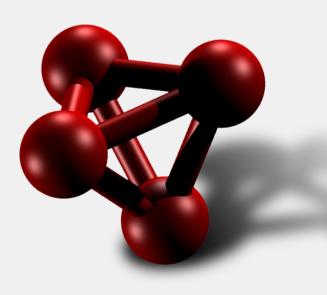
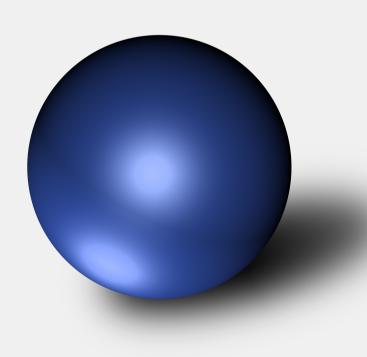
ESPResSo Summer School 2012

ESPResSo++ and AdResS



Staš Bevc

National Institute of Chemistry, Slovenia



Acknowledgements

 Special thanks to the ESPResSo++ developer team

Current developers:

Torsten Stuehn (Max Planck Institute for Polymer Research, Germany) Vitalii Starchenko (Max Planck Institute for Polymer Research, Germany) Konstantin Koschke (Max Planck Institute for Polymer Research, Germany) Livia Moreira (Max Planck Institute for Polymer Research, Germany) Raffaello Potestio (Max Planck Institute for Polymer Research, Germany) Karsten Kreis (Max Planck Institute for Polymer Research, Germany)

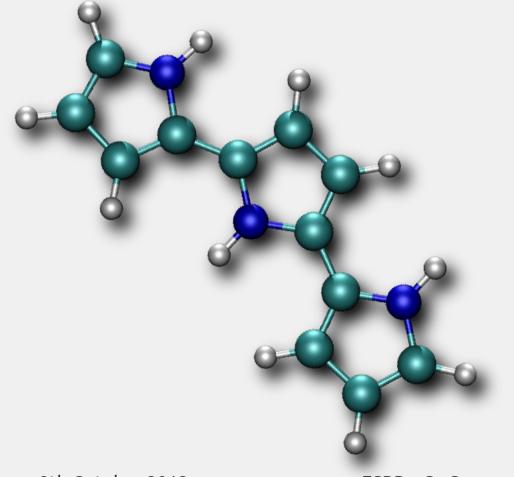
Former developers:

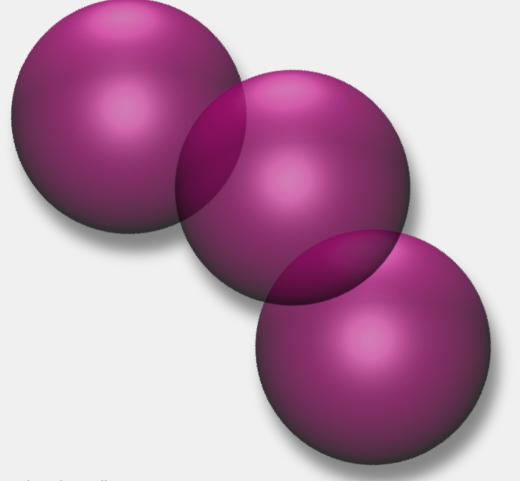
Thomas Brandes (Fraunhofer Institute SCAI, Germany) Dirk Reith (Fraunhofer Institute SCAI, Germany) Jonathan Halverson (Brookhaven National Laboratory, USA) Axel Arnold (Institute for Computational Physics, Uni-Stuttgart, Germany) Olaf Lenz (Institute for Computational Physics, Uni-Stuttgart, Germany) Christoph Junghans (Los Alamos National Laboratory, USA) Victor Ruehle (University of Cambridge, UK)

Outline

- Introduction to AdResS
- AdResS implementation in ESPResSo++
- Tetrahedral molecule example

Motivation





Motivation

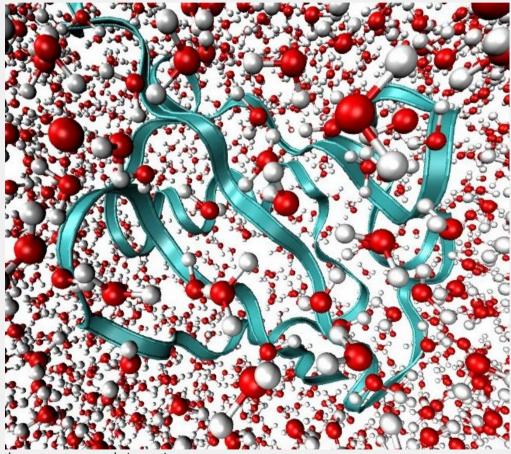
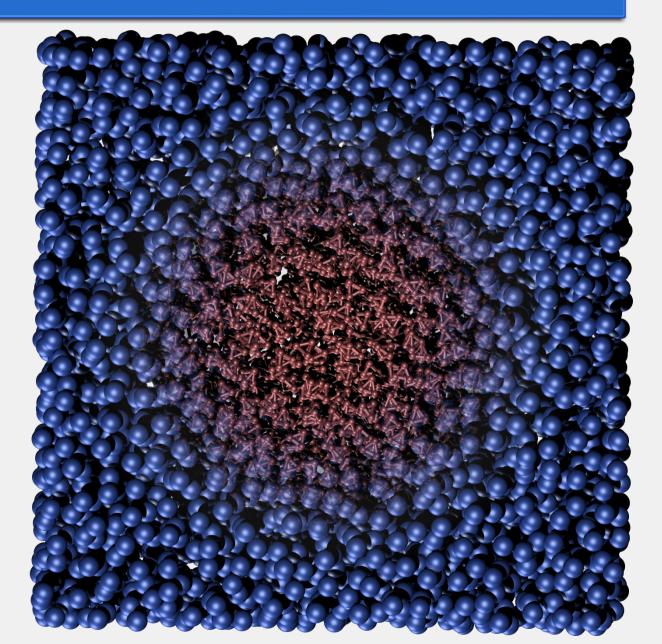
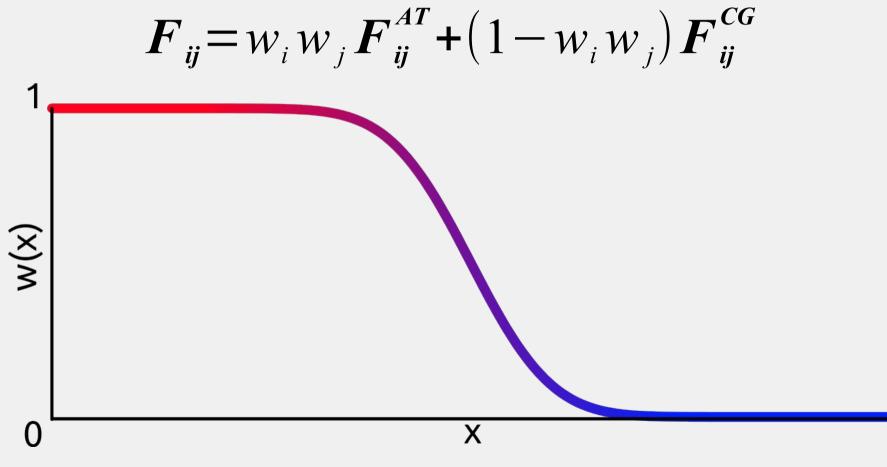


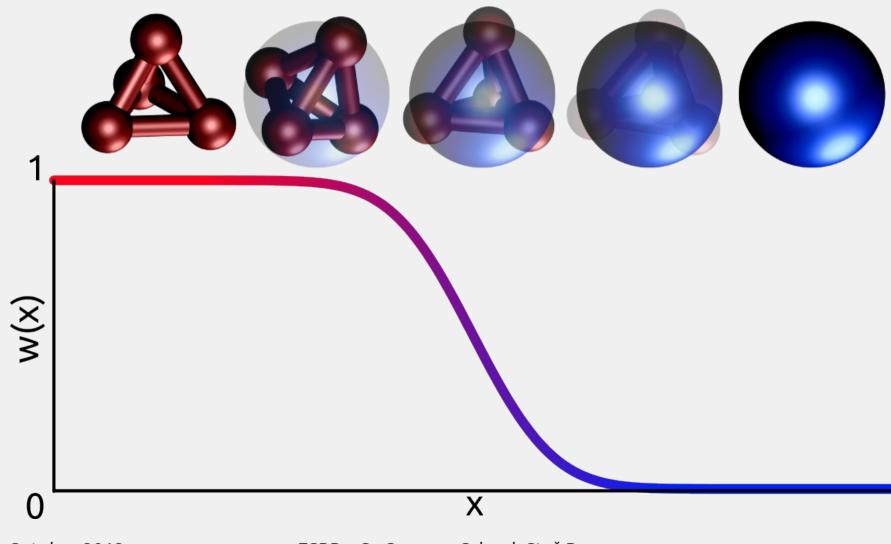
image source: Internet



9th October 2012

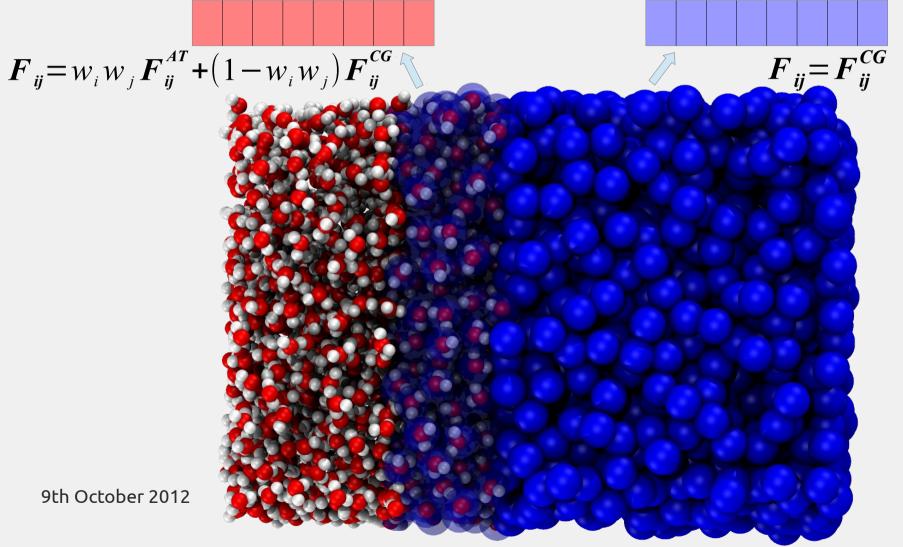
• Force calculation:





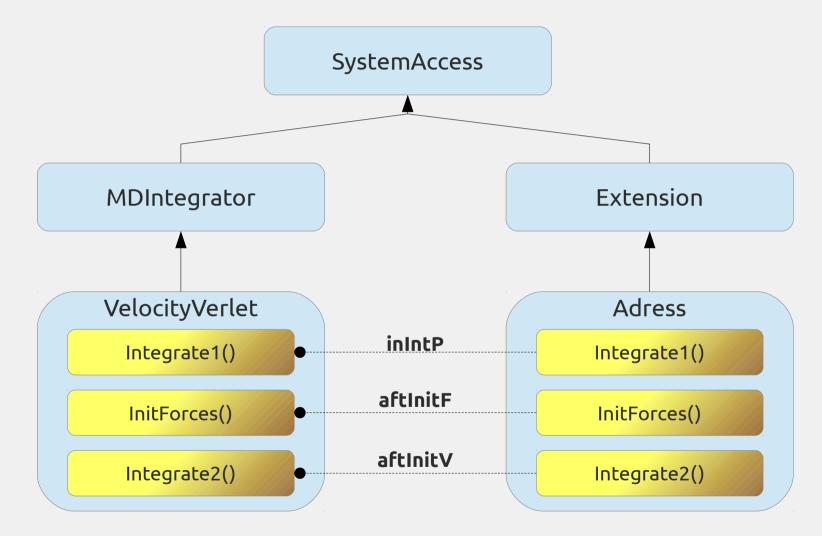
AdResS implementation in ESPResSo++

• General idea: we build two Verlet lists



- General idea: atomistic particles act as an additional property of the coarse grained particles
 - there is a mapping from a CG particle to a list
 of AT particles (via the TupleList)
 std::map<Particle*, std::vector<Particle*> >
 - atomistic particles are always present

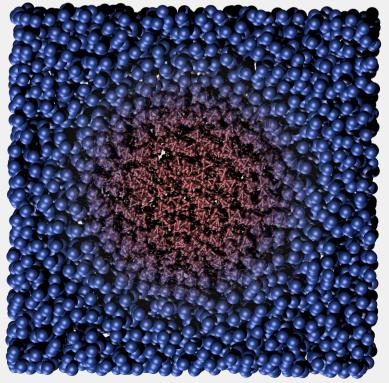
- AdResS is an extension of the integrator
 - but changes to other code and new classes were also introduced
- The new (atomistic) particles have to be integrated
 - we connect to 3 integrator signals:
 - inIntP
 - afterInitF
 - afterIntV



- New classes
 - VerletListAdress
 - VerletListAdressInteractionTemplate
 - DomainDecompositionAdress
 - FixedTupleList
 - FixedPair/Triple/QuadrupleListAdress
- New functions and data structures
 - Storage
 - Particle

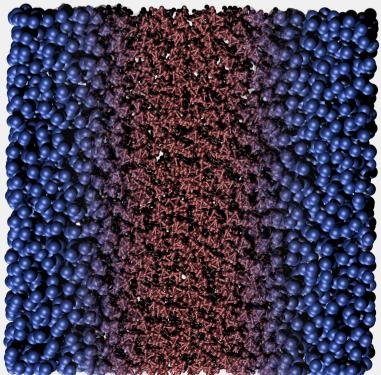
- Spherical geometry by default, but can be planar with minor code changes
- For spherical

Real3D dist = pos() - ref(); real distsq = dist.sqr();



• For planar

real dist = pos()[0] - ref()[0]; real distsq = dist.sqr();



- What about the time?
- Is there any speedup with AdResS?

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• The force calculation is faster

- What about the time?
- Is there any speedup with AdResS?

- The force calculation is faster
- But in total it is slower

Tetrahedral molecule example

AdResS domain decomposition system.storage = espresso.storage.DomainDecompositionAdress(system, nodeGrid, cellGrid)

```
# create CG particles from center of mass
for pidCG in range(num_particlesCG):
  cmp = [0,0,0]
  cmv = [0,0,0]
  tmptuple = [pidCG+num_particles]
  # com calculation
  for pidAT in range(4):
     pid = pidCG*4+pidAT
     tmptuple.append(pid)
     pos = (allParticlesAT[pid])[1]
     vel = (allParticlesAT[pid])[2]
     for i in range(3):
       cmp[i] += pos[i] # masses are 1.0 so we skip
multiplication
       cmv[i] += vel[i]
  for i in range(3):
     cmp[i] /= 4.0 # 4.0 is the mass of molecule
     cmv[i] /= 4.0
```

create CG particles from center of mass
for pidCG in range(num_particlesCG):

. . .

```
allParticles.append([pidCG+num_particles, # CG particle first!
            Real3D(cmp[0], cmp[1], cmp[2]), # pos
            Real3D(cmv[0], cmv[1], cmv[2]), # vel
            Real3D(0, 0, 0), # f
            (0, 4.0, 0]) # type, mass, is not AT particle
for pidAT in range(4):
    pid = pidCG*4+pidAT
    allParticles.append([pid, # now the AT particles can be added
              (allParticlesAT[pid])[1], # pos
              (allParticlesAT[pid])[2], # vel
              (allParticlesAT[pid])[3], # f
              (allParticlesAT[pid])[4], # type
              (allParticlesAT[pid])[5], # mass
              (allParticlesAT[pid])[6]]) # is AT particle
```

create CG particles from center of mass
for pidCG in range(num_particlesCG):

. . .

```
allParticles.append([pidCG+num_particles, # CG particle first!
            Real3D(cmp[0], cmp[1], cmp[2]), # pos
            Real3D(cmv[0], cmv[1], cmv[2]), # vel
            Real3D(0, 0, 0), # f
            (0, 4.0, 0]) # type, mass, is not AT particle
for pidAT in range(4):
    pid = pidCG*4+pidAT
    allParticles.append([pid, # now the AT particles can be added
              (allParticlesAT[pid])[1], # pos
              (allParticlesAT[pid])[2], # vel
              (allParticlesAT[pid])[3], # f
              (allParticlesAT[pid])[4], # type
              (allParticlesAT[pid])[5], # mass
              (allParticlesAT[pid])[6]]) # is AT particle
```

add particles
system.storage.addParticles(allParticles, "id", "pos", "v", "f", "type", "mass",
"adrat")

add tuples
ftpl = espresso.FixedTupleList(system.storage)
ftpl.addTuples(tuples)
system.storage.setFixedTuples(ftpl)

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

AdResS Verlet list
vl = espresso.VerletListAdress(system, cutoff=rc+skin,
adrcut=rc+skin,

dEx=ex_size, dHy=hy_size, adrCenter=[18.42225, 18.42225, 18.42225])

non-bonded potentials # LJ Capped WCA between AT and tabulated Morse between CG particles interNB = espresso.interaction.VerletListAdressLennardJonesCapped(vl, ftpl) potWCA = espresso.interaction.LennardJonesCapped(epsilon=1.0, sigma=1.0, shift=True, caprad=0.27, cutoff=rca) potMorse = espresso.interaction.Tabulated(itype=2, filename=tabMorse, cutoff=rc) # CG interNB.setPotentialAT(type1=1, type2=1, potential=potWCA) # AT interNB.setPotentialCG(type1=0, type2=0, potential=potMorse) # CG system.addInteraction(interNB)

bonded potentials # FENE and LJ potential between AT particles potFENE = espresso.interaction.FENE(K=30.0, r0=0.0, rMax=1.5) potLJ = espresso.interaction.LennardJones(epsilon=1.0, sigma=1.0, shift=True, cutoff=rca) interFENE = espresso.interaction.FixedPairListFENE(system, fpl, potFENE) interLJ = espresso.interaction.FixedPairListLennardJones(system, fpl, potLJ) system.addInteraction(interFENE) system.addInteraction(interLJ)

VV integrator integrator = espresso.integrator.VelocityVerlet(system) integrator.dt = timestep

add AdResS extension
adress = espresso.integrator.Adress(system)
integrator.addExtension(adress)

add Langevin thermostat extension langevin = espresso.integrator.LangevinThermostat(system) langevin.gamma = gamma langevin.temperature = temp langevin.adress = True # enable AdResS! integrator.addExtension(langevin)

- Things to try out
 - Run the simulation with different explicit and hybrid sizes (ex_size, hy_size)
 - Also try CG and AA
 - Compare the times it takes to finish the simulation
 - Modify the code to get 1-dimensional splitting