

LAB: Point defects Molecular dynamics simulations

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Logging in to the server



File transfer from your computer:

Use e.g. FileZilla or WinSCP (both free)

Make sure to edit input files directly **on the server**, not on your windows machine.

Output: use as you like!

- The server is running ubuntu
- ssh to 130.237.70.27
- Username: sh2605
- Password: SH2605
 - Do not spread IP, username or password (do not email about them, SMS, Whatsapp, whatever, please...)
- Login and enter the folder MD
 - cd MD
- Then enter the folder with your name
 - cd Yourname (e.g. cd Par for me)
- There you have all the initial files, the executables etc
 - To see files: Is (or Is -I)
 - To edit files: nano file

Elastic scattering – point defects

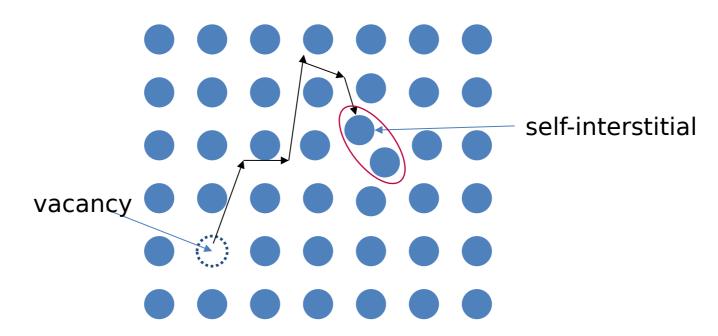
A fast neutron interacts with matter and displaces an atom from its equilibrium lattice site:



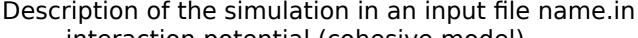
What is the maximum kinetic energy of an Fe atom hit by a 1 MeV neutron?

$$(E_{in} = E_{out}; P_{in} = P_{out})$$

If the atom is given a large enough impulse, it will leave its place in the lattice and create a Frenkel pair, i.e. a vacancy and a self-interstitial.



LAMMPS



- interaction potential (cohesive model)
- crystal
- definition of output files
- simulation (minimization (e.g. quench), molecular dynamics, ...)
- analysis of results



- lammps < name.inp
- lammps < name.in > name.out

To get the wanted information:

- grep "flag" name.out
- or check the other output files!

Example of a simple quench algorithm: if scalar product of velocity and displacement < 0 then set velocity = 0



a) Lattice parameter



- 1. Read an EAM potential for Fe
- 2. Construct a crystal
- 3. Calculate the energy as a function of the lattice parameter
- What is the equilibrium lattice parameter of Fe?
- What is the cohesive energy of Fe?

You may need to change the interval and stepping to "zoom in" on the equilibrium

Extra: Use the input.rose.in to plot the Rose curve → more complete equation of state

b) Vacancy



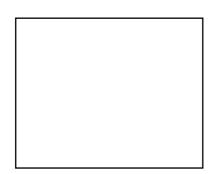
Create a vacancy by removing one atom

- What is the formation energy of a vacancy?
 - Unrelaxed?
 - Relaxed?
 - What is the effect of relaxation on $E_{formation}$?

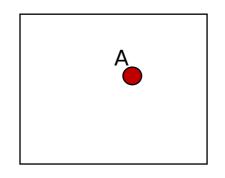
What is the vacancy concentration at 300K, 600K, 900K, 1200K? (from theory, with this E_f)

Formation energy





- N atoms
- E_{ref}



Two cases:

- 1. Relaxation
- 2. Static calculation

- Vacancy: (N 1) atoms: E_{vac}
- Self-interstitial: (N+1) atoms: E_{int}

$$E_{for}(vac) = E_{vac} - E_{ref}(N - 1)/N$$

$$E_{for}(int) = E_{int} - E_{ref}(N+1)/N$$

c) Di-vacancy

Create a vacancy by removing one atom, then a second one in first nearest neighbour position

- What is the formation energy of the di-vacancy?
- What is the binding energy of the di-vacancy?

Change the input in order to setup 2nd, 3rd, etc nearest neighbour di-vacancies

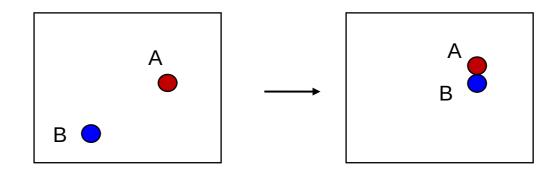
- What is the binding energy of the 2nn di-vacancy?
- What is the binding energy of the 3nn di-vacancy?
- How far do you have to separate the vacancies to have zero interaction? (4nn, 5nn, ...)



Binding energy

Definition:





 $E_{binding} = E(defects A and B far separated) - E(defects A and B close)$

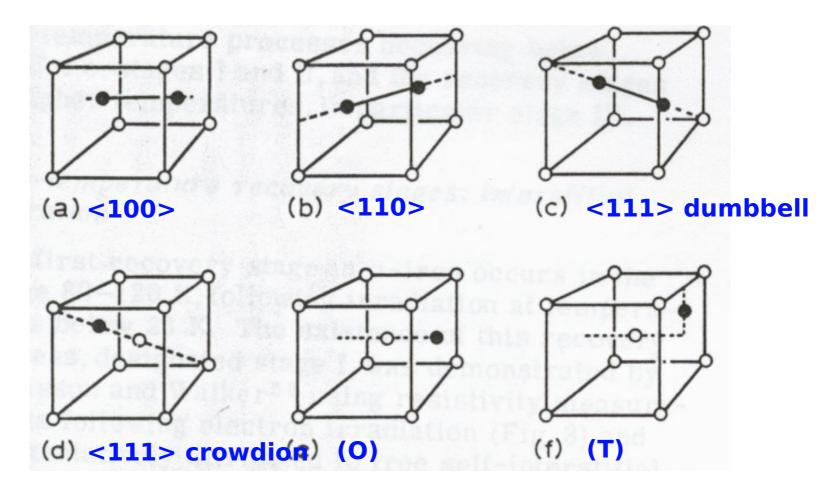
$$E_{binding} > 0$$
: attraction; < 0 : repulsion

Alternate way of calculating binding energy:

d) Self-insterstitial defect configurations

Example: self-interstitial configurations in bcc Fe





d) Self-interstitials

Create a (110) self-nterstitial



- What is the formation energy?
- What is the effect of relaxation? (geometrically)

Copy the input to a new file Modify it in order to create (111) and (100) interstitials

- What are the formation energies?
- Which configuration is the most stable?

Compare to the formation energy of a vacancy

 What is the self-interstitial concentration at 300K, 600K, 900K, 1200K? (from analytical theory, with these values)

e) Threshold energy



Estimate the threshold energy for a stable Frenkel pair in Fe.

 How do the different crystal directions differ? (Investigate (100), (110), (111), (135))

Why should one take care <u>not</u> to use a cubic simulation cell for the (111) direction?