

Modern Tools for Spectroscopy on Advanced Materials

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Jens Jørgen Mortensen

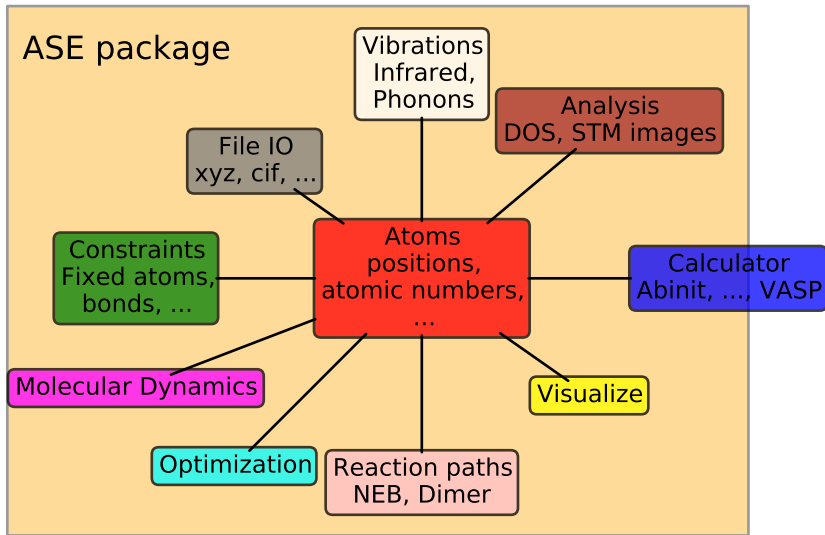
Department of Physics

Technical University of Denmark

- Overview of ASE
- The Python language
- The `Atoms` object
- Building stuff (molecules, surfaces, ...)
- Doing stuff with atoms (MD, optimization, ...)
- Databases
- Installing ASE
- Development of ASE

What is ASE?

ASE is a Python library for working with atoms:



<http://wiki.fysik.dtu.dk/ase>

A recipe for your atoms

```
from ase import Atoms
from ase.optimize import BFGS
from ase.calculators.nwchem import NWChem
from ase.io import write
h2 = Atoms('H2',
           positions=[(0, 0, 0),
                      (0, 0, 0.7)])
h2.calc = NWChem(xc='PBE')
opt = BFGS(h2)
opt.run(fmax=0.02)
write('H2.xyz', h2)
```

Units: eV and Å.

An interactive recipe for your atoms

```
>>> from ase import Atoms
>>> from ase.optimize import BFGS
>>> from ase.calculators.nwchem import NWChem
>>> from ase.io import write
>>> h2 = Atoms('H2',
               positions=[(0, 0, 0),
                           (0, 0, 0.7)])
>>> h2.calc = NWChem(xc='PBE')
>>> opt = BFGS(h2)
>>> opt.run(fmax=0.02)
BFGS:   0  19:10:49   -31.435229   2.2691
BFGS:   1  19:10:50   -31.490773   0.3740
BFGS:   2  19:10:50   -31.492791   0.0630
BFGS:   3  19:10:51   -31.492848   0.0023
>>> write('H2.xyz', h2)
>>> h2.get_potential_energy()
-31.492847800329216
```

What is Python?

`http://www.python.org`

- Interpreted language
- Dynamically typed
- Easy to read and write, and very expressive
- Great for scripting a calculation
- Great for small and large programs
- Optimized for programmer productivity
- Can be extended in C/C++/Fortran
- General purpose language

Numpy: Base N-dimensional array package

Scipy: Fundamental library for scientific computing

Matplotlib: Comprehensive 2D Plotting

Sympy: Symbolic mathematics.

- Python modules and sub-modules:

constraint, data, db, dft, dimer, ga, infrared, io,
md, neb, optimize, phonons, structure,
thermochemistry, transport, units, vibrations
and visualize.

- Command-line tools:

- ase-gui: Graphical user interface

- ase-db: Manipulate ASE-database

- ase-build: Build molecules or crystals

- ase-run: Run simple calculation

The Atoms object



What's inside?

- Positions, atomic numbers, velocities, masses, initial magnetic moments, charges, tags
- Unit cell
- Boundary conditions
- (Calculator)
- (Constraints)

Methods and attributes of the Atoms object

List methods:

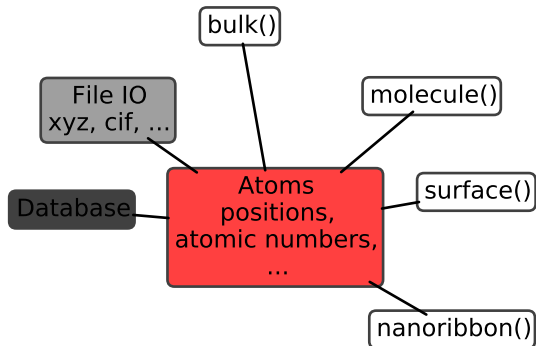
```
a.pop() # remove last atom
a.append('Xe') # append xenon atom
a.extend(Atoms('H2')) # append 2 hydrogens
a[1] # second atom
del a[-3:] # delete three last atoms
```

Other methods: repeat(), translate(), center(), rotate(), get_distance(), get_angle(), get_moments_of_inertia(), ...

Special attributes: positions, numbers, cell, pbc (Numpy n-d arrays)

```
x1, x2 = a.positions[-2:, 0] # x-coords of
                             # last two atoms
a.positions[-2:, 0] += 0.1 # translate atoms
a.positions[:, 2].max() # maximum z-coord
```


Building the Atoms object



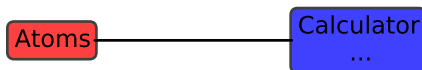
Building the Atoms object ...

```
from ase import Atoms
h2 = Atoms('H2', [(0, 0, 0), (0, 0, 0.74)])
from ase.structure import molecule
water = molecule('H2O')
from ase.lattice import bulk
si2 = bulk('Si')
from ase.lattice.spacegroup import crystal
rutile = crystal(['Ti', 'O'],
                 basis=[(0, 0, 0), (u, u, 0)],
                 spacegroup=136,
                 cellpar=[a, a, c,
                          90, 90, 90])
from ase.lattice.surface import fcc110
slab = fcc110('Pt', (2, 1, 7), a=4.0,
              vacuum=6.0)
from ase.lattice.surface import add_adsorbate
add_adsorbate(slab, 'H', site='hollow')
```

Building the Atoms object, continued ...

```
from ase.lattice.surface import surface
nasty_cut = surface(rutile, (1, 1, 0),
                   layers=5)

from ase.io import read
q = read('quarts.cif')
import ase.db
con = ase.db.con('mystuff.db')
atoms = con.get_atoms(foo='bar', H=0)
# same as this:
atoms = read('mystuff.db@foo=bar,H=0')
```



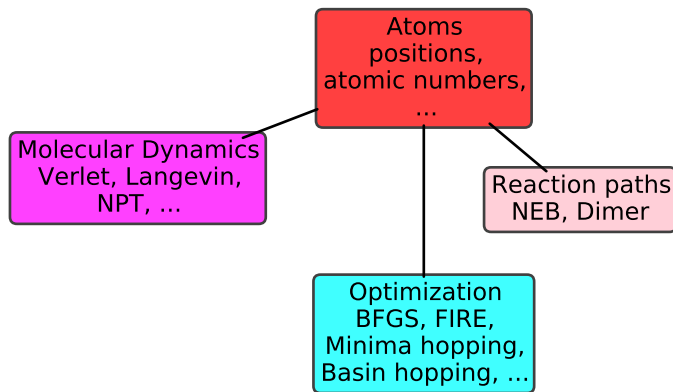
DFT, *ab initio*:

Abinit, Castep, ELK, Exciting, FHI-Aims, Fleur, Gaussian, GPAW, Jacapo, JDFTx, NWChem, QuantumEspresso, SIESTA, Turbomole, VASP

(Semi-)empirical potentials:

ASAP, DFTB+, EAM, EMT, Gromacs, Hotbit, LAMMPS, Lennard-Jones, MOPAC, Morse

Optimization, MD and minimum energy paths



In each row we have:

Taxonomy:

- Atoms object (positions, atomic numbers, ...)
- ID, user-name, creation and modified time
- Constraints
- Calculator name and parameters
- Energy, Forces, Stress tensor, dipole moment, magnetic moments

Folksonomy:

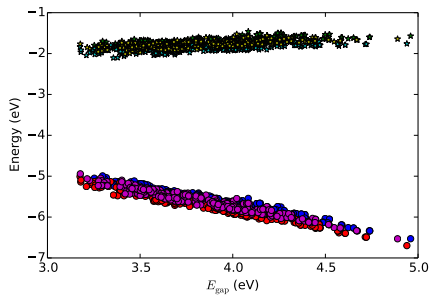
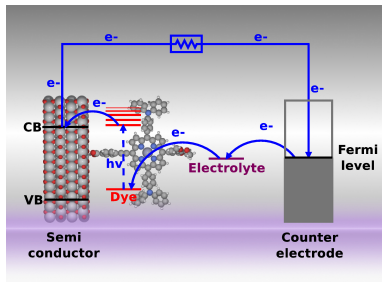
- Key-value pairs
- Keywords

Additional stuff:

- extra data (band structure, ...)

Back-ends: JSON, SQLite3 and PostgreSQL.

Dye sensitized solar cells



¹Kristian B. Ørnsø, Christian S. Pedersen, Juan M. Garca-Lastra and Kristian S. Thygesen *Optimizing porphyrins for dye sensitized solar cells using large-scale ab initio calculations* Phys. Chem. Chem. Phys., 2014, 16, 16246-16254

A figure from Ørnsø *et al.* (almost)

```
import matplotlib.pyplot as plt
import ase.db
con = ase.db.connect('dssc.db')
for M in ['Zn', 'TiO', 'H2']:
    gap = []
    homo = []
    lumo = []
    for dct in con.select(metal=M,
                          anchor='EthynPhA'):
        gap.append(dct.E_gap)
        homo.append(dct.E_HOMO)
        lumo.append(dct.E_LUMO)
    plt.plot(gap, homo, 'o')
    plt.plot(gap, lumo, '*')
plt.ylabel(r'Energy (eV)')
plt.xlabel(r'$E_{\mathrm{gap}}$ (eV)')
plt.savefig('homolumo.svg')
```


Three types of calculators

- 1)
 - a) ASE writes input file
 - b) ASE starts Fortran/C/C++ code (maybe in parallel)
 - c) ASE waits for Fortran/C/C++ code to finish
 - d) ASE reads output file
- 2) Same as above, except that the Fortran code doesn't stop - it waits for a new input file with new coordinates (QuantumEspresso, Dacapo).
- 3) Python solution (EMT, EAM, ...). For parallel calculations, the Python interpreter runs on all processes (GPAW, ASAP)

Installing ASE

Requirements: Python and NumPy (ase-gui: PyGTK and Matplotlib)

```
$ SVN=https://svn.fysik.dtu.dk/projects  
$ svn co $SVN/ase/trunk ase  
$ cd ase  
$ python setup.py install --user  
$ python setup.py test # takes 1 min.
```

Where is what?

ase/:

Source code.

tools/:

Command-line tools

ase/doc/:

Documentation (source for web-page)

How to get started?

- <http://wiki.fysik.dtu.dk/ase>
- <http://docs.scipy.org/doc/numpy>
- <http://www.python.org>

How to get help?

Write to the ase-users mailing list.

How is ASE developed?

Mostly by volunteers. Coordination and discussions on the ase-developers mailing list.

How to contribute?

Please send us bug-reports, patches, code and ideas.

License?

LGPLv2.1+

Thanks

Andrew Peterson, Anthony Goodrow, Ask Hjorth Larsen, Carsten Rostgaard, Christian Glinsvad, David Landis, Elvar Örn Jónsson, Eric Hermes, Felix Hanke, Gaël Donval, George Tritsarlis, Glen Jenness, Heine Anton Hansen, Ivano Eligio Castelli, Jakob Blomquist, Jakob Schiøtz, Janne Blomqvist, Janosch Michael Rauba, Jens Jørgen Mortensen, Jesper Friis, Jesper Kleis, Jess Wellendorff Pedersen, Jingzhe Chen, John Kitchin, Jon Bergmann Maronsson, Jonas Bjork, Jussi Enkovaara, Karsten Wedel Jacobsen, Kristen Kaasbjerg, Kristian Baruel Ørnsø, Lars Grabow, Lars Pastewka, Lasse Vilhelmsen, Marcin Dulak, Marco Vanin, Markus Kaukonen, Mattias Slabanja, Michael Walter, Mikkel Strange, Poul Georg Moses, Rolf Würdemann, Steen Lysgaard, Stephan Schenk, Tao Jiang, Thomas Olsen, Tristan Maxson, Troels Kofoed Jacobsen, ...

How to use `sys.argv`, other ASE functions, for-loops and string interpolation:

```
import sys
from ase.lattice.surface import fcc111
from gpaw import GPAW, PW
a = float(sys.argv[1])
k = int(sys.argv[2])
for n in range(1, 5):
    slab = fcc111('Cu', size=(1, 1, n),
                  a=a, vacuum=5.0)
    slab.calc = GPAW(mode=PW(ecut=400),
                     kpts=(k, k, 1),
                     txt='Cu%d.txt' % n)
    slab.get_potential_energy()
```