## NeNa

### Untertitel

## Projektarbeit

im Fachbereich Physik



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# Zusammenfassung

Zugsamenfassung: Ziel, Methoden, Ergebnisse, Schlussfolgerungen

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## **List of Abbreviations**

**LAMMPS** Large-scale Atomic/Molecular Massively Parallel Simulator

**HF** Hartree-Fock

**BSSE** Basis Set Superposition Error

fcc face-centered-cubic

# Symbolverzeichnis

Symbol	Bedeutung	Einheit
$\overline{B}$	magnetische Flussdichte	T
D	Elektrische Flussdichte	$\mathrm{Asm^{-2}}$

# 1 Einleitung

Einleitung: Wieso sollte diese Arbeit gelesen werden?

## 2 Aufgabenstellung

We are here trying to calculate optical absorbtion spectra with vasp. The atomic configuration is a cavity inside a neon crystal that was proposed to be filled with either a single sodium atom or two sodiumm atoms (a sodium dimer). Now the first step was to find configurations of cavities, i.e. how many neon atoms will be replaced by an inserted sodium atom and subsequently by a sodium dimer. This substitution number will be denoted as S in the following thesis. For these specific configurations the optical absorbtion spectra will be calculated and compared with the experiment.

### 3 Theory

#### 3.1 Statistical Mechanics

First we consider the system of the crystal of Neon Atoms as a canonical ensemble, this means it will be described under the assumption of an infinitely large heat bath by Boltzmann statistics. The probability distribution of members of the ensemble will then be

$$p(\vec{\mathbf{q}}, \vec{\mathbf{p}}) = \frac{e^{-\beta H(\vec{\mathbf{q}}, \vec{\mathbf{p}})}}{\int \cdots \int e^{-\beta H(\vec{\mathbf{q}}, \vec{\mathbf{p}})} d^{3N} \vec{\mathbf{q}} d^{3N} \vec{\mathbf{p}}},$$
(3.1)

with N being the number of particles.

Now an ensemble is to be understood as many fictional copies of the same system representing the states that a system will explore when propagating in time according to Hamilton's equation of motion. The ensemble distribution represents the relative number of times that an ergodic system will come by a given state after an infinite amount of time has passed. Now for a crystal in principle this still holds. At low temperatures though the phase space that is being explored can be abstracted in such cases. The reason is that the system will for a given finite time usually explore only nearby points in the phase space. This is the crystal configuration but allowing for dynamics such as lattice vibrations and thermal movement of the atoms with respect to their lattice site. So for a given initial point in phase space the system will mostly explore its proximity until it probabilistically jumps to another volume of phase space whose proximity then will be explored for some further time. Now this means the systems lattice configuration has changed which at low but non zero temperature should be reasonable. We now consider these volumes of phase space (being approximately confined regions, i.e. configurations of e.g. a crystalline structure) to be discrete states that the system can be in. So our canonical distribution now is a discrete one, each state representing such a configuration volume. What's left is to figure out what state of the still continuously possible states inside will be used to represent each configuration. We in this work will use the state  $(\vec{\mathbf{q}}_{0i}, \vec{\mathbf{p}}_{0i})$  that locally minimizes  $H(\vec{\mathbf{q}}, \vec{\mathbf{p}})$ . Such a minimum can be found by employing a relaxation calculation in LAMMPS. The Boltzmann distribution now follows to be:

$$p(\vec{\mathbf{q}}_{0i}, \vec{\mathbf{p}}_{0i}) = \frac{e^{-\beta H(\vec{\mathbf{q}}_{0i}, \vec{\mathbf{p}}_{0i})}}{\sum_{n} e^{-\beta H(\vec{\mathbf{q}}_{0n}, \vec{\mathbf{p}}_{0n})}}.$$
(3.2)

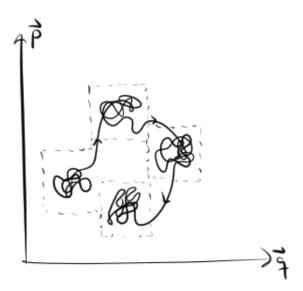


Figure 3.1: state evolution  $(\vec{\mathbf{q}}(t), \vec{\mathbf{p}}(t))$  through 6N-dimensional phase space

grand canonical ensemble with two chemical components sub term of grand canonical ensemble with only ne1 and ne2 Explanation for cohesive energy

### 3.2 Relaxation in LAMMPS

relaxation theory form lammps described in [1] Bitzek, Koskinen, Gahler, Moseler, Gumbsch, Phys Rev Lett, 97, 170201 (2006). Testcite Griffiths[2]

### 3.3 Simulated Annealing

The object that is being annealed is a state vector containing binary information for every lattice site. It refers to the ideal static lattice of the neon structure. A positive value or 1 refers to an atom being present in the initial structure and a negative value or 0 means the neon atom is vacant at this site. So every digit of the binary state refers to a certain specific lattice site. The energy functional of the annealing process will be the energy after minimization (i.e. relaxation) of the above described state vector. This state vector therefore refers to an initial pre-relaxation configuration where sites are simply removed with LAMMPS.

Energy functional contains cohesive Energy/chemical potential, not just hamiltonian!!

Listing 3.1: Simulated annealing algorithm

```
stateVector = [1,1,1,1,1,1,1,1,1,1,1,1,1,1,1]
   alpha = (Tend / Tstart) ** (1 / N)
2
   for i in N:
4
     T = Tstart * (alpha**i)
     n = randomInt(1,len(stateVector))
     stateVector[n] = 1 - stateVector[n]
9
     stateVectorOld = stateVector
10
11
     E0ld = minimizeEnergy(stateVectorOld)
12
     E = minimizeEnergy(stateVector)
13
14
     if E >= E01d:
16
       stateVector = stateVectorOld
     elif random(0,1)<=exp(-1/T*(E-E0ld)):
17
       stateVector = stateVectorNew
18
19
   S = 1
20
   for site in stateVector:
21
   if site == 0:
22
       S += 1
23
```

### 3.4 Symmetry optimization

In the case of a single sodium atom inserted in the neon crystal the highly symmetrical structure was exploited to reduce redundant minimization calls in the annealing routine, by identifying symmetrically equivalent structures and caching them. Applying rotation and reflection matrices of the octahedral symmetry group to a single configuration was used to quickly create all equivalent configurations. By sorting them lexicographically and picking e.g. the first, one ensures every redundant set of configurations is always represented by the same configuration.

### 3.5 DFT optical spectra

#### Listing 3.2: lexicographically sorting point group symmetries

```
equivs = np.transpose(np.einsum("klj,ij", oh_group, pos), (1, 0, 2))

for k in range(equivs.shape[0]):
equivs[k] = equivs[k][np.lexsort(equivs[k].T[::-1])].copy()
sorting_idx = np.lexsort(
np.transpose(equivs, (2, 1, 0)).reshape(pos.size, oh_group.shape[0])

np.transpose(equivs, (2, 1, 0)).reshape(pos.size, oh_group.shape[0])

pos = equivs[sorting_idx[0], :, :]
```

### 4 Finding Minima of Neon Replacements

#### 4.1 Pair Potential Interaction

### 4.1.1 Hartree-Fock (HF) equations

We start with the pair potentials of Ne-Ne, Ne-Na and Na-Na to mimic the interaction of long distance van der Vaals + short distance repulsion forces. The pair potentials have been calculated using a HF self consistent field cycle by varying the atom distance. Now selfconsistently solving the equations:

$$\left[ -\frac{\hbar^2}{2m} \vec{\nabla}^2 - \frac{Ze^2}{4\pi\varepsilon_0} - \frac{e^2}{4\varepsilon_0} \sum_{\nu,\sigma}^{(\mu\sigma)\neq(\nu\sigma)} \iiint_{\mathbb{R}^3} \frac{\varphi_{\nu\sigma'}(\tilde{\vec{r}})}{\|\vec{r} - \tilde{\vec{r}}\|} d^3 \tilde{\vec{r}} - \hat{A}_{\mu\sigma}(\vec{r}) \right] \varphi_{\mu\sigma}(\vec{r}) = \varepsilon_{\mu\sigma} \varphi_{\mu\sigma}(\vec{r}),$$
(4.1)

with  $\hat{A}_{\mu\sigma}$  being the exchange correlation term leads to the eigenenergies  $\varepsilon_{\mu\sigma}$ . Since HF minimizes a slater determinant, the energies correspond to the factors of the slater product state, meaning the systems energy is the sum of the energy of the single electrons. From that, to get the actual energy of the whole system, one needs to remove inter-electron repulsion and exchange correlation from the sum of all eigenenergies  $\varepsilon_{\mu\sigma}$ :

$$E_{HF}^{0} = \sum_{\mu,\sigma} \varepsilon_{\mu\sigma} - \frac{1}{2} \sum_{\substack{\mu,\nu\\\sigma,\sigma'}}^{(\mu\sigma\neq\nu\sigma)} \left[ C_{\mu\sigma}^{\nu\sigma'} - A_{\mu\sigma}^{\nu\sigma} \delta_{\sigma\sigma'} \right]$$
(4.2)

with  $C^{\nu\sigma'}_{\mu\sigma}, A^{\nu\sigma}_{\mu\sigma}$  being defined as:

$$C_{\mu\sigma}^{v\sigma'} = \frac{e^2}{4\pi\varepsilon_0} \iint \frac{|\varphi_{\mu\sigma}(r)|^2 |\varphi_{v\sigma'}(r')|^2}{|r - r'|} d^3r d^3r', \tag{4.3}$$

$$A_{\mu\sigma}^{v\sigma} = \frac{e^2}{4\pi\varepsilon_0} \iint \frac{\varphi_{\mu\sigma}^*(r)\varphi_{v\sigma}^*(r')\varphi_{\mu\sigma}(r')\varphi_{v\sigma}(r)}{|r - r'|} d^3r d^3r'.$$
 (4.4)

Since solving the HF equations in real space is too costly one restricts the variations to coefficients of a superposition of fixed basis vectors. These lead to the Roothaan-Hall equations, which are a discrete Matrix representation of the HF equations:

$$FC = SC\epsilon, \tag{4.5}$$

with  $\epsilon$  being a diagonal matrix of single particle energies on the diagonal, **S** the basis overlap matrix, **C** the coefficient matrix and finally **F** the Fock operator.

#### 4.1.2 Moeller-Plesset Perturbation

-go from pair HF to moller plesset

#### 4.1.3 Basis Set Superposition Error (BSSE)

Now the the superpositions of a fixed basis leads to the BSSE. These errors are being corrected for by using 'ghost calculations', essentially the same calculation with one atom being removed (i.e. the extended two atom centered basis stays the same). This means the basis will have access to the second electron states, but the nucleus and additional electrons are not accounted for. These 'ghost energies' will then subsequently be removed from the HF approximation to correct for the BSSE. This is know as the counterpoise-corrected interaction energy formula or Boys–Bernardi method:

$$E_{\text{int}}^{\text{CP}}(R) = E_{AB}(R) - E_A^{\text{ghost}}(R) - E_B^{\text{ghost}}(R). \tag{4.6}$$

#### 4.1.4 Interpolation

- -explain exact interpolation
- -paper citation where brute force plot shows up as well
- -which basisi is being used in the roothaan equations
- -explain setup of calculation with ball being carved out
- -maybe here explain the nearest neighbours

lammps gets linear interpolation

-since plot is in priniple linearly interpolated this is the exact pair interactins that lammps will read

### 4.2 Geometric LAMMPS Setup

The simulation models a finite spherical cluster of atoms extracted from a bulk facecentered cubic lattice. To balance computational efficiency with physical realism, the system is partitioned into two concentric spherical regions defined in units of the lattice constant. The inner spherical region, with radius  $n_{inner}$  contains atoms that are free to move and evolve dynamically. This region represents the core of the cluster where

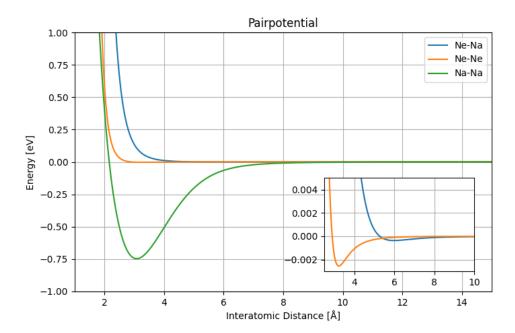


Figure 4.1: Plots of interpolations of pair potentials of Ne-Ne, Na-Na, Ne-Na calculated with Moeller Plesset 2 perturbation theory.

physical process and in this case the energy minimization via a relaxation calculation of LAMMPS takes place.

Sourrounding this core is an immobilized shell with thickness  $n_{outer}$  Atoms within this outer shell are constrained by setting their forces to zero, effectively fixing them in space. This fixed shell act as a rigid boundary that suppresses surface effects and mimics the presence of a an extended bulk lattice, thereby reducing finite size artifact in the dynamics of the inner region.

Consequently the total cluster is confined within a spherical domain of radius  $R_{tot} = (n_{inner} + n_{outer}) \cdot a$  with a being the lattice constant of the pure neon face-centered-cubic (fcc) lattice.

Technically the simulation is defined as a cubic volume large enough to contain the entire spherical cluster, with half-length at least  $R_{tot}$ . For the relaxation mechanics atomic positions are set according to 4.2 on a perfect fcc lattice no matter their role (sodium defect, fixed neon, dynamic neon) as long as they are inside the cutoff radius  $R_{tot}$ . This approach is commonly employed in molecular dynamics studies to simulate nanoparticles or finite clusters embedded in bulk-like surroundings.

The centered red dots in 4.2 represent sodium atoms. They represent the initial prerelaxation positions of sodium inside the dynamic inner sphere. These initial positions

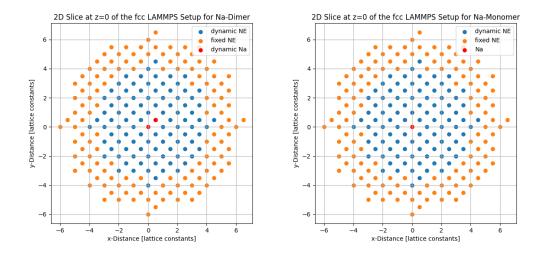


Figure 4.2: Initial setup for the LAMMPS relaxation. Only the inner blue sphere is allowed to relax, forces on the orange atom positions are overridden to 0.

are the same for every relaxation step of the simulated annealing. Note that the minimum energy of the pair-potential in 4.1 is roughly at 3.1Å which is the nearest neighbor distance in the neon fcc lattice. With a = 4.4637Å:

$$\vec{a}_1 = \begin{bmatrix} a \\ 0 \end{bmatrix} \qquad \vec{a}_2 = \begin{bmatrix} 0 \\ a \end{bmatrix} \tag{4.7}$$

$$\Rightarrow \|\frac{1}{2}(\vec{a}_1 + \vec{a}_2)\| = \frac{1}{\sqrt{2}} \cdot a = 3.1563\text{Å}.$$
 (4.8)

We therefore put the second sodium atom in the dimer calculations at the nearest neighbor lattice site.

#### 4.3 Sodium Monomer

Now the annealing was done as described in the chapters before for a single sodium atom at the center. Since the known energetically minimal configurations for every S and it's corresponding energy was known due to a brute force calculation, this calculation serves as a benchmark for the annealing algorithm. The brute force minimal energies for the vacancies are shown in 4.3 on the right axis. The left axis shows the relative counts (with respec to the total counts) of one annealing sweep consisting of itself 1000 random removals of neon atoms from the state vector. This annealed approach quickly converges to a statistical distribution shown in the same figure 4.3 on the left y-axis. The pattern surfaces quickly after about 50 sweeps, the total amount of sweeps for the plot were

736. The true local minima at S=10 and S=13 could be picked up upon quite sharply. At S=8 the annealing algorithm seems to pick up on the notch, that is not quite an actual minima but could look like a local minima if approached from many directions in the high dimensional phase space.

- -Not feasible to brute force dimer with 3rd nearest neighbors
- -explain nearest neighbors
- -even with full 48 cubic point symmetry of the host of the defect which no structure can exceed ...
- -explain confirmation of algorithm with brute force for mono sodium
- -explain how plot is created with sweeps
- -more complicated symmetries
- -compare figures
- -calculation for dimer
- -discuss noteworthy structure (e.g. inner shell carved out)
- -citation of paper that has the same plot Now first by brute force search we know the minima for a single sodium atom.

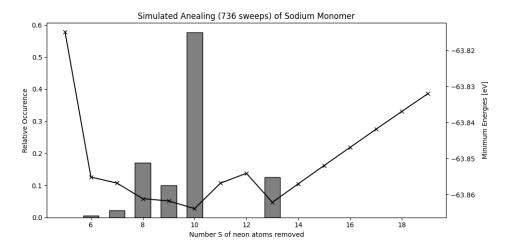


Figure 4.3: Simulated Annealing for single sodium atom inserted, compared to a brute force minima search for a fixed number of removed atoms S.

Since 4.3 shows the simulated annealing algorithm accurately picks up on the location of the minima (albeit with no quantitative measure and misleading local minima being blown out of proportion (See S=8)) we try the same approach for the sodium dimer. Here the energetically optimal S replacement will exceed the number of available sites as in the monomer calculation (i.e. up to second nearest neighbor). The sweeps and their random removal of neon atoms now act upon an extended state vector containing every lattice site up to the 3rd nearest neighbor. This is also the very reason a brute force

approach is unfeasable, since the volume of phasespace to be explored grows quickly. This is the reason the statistical approach was chosen in the first place.

-brute force lexiciraphically sorting algorithm here

### 4.4 Sodium Dimer

- -in setup exxplain that dimer binding position is roughly first nearest neighbour of ne lattice
- -3rd nearest neighbors, why? because minima is more the 2 shells -rough estimate of time used by brute forcing
- -even with full cubic symmetry this is too long (divide b 64)
- -maybe think about symmetry
- -same graphics as for the monomer case
- -heuristically take the maxima of these plots
- -give structures
- -discuss structures (inner shell carved out)

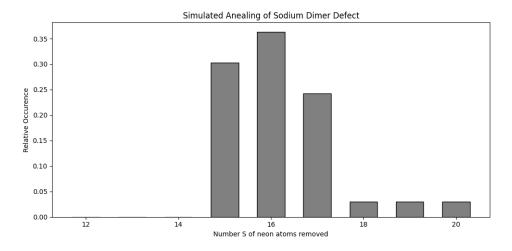


Figure 4.4: Simulated annealing results and their relative occurrence after x sweeps

# 5 DFT optical spectra results

Zweites Kapitel

# 6 Conclusion

Unfortunately no confidence or error can be estimated since the approach was purley heuristcal. Optical spectrum does not depend much on replacements and precise structure.

# 7 Ausblick

Ausblick

# **Bibliography**

- [1] P. Bitzek et al. "Structural Relaxation Made Simple". In: *Physical Review Letters* 97.17 (2006), p. 170201. DOI: 10.1103/PhysRevLett.97.170201.
- [2] David J. Griffiths. Introduction to Quantum Mechanics. 3rd ed. Cambridge University Press, 2018. ISBN: 9781107189638.

# A Anhang

# A.1 Inhalt des beigefügten Datenträgers



## Eidesstattliche Erklärung

Ich, Gabriel Sommer, versichere hiermit, dass ich die vorliegende Arbeit selbstständig verfasst und keine anderen als die angegebenen Quellen und Hilfsmittel benutzt habe, wobei ich alle wörtlichen und sinngemäßen Zitate als solche gekennzeichnet habe.

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## **Declaration of Authorship**

I, Gabriel Sommer, hereby declare that I have written the present thesis independently and have used no sources or aids other than those explicitly stated. All direct quotations and all paraphrased ideas have been clearly marked as such.

This work has not been submitted, either in the same or a substantially similar form, to any other examination board, nor has it been published elsewhere.

Vienna, August 10, 2025	
Garriel Sommer	