

Computational Science

Molecule Dynamic simulation

Task:

Implementation of an MD-simulation for Lennard-Jones particles/ ions/ water in 2D and 3D. →interaction between the particles an thereby causes spatial motion by iterativ calculations

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Concept:

Data storage: ??

Global variables (cutoff, Energieparameter, Tensoren für Speicherung von Zwischenergebnissen Oder nur Umwandlung von Metadata und Ergebnissen), Formate (vectorisation np.arrays), labeling
Parameter (simulation size N, timestep, and total time duration)

Parameters?

Sampling:

creation of particle type (LJ-particle, ion, water), generation of particle positions, assume a distribution of the particle, Microcanonical ensemble (NVE), Canonical ensemble (NVT)?

- Markov Chain Monte Carlo
- Metropolis Monte Carlo

Motion of the particle distribution

Visualisation: plotting, analysing
Mathplotlib

and

Testing
pytest

Distance model:

Calculation of the respective distance (x, y, z) of the particle (N)
→ $x_1:(x_2, \dots, x_N)$ etc.; Periodic/Non-periodic
Short-range interaction:

- Cell list
- Verlet list / Neighbour list

Long-range interaction:

- Ewald summation
- Particle mesh Ewald summation

Potential mode:

Creation of potential/ gradient function an its optimisation

- Coulomb-Potential (-> Particle-Mesh)
- Lennard-Jones
- Particle-mesh Ewald
- Gradient descent

Integrator mode:

numerical method to solve Newton's EOM → to calculate trajectories of particles

- Euler
- Langevin (Thermostat)
- Velocity-verlet/Rungakutta methods (symplectic/ dynamics)

Vectorized particle distances

Potential functions

Vectorized particle positions

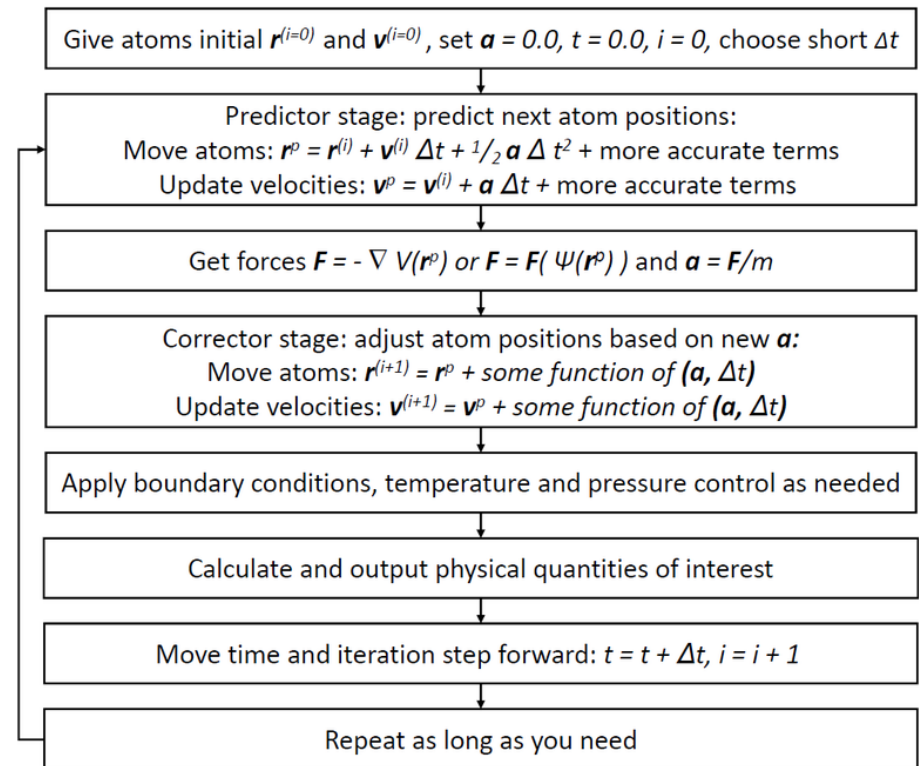
Vectorized energies and forces

Workflow

So, 13.01.19

- finale aufgaben Verteilung
- modules diskutieren
- Labeling festlegen (formate, Bezeichnungen, Abkürzungen, Parameter)
- Input und Output formate und Benennungen definieren
- Ziele festlegen für Fortschritt bis zum nächsten Treffen
- Parallelization strategies ? Domain decomposition method
- (Distribution of system data for parallel computing)
-

Simplified schematic of the molecular dynamics algorithm



Selbstständige Arbeitsphase

- Ziele umsetzen für Fortschritt bis zum nächsten Treffen

So, 20.01.19?

- Fortschritte präsentieren Fehler Probleme diskutieren

-

Overview:

Aufgabe für (Person): Phillip (P), Felix (F), Malte (M), Joana (J)

MODUL	INPUT	TASK	OUTPUT	TESTING	VISUALISATION
DATA STORAGE		? Configuration ?			
SAMPLING 13.01.19 (P, F, M, J)	N = number of particle - generate configuration - -	- configuration ? - Dann in: ? Markov Chain Monte Carlo Metropolis Monte Carlo	2D: Np.array([x_position 1:N],[y_position 1:N]) 3D: Np.array([x_position 1:N],[y_position 1:N],[z_position 1:N])	-	- distribution of particle plotten -
DISTANCE	2D: Np.array([x_position 1:N],[y_position 1:N]) 3D: Np.array([x_position 1:N],[y_position 1:N],[z_position 1:N])		2D: Np.array([x_i 1:N],[dis_x_i+1 1:N],...,[dis_x_N 1:N]); wie y, z 3D:...		
POTENTIAL				- energy consistency - Use simple potential with work and compare	
INTEGRATOR					

Information about modules

DATA STORAGE MODULE

???

SAMPLING MODULE

Initial configuration (markow chain, Particles equispaced in a box; 2D/ 3D)

MONTE CARLO

INPUT	OUTPUT
initial configuration $\mathbf{r}(0)$, potential, size, beta, step	Position \mathbf{r} , potential(\mathbf{r})

(1) Start with initial configuration $\mathbf{r}^{(0)}$ and set $k = 0$.

(2) Fork=1,...,K:

- (a) Sample random vector $\eta \in \mathbb{R}^N$ with $\eta_i \sim N(0, \sigma^2)$, i.e. an isotropic multivariate Gaussian distribution/ or function with variances σ^2 .
- (b) Propose new configuration $\mathbf{r}' = \mathbf{r}^{(k)} + \eta$.

(c) Accept new configuration with probability $p_{\text{acc}} = \min \left\{ 1, \exp \left(\frac{\phi(\mathbf{r}') - \phi(\mathbf{r}^{(k)})}{k_B T} \right) \right\}$

If accepted, set $\mathbf{r}^{(k+1)} = \mathbf{r}'$, otherwise $\mathbf{r}^{(k+1)} = \mathbf{r}^{(k)}$.

Markov Chain Monte Carlo

- create samples from a possibly multi-dimensional continuous random variable, with probability density proportional to a known function.
- these samples can be used to evaluate an integral over that variable, as its expected value or variance.
- in Markov chain Monte Carlo methods are autocorrelated.
- a equilibrium distribution which is proportional to the function given.

Metropolis Monte Carlo

- Konfigurationen gemäß ihrer Boltzmann-Wahrscheinlichkeit + einfachen arithmetischen Durchschnitt
- zuerst eine zufällige Bewegung ausgeführt
- dann die Boltzmann-Wahrscheinlichkeit einer solchen Bewegung ausgewertet wird und die Wahrscheinlichkeit mit einer Zufallszahl verglichen wird.
- Ist die Boltzmann-Wahrscheinlichkeit der Bewegung größer als die Zufallszahl, wird die Bewegung akzeptiert. Andernfalls wird das System auf seine ursprüngliche Konfiguration zurückgesetzt.

Vor- und Nachteile: ?

- not interested in the statistics of momenta anymore, so we integrate them out
- sampling the probability density.
- simple version of Monte Carlo is Metropolis Monte Carlo

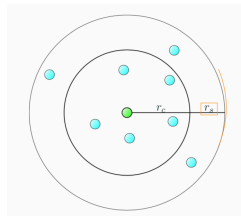
DISTANCE MODULE

INPUT	OUTPUT
Position r , potential(r)	Matrix distances $x_1 \ x_2 \ \dots \ x_N$

- calculation of distances between particles in MD Energy and force Calculation
- Introduce a cutoff and considering only the nearest neighbours within the radius; for LJ.Potential $r_c = 2.5 \sigma$
 - Reduce effort: loop over all pairs of particle $(i,j) = (j,i)$

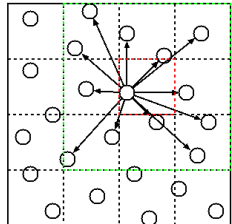
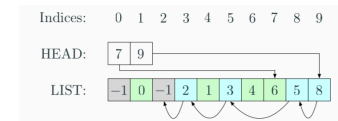
Verlet list

- For each particle i , a Verlet list has lists all other particles j within a smaller distance than $r_c + r_s$
- Update the neighbour list every N_s timesteps
- $r_s > N_s v_{typ} \delta t$ type speed of particles
- Choice between hash table and array



Cell-linked list : get complexity down

- domain into cells with an edge length greater than or equal to the cut-off radius of the interaction to be computed
- particles sorted into cells and the interactions are computed between particles in the same or neighbouring cells
- Needed array HEAD and array LIST
- Think about parallelisation



Cell decomposition approach

- partition the domain into cells and build a cell-linked list
- Verlet list by searching only in the neighbouring cells instead of considering all particles
- Dirty flag ?

POTENTIAL MODULE

description how the particles will interact in the simulation; two main approximations: Born–Oppenheimer approximation (dynamics of electrons), point particles that follow classical Newtonian dynamics (nuclei, which are much heavier than electrons)

Coulomb-Potential (-> Particle-Mesh)

$$\phi_i(\mathbf{r}) = \frac{1}{4\pi\epsilon_0} \frac{q_i}{|\mathbf{r} - \mathbf{r}_i|}$$
$$\frac{1}{4\pi\epsilon_0} = c_0^2 \mu_0 = c_0^2 \times 10^{-7} \text{ H}\cdot\text{m}^{-1} = 8.987 \times 10^9 \text{ Nm}^2\text{C}^{-2}$$

Lennard-Jones

Particle-mesh Ewald

INTEGRATOR MODULE

Präsentation

