedTripleListTabulatedAngular.setPotential(potential)
         Parameters potential (espressopp.interaction.Potential) - The potential
             object.
espressopp.interaction.FixedTripleListTypesTabulatedAngular(system, ftl)
         Parameters
              • system (espressopp.System) – The Espresso++ system object.
```

• ftl (espressopp.FixedTripleList) - The FixedTriple list.

espressopp.interaction.FixedTripleListTypesTabulatedAngular.setPotential(type1, type2, type3, po-ten-

tial)

Defines angular potential for interaction between particles of types type1-type2-type3.

Parameters

- type1 (int) Type of particle 1.
- type2 (int) Type of particle 2.
- type3 (int) Type of particle 3.
- potential (espressopp.interaction.AngularPotential) The potential to set up.

 ${\bf class} \ {\bf espressopp.interaction.} Tabulated {\bf Angular.} {\bf Tabulated Angular potential.}$

4.20.43 espressopp.interaction.TabulatedDihedral

espressopp.interaction.**TabulatedDihedral**(itype, filename)

Parameters itype – The interpolation type: 1 - linear, 2 - akima spline, 3 - cubic spline :param filename: The tabulated potential filename. :type itype: int :type filename: str

espressopp.interaction.FixedQuadrupleListTabulatedDihedral (system, fql, potential)

Parameters

- **system** (espressopp.System) The Espresso++ system object.
- fql (espressopp.FixedQuadrupleList) The FixedQuadrupleList.
- potential (espressopp.interaction.Potential) The potential.

espressopp.interaction.FixedQuadrupleListTabulatedDihedral.setPotential(potential)

Parameters potential (espressopp.interaction.Potential) - The potential object.

espressopp.interaction. Fixed Quadruple List Types Tabulated Dihedral (system, fql)

Parameters

- **system** (espressopp.System) The Espresso++ system object.
- ftl (espressopp.FixedQuadrupleList) The FixedQuadrupleList list.

espressopp.interaction.FixedQuadrupleListTypesTabulatedDihedral(system, ftl)

Parameters

- **system** (espressopp.System) The Espresso++ system object.
- ftl (espressopp.FixedQuadrupleList) The FixedQuadruple list.

 $\verb|espressopp.interaction.FixedQuadrupleListTypesTabulatedDihedral.setPotential| (type I, type 2, typ$

type3, type4,

poten-

tial)

Defines dihedral potential for interaction between particles of types type1-type2-type3-type4.

Parameters

- type1 (int) Type of particle 1.
- type2 (int) Type of particle 2.
- type3 (int) Type of particle 3.
- type4 (int) Type of particle 4.

• potential (espressopp.interaction.DihedralPotential) - The potential to set up.

class espressopp.interaction.TabulatedDihedral.TabulatedDihedral
 The TabulatedDihedral potential.

4.20.44 espressopp.interaction.TersoffPairTerm

This class provides methods to compute forces and energies of 2 body term of Tersoff potential.

if
$$d_{12} > R + D$$

$$U = 0$$

if $d_{12} < R - D$

$$U = Ae^{-\lambda 1d_{12}}$$

else

$$U = \frac{1}{2} \left(1 - \sin \left(\frac{\pi}{4D} (d_{12} - R) \right) \right) A e^{-\lambda_1 d_{12}}$$

espressopp.interaction.TersoffPairTerm(A, lambda1, R, D, cutoff)

Parameters

- A –
- lambda1 -
- R -
- D -
- **cutoff** (default: infinity)

espressopp.interaction.VerletListTersoffPairTerm(vl)

Parameters v1 -

espressopp.interaction.VerletListTersoffPairTerm.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

 $\verb|espressopp.interaction.VerletListTersoffPairTerm.getVerletList||()$

Return type A Python list of lists.

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.CellListTersoffPairTerm(stor)

Parameters stor -

espressopp.interaction.CellListTersoffPairTerm.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.FixedPairListTersoffPairTerm(system, vl, potential)

Parameters

- system -
- v1 -
- potential -

 $\verb|espressopp.interaction.FixedPairListTersoffPairTerm.\textbf{setPotential}| (\textit{potential})|$

Parameters potential -

class espressopp.interaction.TersoffPairTerm.TersoffPairTerm
The Lennard-Jones potential.

4.20.45 espressopp.interaction.TersoffTripleTerm

This class provides methods to compute forces and energies of the Tersoff Triple Term potential.

$$U = f_{C_j} f_A \left(1 + \left(\beta f_{C_k} \gamma \left(1 + \frac{c_2}{d_2} - \frac{c_2}{d_2 + \left(\frac{r_{12} r_{32}}{|r_{12}||r_{32}|} - \cos(\theta_0) \right)^2} \right) \left(e^{\lambda_3 (|r_{12}| - |r_{32}|)} \right)^m \right)^{n} \right)^{-\frac{1}{2n}}$$

espressopp.interaction.VerletListTersoffTripleTerm(system, vl3)

Parameters

- system -
- v13 -

Parameters

- type1 -
- type2 -
- type3 -

Return type

 $\verb|espressopp.interaction.VerletListTersoffTripleTerm.getVerletListTriple()|$

Return type A Python list of lists.

espressopp.interaction.VerletListTersoffTripleTerm.setPotential(type1, type2,

type3,

potential)

Parameters

• type1 -

```
• type2 -
```

- type3 -
- potential -

espressopp.interaction.FixedTripleListTersoffTripleTerm(system, ftl, potential)

Parameters

- system -
- ft1 -
- potential -

 $\verb|espressopp.interaction.FixedTripleListTersoffTripleTerm.getFixedTripleList(|)|$

Return type A Python list of lists.

 $\verb|espressopp.interaction.FixedTripleListTersoffTripleTerm.setPotential| (\textit{type1}, \textit{type1}, \textit{type2}, \textit{type1}, \textit{type2}, \textit{type3}, \textit$

type2,

type3,

po-

ten-

tial)

Parameters

- type1 -
- type2 -
- type3 -
- potential -

4.20.46 espressopp.interaction.VSpherePair

This class provides methods to compute forces and energies of the VSpherePair potential.

$$V(r_{ij}, \sigma_{ij}) = \frac{\varepsilon}{\beta} \left(\frac{2\pi}{3}\right) \sigma_{ij}^{-\frac{3}{2}} e^{-\frac{3}{2} \frac{r_{ij}^2}{\sigma_{ij}}}, r_{ij} = |\vec{r_i} - \vec{r_j}|, \sigma_{ij} = \sigma_i^2 + \sigma_j^2$$

espressopp.interaction.**VSpherePair**(*epsilon*, *cutoff*, *shift*)

Parameters

- epsilon (real) (default: 1.0)
- cutoff (default: infinity)
- **shift** (default: "auto")

espressopp.interaction.VerletListVSpherePair(vl)

Parameters v1 -

espressopp.interaction.VerletListVSpherePair.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListVSpherePair.getVerletList()

Return type A Python list of lists.

espressopp.interaction.VerletListVSpherePair.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

class espressopp.interaction.VSpherePair.VSpherePair
The Lennard-Jones potential.

4.20.47 espressopp.interaction.VSphereSelf

This class provides methods to compute forces and energies of the VSphereSelf potential.

$$U = e_1 \left(\frac{4}{3}\pi\sigma_2\right)^{\frac{3}{2}} + \frac{a_1 N_b^3}{\sigma_2^3} + \frac{2a_2}{N_b}\sigma_2$$

espressopp.interaction.**VSphereSelf** (e1, a1, a2, Nb, cutoff, shift)

Parameters

- **e1** (real) (default: 0.0)
- **a1** (real) (default: 1.0)
- **a2** (real) (default: 0.0)
- **Nb** (*int*) (default: 1)
- cutoff (default: infinity)
- **shift** (real) (default: 0.0)

espressopp.interaction.SelfVSphere(system, potential)

Parameters

- system -
- potential -

espressopp.interaction.SelfVSphere.getPotential()

Return type

espressopp.interaction.SelfVSphere.setPotential(potential)

Parameters potential -

class espressopp.interaction.VSphereSelf.VSphereSelf
The VSphereSelf potential.

4.20.48 espressopp.interaction.Zero

This class provides methods for a zero potential no interactions between particles, mainly used for debugging and testing

```
espressopp.interaction.Zero()
espressopp.interaction.VerletListZero(vl)
```

Parameters v1 -

espressopp.interaction.VerletListZero.getPotential(type1, type2)

Parameters

- type1 -
- type2 -

Return type

espressopp.interaction.VerletListZero.setFixedTupleList(ftpl)

Parameters ftpl -

espressopp.interaction.VerletListZero.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListAdressZero(vl)

Parameters v1 -

 $\verb|espressopp.interaction.VerletListAdressZero.setFixedTupleList| (ftpl)$

Parameters ftpl -

espressopp.interaction.VerletListAdressZero.setPotentialAT(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListAdressZero.setPotentialCG(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListHadressZero(vl, fixedtupleList)

Parameters

- **vl** -
- fixedtupleList -

 $\verb|espressopp.interaction.VerletListHadressZero.setFixedTupleList| (ftpl)$

Parameters ftpl -

espressopp.interaction.VerletListHadressZero.setPotentialAT(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.VerletListHadressZero.setPotentialCG(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.CellListZero(stor)

Parameters stor -

espressopp.interaction.CellListZero.setPotential(type1, type2, potential)

Parameters

- type1 -
- type2 -
- potential -

espressopp.interaction.FixedPairListZero (system, vl, potential)

Parameters

- system -
- v1 -
- potential -

espressopp.interaction.FixedPairListZero.setPotential(potential)

Parameters potential -

class espressopp.interaction.Zero.Zero
The Zero potential.

4.20.49 espressopp.interaction.LennardJones93Wall

This class defines a Lennard-Jones 9-3 SingleParticlePotential in the direction x.

$$V(r) = \epsilon \left(\left(\frac{\sigma}{r} \right)^9 - \left(\frac{\sigma}{r} \right)^3 \right)$$

where r is the distance from the lower or upper wall in the x direction. V(r) = 0 after a distance sigmaCutoff.

The parameters have to be defined for every species present in the system with *setParams* and can be retrieved with *getParams*.

Example:

```
>>> LJ93 = espressopp.interaction.LennardJones93Wall()
>>> LJ93.setParams(0, 6., 1., wall_cutoff)
>>> SPLJ93 = espressopp.interaction.SingleParticleLennardJones93Wall(system, LJ93)
>>> system.addInteraction(SPLJ93)
```

espressopp.interaction.LennardJones93Wall()

espressopp.interaction.LennardJones93Wall.getParams(type_var)

Parameters type_var -

Return type

espressopp.interaction.LennardJones93Wall.setParams($type_var$, epsilon, sigma, sigmaCutoff, r0)

Parameters

```
• type_var -
```

- epsilon -
- sigma -
- sigmaCutoff -
- r0 -

espressopp.interaction.SingleParticleLennardJones93Wall (system, potential)

Parameters

- system -
- potential -

espressopp.interaction.SingleParticleLennardJones93Wall.setPotential(potential)

Parameters potential -

class espressopp.interaction.LennardJones93Wall.LennardJones93Wall
 The LennardJones93Wall potential.

4.21 io

4.21.1 DumpGRO - IO Object

- *dump()* write configuration to trajectory GRO file. By default filename is "out.gro", coordinates are folded. Properties
- filename Name of trajectory file. By default trajectory file name is "out.gro"
- unfolded False if coordinates are folded, True if unfolded. By default False
- append True if new trajectory data is appended to existing trajectory file. By default True
- *length_factor* If length dimension in current system is nm, and unit is 0.23 nm, for example, then length_factor should be 0.23
- length_unit It is length unit. Can be LJ, nm or A. By default LJ

usage:

writing down trajectory

writing down trajectory using ExtAnalyze extension

Both exapmles will give the same result: 200 configurations in trajectory .gro file.

setting up length scale

For example, the Lennard-Jones model for liquid argon with $\sigma = 0.34[nm]$

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```
>>> dump_conf_gro = espressopp.io.DumpGRO(system, integrator, filename='trj.gro',_

ounfolded=False, length_factor=0.34, length_unit='nm', append=True)
```

will produce trj.gro with in nanometers

espressopp.io.DumpGRO (system, integrator, filename, unfolded, length_factor, length_unit, append)

Parameters

- system -
- integrator -
- **filename** (default: 'out.gro')
- unfolded (default: False)
- length_factor (real) (default: 1.0)
- length_unit (default: 'LJ')
- append (default: True)

espressopp.io.DumpGRO.dump()

Return type

4.21.2 DumpXYZ - IO Object

• *dump()* write configuration to trajectory XYZ file. By default filename is "out.xyz", coordinates are folded. DumpXYZ works also for Multiple communicators.

Properties

- filename Name of trajectory file. By default trajectory file name is "out.xyz"
- unfolded False if coordinates are folded, True if unfolded. By default False
- append True if new trajectory data is appended to existing trajectory file. By default True
- *length_factor* If length dimension in current system is nm, and unit is 0.23 nm, for example, then length_factor should be 0.23 Default: 1.0
- length_unit It is length unit. Can be LJ, nm or A. By default LJ
- *store_pids* True if you want to store pids as fastwritexyz does. False otherwise (standard XYZ) Default: False
- *store_velocities* True if you want to store velocities. False otherwise (XYZ doesn't require it) Default: False

usage:

writing down trajectory

writing down trajectory using ExtAnalyze extension

Both exapmles will give the same result: 200 configurations in trajectory .xyz file.

setting up length scale

For example, the Lennard-Jones model for liquid argon with $\sigma = 0.34[nm]$

will produce trj.xyz with in nanometers

```
espressopp.io.DumpXYZ(system, integrator, filename, unfolded,
length_factor, length_unit, store_pids,
store_velocities, append)

param system

param integrator

param filename (default: 'out.xyz')

param unfolded (default: False)

param length_factor (default: 1.0)

param length_unit (default: 'LJ')
```

Parameters

- store_pids (bool) (default: False)
- **store_velocities** (bool :type append:) (default: False) :param append: (default: True) :type system: :type integrator: :type filename: :type unfolded: :type length_factor: real :type length_unit:

espressopp.io.DumpXYZ.dump()

Return type

4.22 standard_system

4.22.1 espressopp.standard_system.Default

espressopp.standard_system.**Default** (box, rc, skin, dt, temperature)

Parameters

- box -
- rc (real) (default: 1.12246)
- **skin** (real) (default: 0.3)
- **dt** (real) (default: 0.005)
- temperature (default: None)

Return default system and integrator, no interactions, no particles are set if tempearture is != None then Langevin thermostat is set to temperature (gamma is 1.0)

4.22.2 espressopp.standard_system.KGMelt

espressopp.standard_system.KGMelt(num_chains, chain_len)

Parameters

- num_chains -
- chain_len -

4.22.3 espressopp.standard_system.LennardJones

espressopp.standard_system.**LennardJones** (num_particles, box, rc, skin, dt, epsilon, sigma, shift, temperature, xyzfilename, xyzrfilename)

Parameters

- num_particles -
- **box** (default: (000))
- rc (real) (default: 1.12246)
- **skin** (real) (default: 0.3)
- **dt** (real) (default: 0.005)
- **epsilon** (*real*) (default: 1.0)
- **sigma** (real) (default: 1.0)
- **shift** (default: 'auto')
- temperature (default: None)
- xyzfilename (default: None)
- xyzrfilename (default: None)

return random Lennard Jones system and integrator: if tempearture is != None then Langevin thermostat is set to temperature (gamma is 1.0)

4.22.4 espressopp.standard_system.Minimal

espressopp.standard_system.Minimal(num_particles, box, rc, skin, dt, temperature)

Parameters

- num_particles -
- box -
- rc(real)-(default: 1.12246)
- **skin** (real) (default: 0.3)
- dt (real) (default: 0.005)
- temperature (default: None)

Return minimal system and integrator whithout any interactions defined: particles have random positions in box if tempearture is != None then Langevin thermostat is set to temperature (gamma is 1.0)

4.22.5 espressopp.standard_system.PolymerMelt

```
espressopp.standard_system.PolymerMelt(num_chains, monomers_per_chain, box, bondlen, rc, skin, dt, epsilon, sigma, shift, temperature, xyzfilename, xyzrfilename)
```

Parameters

```
• num chains -
```

• monomers_per_chain -

• **box** – (default: (000))

• **bondlen** (real) – (default: 0.97)

• rc (real) - (default: 1.12246)

• **skin** (real) – (default: 0.3)

• **dt** (real) – (default: 0.005)

• epsilon (real) – (default: 1.0)

• **sigma** (real) - (default: 1.0)

• **shift** – (default: 'auto')

• temperature - (default: None)

• xyzfilename – (default: None)

• xyzrfilename - (default: None)

returns random walk polymer melt system and integrator: if tempearture is != None then Langevin thermostat is set to temperature (gamma is 1.0)

4.23 storage

4.23.1 espressopp.storage.DomainDecomposition

espressopp.storage.DomainDecomposition(system, nodeGrid, cellGrid)

Parameters

- system -
- nodeGrid -
- cellGrid -

espressopp.storage.DomainDecomposition.getCellGrid()

Return type

espressopp.storage.DomainDecomposition.getNodeGrid()

Return type

4.23.2 DomainDecompositionAdress - Object

The DomainDecompositionAdress is the Domain Decomposition for AdResS and H- AdResS simulations. It makes sure that tuples (i.e. a coarse-grained particle and its corresponding atomistic particles) are always stored together on one CPU. When setting DomainDecompositionAdress you have to provide the system as well as the nodegrid and the cellgrid.

Example - setting DomainDecompositionAdress:

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```
>>> system.storage = espressopp.storage.DomainDecompositionAdress(system, nodeGrid, cellGrid)
```

espressopp.storage.DomainDecompositionAdress(system, nodeGrid, cellGrid)

Parameters

- system -
- nodeGrid -
- cellGrid -

4.23.3 espressopp.storage.DomainDecompositionNonBlocking

espressopp.storage.DomainDecompositionNonBlocking (system, nodeGrid, cellGrid)

Parameters

- system -
- nodeGrid -
- cellGrid -

4.23.4 Storage - Storage Object

This is the base class for all storage objects. All derived classes implement at least the following methods:

• decompose()

Send all particles to their corresponding cell/cpu

• addParticle(pid, pos):

Add a particle to the storage

• removeParticle(pid):

Remove a particle with id number *pid* from the storage.

```
>>> system.storage.removeParticle(4)
```

There is an example in examples folder

• getParticle(pid):

Get a particle object. This can be used to get specific particle information:

```
>>> particle = system.storage.getParticle(15)
>>> print "Particle ID is : ", particle.id
>>> print "Particle position is : ", particle.pos
```

you cannot use this particle object to modify particle data. You have to use the modifyParticle command for that (see below).

• addAdrParticle(pid, pos, last_pos):

Add an AdResS Particle to the storage

- *setFixedTuplesAdress(fixed_tuple_list)*:
- addParticles(particle_list, *properties):

This routine adds particles with certain properties to the storage.

param particleList list of particles (and properties) to be added

param properties property strings

Each particle in the list must be itself a list where each entry corresponds to the property specified in properties.

Example:

```
>>> addParticles([[id, pos, type, ...], 'id', 'pos', 'type', ...

→)
```

• modifyParticle(pid, property, value, decompose='yes')

This routine allows to modify any properties of an already existing particle.

Example:

```
>>> modifyParticle(pid, 'pos', Real3D(new_x, new_y, new_z))
```

• removeAllParticles():

This routine removes all particles from the storage.

• 'system':

The property 'system' returns the System object of the storage.

Examples:

espressopp.storage.Storage.addAdrATParticle(pid, *args)

Parameters

- pid-
- *args -

Return type

espressopp.storage.Storage.addParticle(pid, pos)

Parameters

- pid-
- pos -

Return type

espressopp.storage.Storage.addParticles(particleList, *properties)

Parameters

- particleList -
- *properties -

Return type

```
espressopp.storage.Storage.clearSavedPositions()
```

Return type

espressopp.storage.Storage.getParticle(pid)

Parameters pid-

Return type

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```
espressopp.storage.Storage.getRealParticleIDs()
        Return type
espressopp.storage.Storage.modifyParticle(pid, property, value)
        Parameters
             • pid-
             • property -
             • value -
        Return type
espressopp.storage.Storage.particleExists(pid)
        Parameters pid-
        Return type
espressopp.storage.Storage.printRealParticles()
        Return type
espressopp.storage.Storage.removeAllParticles()
        Return type
espressopp.storage.Storage.removeParticle(pid)
        Parameters pid-
        Return type
espressopp.storage.Storage.restorePositions()
        Return type
espressopp.storage.Storage.savePositions(idList)
        Parameters idList -
        Return type
espressopp.storage.Storage.setFixedTuplesAdress(fixedtuples)
        Parameters fixedtuples -
```

4.24 Logging mechanism

ESPResSo++ uses Loggers

Logging can be switched on in your python script with the following command:

```
>>> logging.getLogger("*name of the logger*").setLevel(logging.*Level*)
```

Level is one of the following:

| ERROR | for errors that might still allow the application to continue |
|-------|---|
| WARN | for potentially harmful situations |
| INFO | informational messages highlighting progress |
| DEBUG | designates fine-grained informational events |

Example:

```
>>> import espressopp
>>> import logging
>>> logging.getLogger("Storage").setLevel(logging.ERROR)
```

To log everything (WARNING: this will produce **lots** of output):

>>> logging.getLogger("").setLevel(logging.DEBUG)

The following loggers are currently available:

- Configurations
- Observable
- Velocities
- BC
- Logger
- FixedListComm
- FixedPairList
- FixedQuadrupleList
- FixedTripleList
- FixedTupleList
- Langevin
- MDIntegrator
- AngularPotential
- DihedralPotential
- Interaction
- InterpolationAkima
- InterpolationCubic
- InterpolationLinear
- InterpolationTable
- Potential
- CellListAllPairsIterator
- DomainDecomposition.CellGrid
- DomainDecomposition
- DomainDecomposition.NodeGrid
- Storage
- DomainDecompositionAdress
- StorageAdress
- VerletList
- VerletList

FREQUENTLY ASKED QUESTIONS

Do I need to learn Python when using ESPResSo++?

The short answer is "no". Most of the example scripts are self-explanatory and can be adapted for your purposes by simple changes. You can also use ESPResSo++ like other MD simulation software, that is driven by some kind of configuration file.

The long answer is "yes". If you want to take advantage of all features of ESPResSo++ you need some knowledge of how the Python interpreter works.

But don't be afraid of learning Python:

- Python is easy to learn
- The ESPResSo++ example simulation scripts gives you a very fast insight of how Python works.
- Writing programs in Python is much easier than writing programs in C++
- Python programs are easier to read than Tcl or Perl programs.

And here are some arguments why it is worth while:

- There are many Python programs you can use in your applications
- Python gives you a flexible way of running MD simulations with ESPResSo++

Do you support other script languages, e.g. Tcl/Tk?

No. We choose the support only Python as ESPResSo++ scripting language. This enables ESPResSo++ users to read and adapt scripts written by other ESPResSo++ users.

Can Tcl scripts converted to Python automatically?

The recommendation is - don't do it! Instead, a Tcl interpreter can be loaded via Python and given the job to do. That is similar to what Tkinter does; Tkinter is a wrapper to use the Tk toolkit from Python.

Why should I use Python if C++ programs are much faster?

Python is the driver of your simulation which will still run in the ESPResSo++ C++ engine.

Python programs are about 30 to 50 times slower than the same programs written in C++.

That is why we use Python to set up and control simulations while the simulation system itself is written in efficient C++ code.

Can I run ESPResSo++ on parallel machines?

Yes. The parallel version uses MPI and is therefore as portable as MPI is. Typical MD simulations scale rather well.

Do I need to write parallel scripts for parallel machines?

No. The Python scripts are executed only by the first processor which will broadcast the ESPResSo++ commands to the other processors automatically using the PMI interface (Parallel Method Invocation). For you, it will look like a serial script. But the particles of the simulation are distributed among the available processors and the commands issued for ESPResSo++ will be executed by each processor.

How efficient is ESPResSo++?

Efficiency is a high priority though less than the extendability of the system. You should expect a good performance but might be that ESPResSo++ is less efficient than other simulation programs that are around.

If you experience that ESPResSo++ is more than 2 times slower than other simulation systems you have found a performance bug.

Do I need the source code distribution or can I use binary version?

Currently, we only provide a source code distribution. This might change in the future. Our major problem to provide binary versions is that there are many different Python versions and the binary versions of the ESPResSo++ and python libraries must not be of mixed versions.

Which build systems are used for ESPResSo++?

Compilation and installation of ESPResSo++ is due to the many shared libraries and loadable modules rather complex and so we use a build system to make our job and maintainability easier.

Currently we support building the system with cmake.

What means extendability?

Each software is in a certain sense extendable by adding some functionality somewhere in the code. But we understand extendability in the following sense:

• You can add functionality without changing existent interfaces. For the object-oriented approach this means in practice: take an available base class and define a new derived class with your needed functionality.

CHAPTER

SIX

GETTING HELP

We currently have two mailing lists that provide help for users and developers of ESPResSo++

If you are a user please subscribe to ESPResSo++ Users List

If you are a developer please subscribe to ESPResSo++ Developers List

SEVEN

DEVELOPER TEAM

Current developers:

- Torsten Stuehn (Max Planck Institute for Polymer Research, Germany)
- Horacio Vargas Guzman (Max Planck Institute for Polymer Research, Germany)
- Nikita Tretyakov (Max Planck Institute for Polymer Research, Germany)
- Aoife Fogarty (Max Planck Institute for Polymer Research, Germany)
- Karsten Kreis (Max Planck Institute for Polymer Research, Germany)
- Jakub Krajniak (KU Leuven, Belgium)
- Pierre de Buyl (KU Leuven, Belgium)
- Christoph Junghans (Los Alamos National Laboratory, USA)
- Lothar Brombacher (Max Planck Institute for Polymer Research, Germany)
- Marc Radu (Max Planck Institute for Polymer Research, Germany)

Former developers:

- Axel Arnold (Institute for Computational Physics, Uni-Stuttgart, Germany)
- Stas Bevc (National Institute of Chemistry, Slovenia)
- Thomas Brandes (Fraunhofer Institute SCAI, Germany)
- Sebastian Fritsch (Max Planck Institute for Polymer Research, Germany)
- Jonathan Halverson (Brookhaven National Laboratory, USA)
- Elena Hoemann (Max Planck Institute for Polymer Research, Germany)
- Konstantin Koschke (Max Planck Institute for Polymer Research, Germany)
- Olaf Lenz (Institute for Computational Physics, Uni-Stuttgart, Germany)
- Livia Moreira (Max Planck Institute for Polymer Research, Germany)
- Dirk Reith (Fraunhofer Institute SCAI, Germany)
- Victor Ruehle (University of Cambridge, UK)
- Vitalii Starchenko (Max Planck Institute for Polymer Research, Germany)

CHAPTER **EIGHT**

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