

Biomolecular modeling

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- ① Total energy with molecular mechanics
 - bonded and non-bonded interactions
 - contributions to the energy
 - parameters and variables in the energy contributions
 - relation of force and energy
- ② Molecular dynamics simulation
 - equations of motion
 - their numerical solution
 - initial conditions
 - Verlet method
 - time step
 - conservation of energy

③ Temperature and MD simulation

- relation of temperature
- kinetic energy and velocities
- scaling of velocities to control temperature
- Berendsen thermostat

④ Setup of molecular system to be simulated

- periodic boundary conditions
- shape of the box
- MM models of water
- treatment of non-bonded interactions (cut-off, Ewald method)

- 5 Analysis of simulation
 - average structure
 - RMSD
 - RMSF
 - radial distribution function
- 6 Coarse-grained simulations
 - in the extent of the slide deck from the class
- 7 Enhanced sampling methods
 - statement of the problem
 - metadynamics
 - principle
 - application of additional energy contribution
 - need of a collective variable
 - free energy as function of CV