

Molecular Simulation HOMEWORK 2: Properties of nitromethane

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9/18/12 Thursday

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1 Molecular Weight

Use ase and python to compute the molecular weight of nitromethane (CH3NO2). Compare your answer to what you compute by hand

```
from ase.data.molecules import molecule
import numpy as np

# Calculate using ase and numpy functions
atoms = molecule('CH3NO2')
masses = atoms.get_masses()
total = masses.sum()
```

```
8  print 'ASE and numpy calculates the molecular weight of nitromethane is {0:1.3f} g/mol'.format(total)
9
10  # Calculate 'by hand' using periodic table
11  C = 12.011
12  H = 1.008
13  N = 14.007
14  O = 15.999
15
16  total = C + 3*H + N + 2*O
17  print 'We calculate "by hand" the molecular weight of nitromethane to be {0:1.3f} g/mol'.format(total)
```

ASE and numpy calculates the molecular weight of nitromethane is 61.040 g/mol We calculate "by hand" the molecular weight of nitromethane to be 61.040 g/mol

2 Center of Mass

Write a python function to compute the center of mass of nitromethane. Compare your answer to the output of ase. Atoms.get_{centerof mass}.

The center of mass is defined as

$$R = \frac{1}{M} \sum m_i r_i \tag{1}$$

where M is the total mass and m_i and r_i is each chemical species' mass and distance from a fixed point. Note that R and r_i are given from the same fixed point, which we will define for simplicity as the origin. We will first write a program that calculates this.

```
from ase.data.molecules import molecule
2 import numpy as np
3
    # Prepare the data
5 atoms = molecule('CH3N02')
   masses = atoms.get_masses()
    positions = atoms.get_positions()
    total_mass = masses.sum()
8
   # Compute the center of mass (COM). First compute the sums of the m*r values
10
   sum = np.array((0., 0., 0.))
11
   for mass, pos in zip(masses, positions):
12
13
        sum += mass * pos
   COM = sum / total_mass
14
   print 'We compute the center of mass to be ({0:1.5f}, {1:1.5f}, {2:1.5f})'.format(COM[0], COM[1], COM[2])
15
16
17
    # Use ASE to compute the center of mass
    COM = atoms.get_center_of_mass()
    print 'ASE computes the center of mass to be ({0:1.5f}, {1:1.5f}, {2:1.5f})'.format(COM[0], COM[1], COM[2])
```

We compute the center of mass to be (0.00619, 0.07989, 0.00000)ASE computes the center of mass to be (0.00619, 0.07989, 0.00000)

3 Moments of Inertia

Compute the moments of inertia for nitromethane

```
from ase.data.molecules import molecule
atoms = molecule('CH3NO2')
I = atoms.get_moments_of_inertia()
print 'The moments of inertia are {0:1.3f}, {1:1.3f}, and {2:1.3f} amu*angstroms^2'.format(I[0], I[1], I[2])
```

The moments of inertia are 42.242, 47.838, and 86.868 amu*angstroms^2

4 Bond Lengths

1.087 Angstroms 1.087 Angstroms

Compute the bond length between the C and each H atom

```
from ase.data.molecules import molecule
1
   atoms = molecule('CH3NO2')
3
    print 'atom symbol'
    print '=====;
   for i, atom in enumerate(atoms):
       print i, ' ' + atom.symbol
    # From here we need to find distances between atom indexes 0-2, 0-3, and 0-4
10 indexes = ((0, 2), (0, 3), (0, 4))
   print 'The C-H bond distances are'
12
   for index in indexes:
       print '{0:1.3f}'.format(atoms.get_distance(index[0], index[1])), 'Angstroms'
    atom symbol
    ========
          C
          N
    1
          Η
          Η
    4
          Η
    5
          0
    The C-H bond distances are
    1.090 Angstroms
```

To understand why they are not equal, we can look at the molecule.

```
from ase.data.molecules import molecule
from ase.visualize import view
from ase.io import write

atoms = molecule('CH3N02')
atoms.set_cell((10, 10, 10))
atoms.center()
write('CH3N02.png', atoms, show_unit_cell=2)
write('CH3N02_rot.png', atoms, show_unit_cell=2, rotation='-90x,-90y, 90z')
```

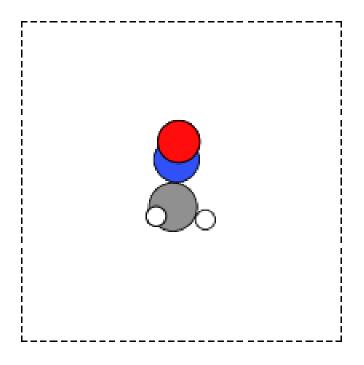


Figure 1: Side view of $\mathrm{CH}_{3\mathrm{NO}2}$

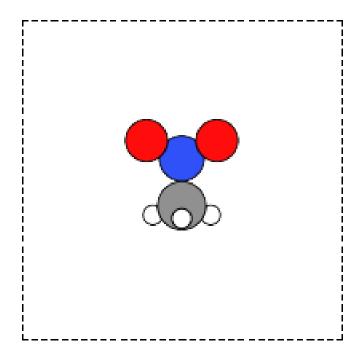


Figure 2: Side view of CH_{3NO2}

We can see that the molecule is symmetric along a plane that splites the two oxygens. These therefore will have the same bond length.

5 Bond angle in the nitro group

Compute the bond angle in degrees between O-N-O in the nitro group. From the previous problem, we know the indexes of the O-N-O bond are $6,\,1,\,\mathrm{and}\,5$

```
from ase.data.molecules import molecule
from numpy import pi

atoms = molecule('CH3N02')
s = 'The angle between O-N-O is {0:1.1f} degrees'
print s.format(atoms.get_angle([6,1,5])*180/pi)
```

The angle between O-N-O is 125.7 degrees

6 Generate an xyz file

Use ase to generate an xyz file of the coordinates of the nitromethane molecule. Include the output of your xyz file in your homework.

```
from ase.data.molecules import molecule
  from ase.io import write
4 atoms = molecule('CH3NO2')
  write('CH3NO2.xyz', atoms)
5
  file = open('CH3NO2.xyz', 'r')
 lines = file.readlines()
  for line in lines:
      print line[0:-1] #We want this to avoid the extra new line at the end of each line
    7
    C
            -0.114282000000000
                                      -1.314565000000000
                                                                  0.00000000000000
    N
             0.00000000000000
                                       0.166480000000000
                                                                  0.00000000000000
             0.899565000000000
    Η
                                      -1.715256000000000
                                                                  0.00000000000000
    Η
            -0.640921000000000
                                      -1.607212000000000
                                                                  0.904956000000000
    Η
            -0.640921000000000
                                      -1.607212000000000
                                                                 -0.904956000000000
    0
             0.066748000000000
                                       0.728232000000000
                                                                 -1.103775000000000
    0
             0.066748000000000
                                       0.728232000000000
                                                                  1.103775000000000
```

7 Create a graphic of nitromethane

The molecule should be centered in a unit cell with dimensions $$10 \times 11.52.1 \text{\AA}$. The unit cell should be visible

```
from ase.data.molecules import molecule
from ase.visualize import view
from ase.io import write

atoms = molecule('CH3N02')
atoms.set_cell((10, 11.5, 12.1))
atoms.center()
write('CH3N02_image.png', atoms, show_unit_cell=2, rotation='-45x,-45y, 45z')
```

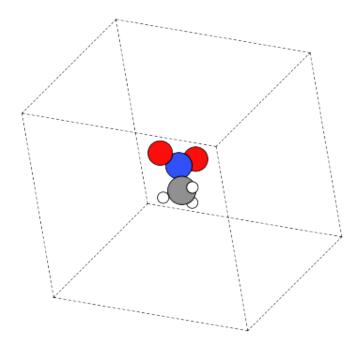


Figure 3: Graphic of nitromethane