

CAMD Summer School 2024

Electronic structure tools: ASE and GPAW and introduction to the computer projects

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Overview

- The Python language
- The Atomic Simulation Environment (ASE)
 - The anatomy of an atomic-scale simulation/calculation
 - -The ASE
- Examples
 - -Almost the simplest possible molecular dynamics simulation.
 - -Almost the simplest possible GPAW calculation.
- Using the DTU "databar" (computer lab).
- Computational projects





Part I

BRIEF INTRODUCTION TO PYTHON

Why Python?

- For the programmer: Python is object-oriented
 - Object oriented and modular: Facilitates writing and maintaining complex problems.
 - Dynamically typed: Flexibility, facilitates code reuse.
 - Easy to write readable code: Code is maintainable.
 - Large libraries available (numerics, plotting, ...)
- For the user: Python is a scripting language.
 - Great for scripting a calculation
 - Great for small programs and prototypes.
 - Great for interactive experimenting.
 - Easy to learn.
 - Objects are powerful in scripts!
- Python can be extended in C/C++/Fortran
 - Solves performance problems of non-compiled languages.

Learning Python

- Don't waste money on Python books!
 - -It's not that hard, and online docs are good.
- Python tutorial: <u>http://docs.python.org/3/tutorial/</u> –More documentation at docs.python.org
- Learn Python + ASE + GPAW by example
 - -Get a simple script, and modify it.
 - -Simple scripts are almost like old-fashioned input files!

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A few places where Python is different

See notebook!





Part II

INTRODUCTION TO ASE AND GPAW

The "anatomy" of an atomic-scale computer simulation









import numpy as np
from ase import Atoms, units
from ase.io.trajectory import Trajectory
from ase.build import bulk
from ase.md.verlet import VelocityVerlet
from asap3 import EMT

Example: Molecular dynamics

```
# Create the atoms
atoms = bulk("Cu", "fcc", cubic=1).repeat([3,3,3])
atoms.set_pbc(False)
atoms.center(vacuum=5.0)
# Give the first atom a non-zero momentum
atoms[0].momentum = np.array([0, -11.3, 0])
```

```
# Prepare to do molecular dynamics, forces described by EMT
atoms.calc = EMT()
dyn = VelocityVerlet(atoms, 5.0*units.fs)
```

```
# Make a trajectory writing output every fifth timestep.
trajectory = Trajectory("MD-output.traj", "w", atoms)
dyn.attach(trajectory, interval=5)
```

```
# Now do 1000 timesteps.
dyn.run(1000)
```

GPAW Example: Atomization energy of Hydrogen (1/2)

```
from ase import Atoms, Atom
from gpaw import GPAW
```

```
# gpaw calculator:
calc = GPAW(h=0.18, nbands=1, xc='PBE', txt='H.out')
atom.calc = calc
```

```
e1 = atom.get_potential_energy()
calc.write('H.gpw')
```



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GPAW Example: Atomization energy of Hydrogen (2/2)

Continued ...

```
# Hydrogen molecule:
d = 0.74 # Experimental bond length
molecule = Atoms('H2',
                 positions = ([c - d / 2, c, c])
                           [c + d / 2, c, c]),
                 cell=(a, a, a))
calc.set(txt='H2.out')
molecule.calc = calc
e2 = molecule.get_potential_energy()
calc.write('H2.gpw')
print 'hydrogen atom energy:
                            %5.2f eV' % e1
print 'hydrogen molecule energy: %5.2f eV' % e2
                            %5.2f eV' % (2 * e1 - e2)
print 'atomization energy:
```

The Atoms object

The Atoms object is the main simulation object. It contains:

- Per-atom data: positions, velocities, charges, magnetic moments, tags.
- Global data: Unit cell, boundary conditions.
- Refs to helper objects: Calculator, constraints, ...

Accessing the data:

r = atoms.get_positions() Or r = atoms.positions
atoms.set_positions(r) Or atoms.positions = r

f = atoms.get_forces() Will trigger a calculation, if needed.

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The Atoms object

Manipulating the Atoms object:

atoms.center(vacuum=5.0)	Centers in unit cell, possibly adjusting the amount of vacuum.
atoms.rotate(45, 'z')	Rotate the atoms
<pre>atoms.repeat([2,2,1])</pre>	Replicate atoms along x,y,z axes.

The Atoms object is a Python sequence:

atoms[i]	The i th atom (starting at zero).
atoms += atoms2	Add more atoms
atoms1 + atoms2	Merge two atoms objects
for a in atoms:	Loop over atoms
print(a.position)	

Making systems

Manually specify the atoms:

```
from ase import Atoms
h2 = Atoms('H2', positions=[(0, 0, 0), (0, 0, 0.74)])
```

Known molecules and bulk structures:

```
from ase.build import molecule, bulk
water = molecule('H2O')
si2 = bulk('Si', 'diamond', a=5.4)
```





Simple surfaces:

```
from ase.build import fcc110
slab = fcc110('Pt', (2, 1, 7), a=4.0, vacuum=6.0)
```



GPAW essentials

• Grid mode – wavefunctions on a real-space grid

```
calc = GPAW (h=0.18, xc='PBE', ...)
```

Parallelizes very well for large systems. High accuracy

• Plane wave mode – wavefunctions in k-space

calc = GPAW (mode=PW(400), xc=..., ...)

Faster than grid mode for small/medium systems. High accuracy

• LCAO mode - wavefunctions in real space; with basis fct.

```
calc = GPAW(mode='lcao', basis='dzp', h=0.18, ...)
```

Fast but less accurate.



Part III

COMPUTER EXERCISES AND THE DTU "DATABAR"

Increased IT security at DTU

- All external log in to DTU systems require two-factor login
- In our case the two factors are
 - -Your password
 - -A cryptographic key on your laptop



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The "databar" and the projects

 Information about the Summerschool projects: wiki.fysik.dtu.dk/gpaw/

Includes detailed instructions on how to log in.

- Changing password in the HPC system ("databar"):
 - -Change at password.dtu.dk (takes up to 1 hour to sync)
 - Download your SSH Secret key with original password!
- **SLIDES** available from the GPAW summerschool pages.

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Projects

- Excited states
 - -Jakob Svaneborg and Jiban Kangsabanik
- Catalysis
 - -Georg Kastlunger and Dipam Patel
- Magnetism
 - -Martin Ovesen and Varun Rajeev Pavizhakumari
- Batteries
 - -William Hansen and Lotte Kortstee
- Machine Learning
 - Jesper Rask Pedersen and Armando Morin Martinez
- Computational workflows
 - -Ask Hjort Larsen and Tara Boland

Project: Excited states

- Calculate electronic band structures and bandgaps
- Calculate optical absorption

- Special methods:
 - -The GW approximation
 - -The Random Phase approximation
 - -Optional: The Bethe-Salpeter Equation

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Project: Catalysis

- DFT calculations of adsorption geometries of molecules on surfaces.
- Calculations of the reaction path and the transition energy
- Special methods:
 - -Nudged Elastic Band (for transition paths)
 - -Optional: Vibrational analysis for adsorption entropy

Project: Magnetism in 2D

- Calculations of the critical temperature of a Crl₃ monolayer
- Calculations of the noncollinear ground state in VI₂
- Autodiscovery of new magnetic monolayers in the Computational 2D Materials Database (C2DB)
- Special methods:
 - -Energy mapping analysis
 - -Noncollinear DFT calculations

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- Study the anode and cathode materials of a Li-ion battery with DFT
- Calculate the intercalation energy of Li in graphite, establish the equilibrium potential of a LiFePO4/C battery and determine important battery characteristics such as Li transport barriers and the voltage profile.
- Special methods:
 - Structure creation and modification with ASE
 - Unit cell relaxation
 - Bayesian error estimation
 - Nudged Elastic Band (NEB) calculations for estimating Li migration barriers

Project: Machine Learning

- Machine Learning methods used on an example database
- Predicting band gaps and heat of formation using a structure "fingerprint"
- Predictions with several methods (e.g., ridge regression, decision tree, and gaussian process)
- Validation of predictions using DFT

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Project: Computational Workflows

- Learn to define tasks and workflows with TaskBlaster
- Use command-line tools to create and manage tasks in a directory tree
- Run a materials workflow on multiple materials

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