

# Python interface for MyMD

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# Design Problems

- MyMD code to be completely reorganized
- No fixed API
- No compiling code to test against
- => Decision: 'API' = input file format

# Getting started

- $4 * 1/2$  day = not a lot
- usable interface = lot of work
- $\Rightarrow$  What's already there?

# Atomic Simulation Environment ¶

The Atomic Simulation Environment (ASE) is the common part of the simulation tools developed at CAMd. ASE provides [Python](#) modules for manipulating atoms, analyzing simulations, visualization etc.

**Note:** The old ASE-2 webpage has moved to <http://wiki.fysik.dtu.dk/ase2>.

Supported [calculators](#):



# Calculator Class

```
"""This package defines an ASE interface to MyMD
```

```
MyMD is an educational project, written during the 2013  
Workshop on Computer Programming and Advanced Tools for Scientific Research Work  
at ICTP Trieste, Italy.  
"""
```

```
import os  
import sys  
import subprocess
```

```
import numpy as np
```

```
from ase.atoms import Atoms  
from ase.calculators.calculator import FileIOCalculator, Parameters, ReadError
```

```
import potentials  
import io
```

```
class FileIOMyMD(FileIOCalculator):
```



# Example

```
""" Example 03
```

```
This example shows how to use the built-in atom viewer  
of ASE as well as data visualization with matplotlib.  
"""
```

```
from ase.lattice.cubic import SimpleCubic  
from ase.md.velocitydistribution import MaxwellBoltzmannDistribution  
import ase.units as units  
import ase.calculators.mymd as mymd  
  
# Set up a cube of 125 Argon atoms in a cube of (15 Angstroms)**3.  
myatoms = SimpleCubic('Ar', latticeconstant=3.0, size=(5,5,5))  
scalef = 8.31446714569e-7      # adjust velocity units  
MaxwellBoltzmannDistribution(myatoms, temp=50*scalef, force_temp=True)
```

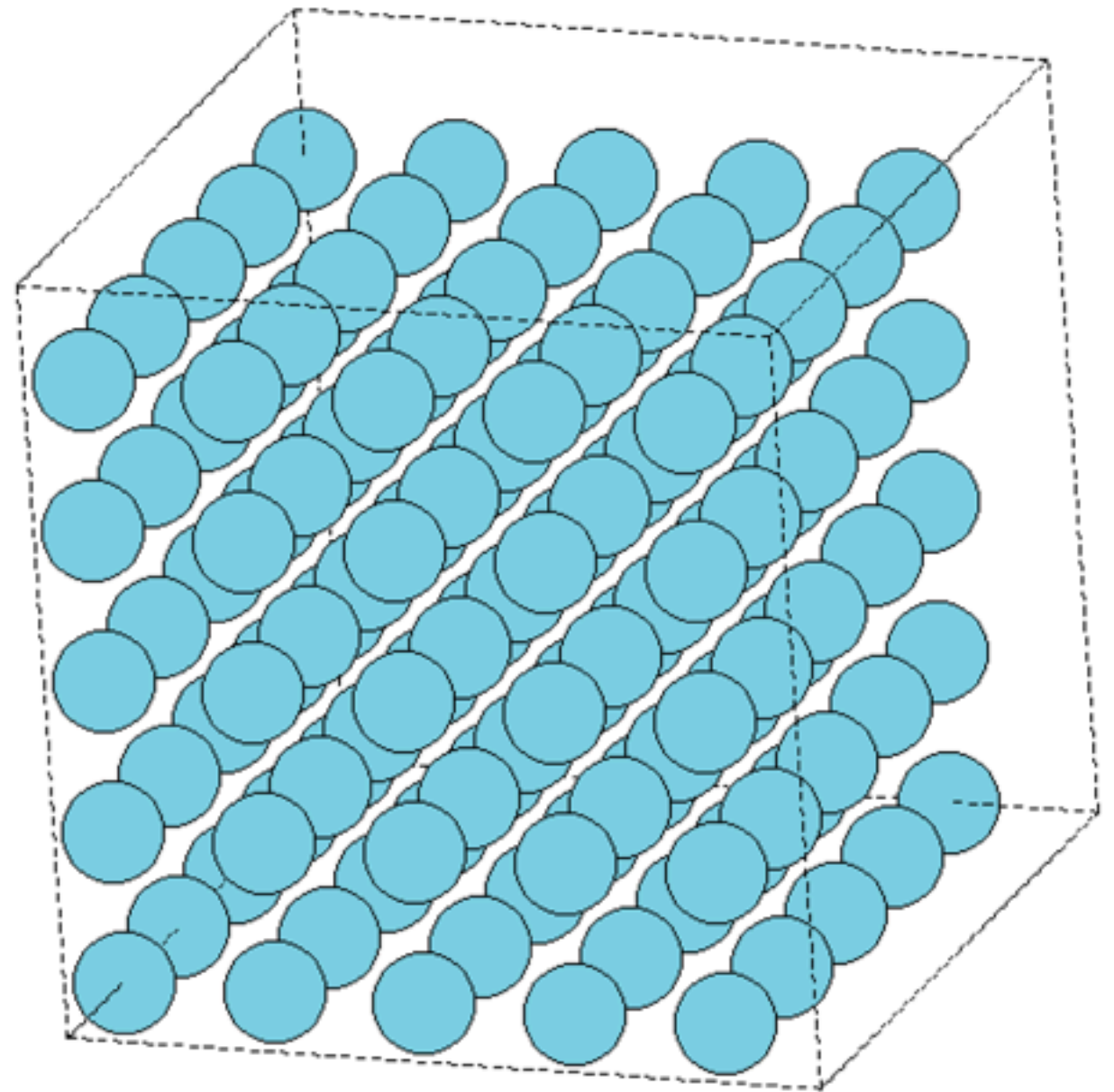
# Example

```
from ase.visualize import view
view(myatoms)

# Set up MyMD calculator and run
calc = mymd.FileIOMyMD(label='mymd',
                        nsteps=10000,
                        dt=2.0,
                        nprint=100)
calc.run_md(myatoms)

# View final state of the atoms
view(calc.state)

# View trajectory of the atoms
trajectory = calc.frames.collect('atoms')
view(trajectory)
```





# Example

```
# Visualize data using matplotlib
import matplotlib.pyplot as plt

n = calc.frames.collect('index')
ekin = calc.frames.collect('ekin')
epot = calc.frames.collect('epot')
etot = calc.frames.collect('etot')

plt.plot(n, epot, label='Potential Energy')
plt.plot(n, ekin, label='Kinetic Energy [kcal/mol]')
plt.plot(n, etot, label='Total Energy [kcal/mol]')
plt.legend()

plt.show()
```

