

# Structural optimizations

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# Acetylene molecule ( $C_2H_2$ )

We want to use pw.x to find the optimized geometry of acetylene.

Let us suppose we have the following guess for the structure (coordinates in atomic units):

C	0.000000000	0.000000000	0.000000000
C	2.100000000	0.000000000	0.000000000
H	-1.600000000	0.400000000	0.000000000
H	3.600000000	-0.400000000	0.000000000

**pw.x** adopts periodic boundary conditions.

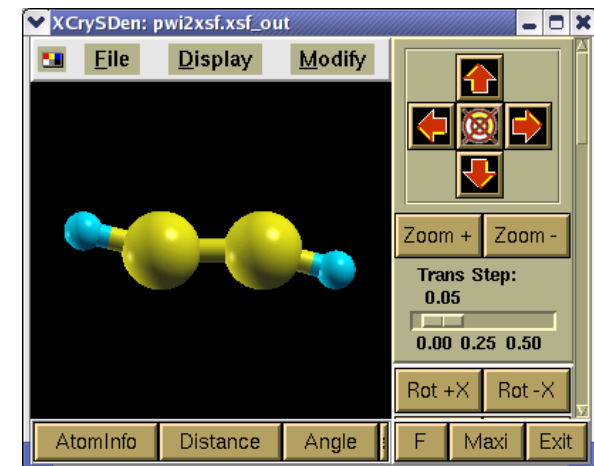
We can put the molecule in a large box, whose size is a convergency parameter.

Here, we choose a cube of side length 12 a.u. for quickness. Then:

1) Prepare the input file for the SCF calculation (see next slide for other details), use PWgui if you prefer

2) Visualize the structure with XcrySDen  
`xcrysdn --pwi acetylene.scf.in`

3) Run the SCF calculation and look at forces on atoms.  
`pw.x < acetylene.scf.in > acetylene.scf.out`



# C<sub>2</sub>H<sub>2</sub> SCF: the input

&CONTROL

```

  calculation = 'scf' ,
  restart_mode = 'from_scratch' ,
  outdir = 'your directory for temporary files/' ,

```

**Change this!**

/

&SYSTEM

```

  ibrav = 1,
  celldm(1) = 12.D0,
  nat = 4,
  ntyp = 2,
  ecutwfc = 25.0D0 ,
  ecutrho = 160.0D0 ,
  nosym = .true. ,
  occupations = 'smearing' ,
  degauss = 0.005D0 ,

```

**Why do we treat acetylene as a metal?**

/

&ELECTRONS

```

  conv_thr = 1.D-7 ,
  mixing_beta = 0.5D0 ,

```

/

ATOMIC\_SPECIES

```

  H 1.00000 H.US_PBE.RRKJ3.UPF
  C 12.00000 C.US_PBE.RRKJ3.UPF

```

**Do you have these pseudopotentials?**

ATOMIC\_POSITIONS bohr

```

  C 0.000000000 0.000000000 0.000000000 0 0 0
  C 2.100000000 0.000000000 0.000000000 1 1 1
  H -1.600000000 0.400000000 0.000000000 1 1 1
  H 3.600000000 -0.400000000 0.000000000 1 1 1

```

**This atom is locked**

K\_POINTS gamma

**Gamma point only is sufficient for the molecule-in-a-box approach**

# C<sub>2</sub>H<sub>2</sub> SCF: the output

From the output file:

Forces acting on atoms (Ry/au):

atom	1	type	2	force =	0.04225946	-0.07108502	0.00000000
atom	2	type	2	force =	-0.29082443	0.14196405	0.00000000
atom	3	type	1	force =	-0.50847780	0.10558208	0.00000000
atom	4	type	1	force =	0.75704277	-0.17646111	0.00000000

Total force = 0.989284

Total SCF correction = 0.000255

Forces on atoms are quite large.

We now optimize the structure by two ways:

A) damped molecular dynamics

B) BFGS minimization

**PAY ATTENTION !**

This number measures the error on forces given by a non perfect self-consistency.

It has to be much smaller than the force itself.

Connected to `conv_thr`.

Copy the input file for SCF to a new file, and open it with PWgui:

```
cp acetylene.scf.in acetylene.damp.in
```

```
pwgui
```

# C<sub>2</sub>H<sub>2</sub> damped dynamics

Modify the new input file `acetylene.damp.in`: use an editor, or PWgui to

1) Specify the following variables in the “Control” namelist:

```
calculation = "relax",  
dt          = 20.D0,
```

2) Set a stricter SCF threshold (“Electrons” namelist):

```
conv_thr = 1.D-8,
```

3) Add “Ions” namelist:

```
&IONS  
ion_dynamics      = "damp",  
pot_extrapolation = "second_order", (these extrapolations make  
wfc_extrapolation = "second_order", the SCF loop shorter)  
/  

```

4) Set equal masses (we are not interested in a real dynamics!)

```
H  1.0  H.US_PBE.RRKJ3.UPF  
C  1.0  C.US_PBE.RRKJ3.UPF
```

5) Save and run:

```
pw.x < acetylene.damp.in > acetylene.damp.out
```

# C<sub>2</sub>H<sub>2</sub> damped dynamics

Examine the output:

1) Scroll the file:

```
less acetylene.damp.out (press q to exit)
```

2) Extract key quantities:

```
total energy: grep ! acetylene.damp.out | nl
```

```
total force: grep "Total force" acetylene.damp.out | nl
```

3) Use XCrySDen to visualize the dynamics:

```
xcrysdn --pwo acetylene.damp.out
```

(remember to select “reduce dimension to 0D” and

“Display All Coordinates as Animation”; type “f” to visualize the forces)

At convergence, forces are smaller than the specified (or default) threshold, and the algorithm stops:

Forces acting on atoms (Ry/au):

atom	1	type	2	force =	-0.00052921	-0.00029862	0.00000046
atom	2	type	2	force =	0.00029473	0.00021132	-0.00000059
atom	3	type	1	force =	0.00006651	-0.00020371	-0.00000019
atom	4	type	1	force =	0.00016798	0.00029100	0.00000032

Total force = 0.000539 Total SCF correction = 0.000021

Damped Dynamics: convergence achieved in 17 steps

# C<sub>2</sub>H<sub>2</sub> BFGS minimization

1) Copy the input file for damped dynamics to a new file, and open it with PWgui:

```
cp acetylene.damp.in acetylene.bfgs.in  
pwgui
```

2) In “Ions” namelist

change:

```
ion_dynamics = 'damp' ,
```

to:

```
ion_dynamics = 'bfgs' ,
```

3) Set a variable SCF threshold

(100 times more accurate closer to the minimum of energy):

In “Electrons” namelist

Change:

```
conv_thr = 1.D-8 ,
```

To:

```
conv_thr = 1.D-6 ,
```

In “Ions” namelist

Add:

```
upscale = 100 ,
```

4) Run the calculation

```
pw.x < acetylene.bfgs.in > acetylene.bfgs.out
```

5) Look at the output

6) Compare final positions, energy, and number of steps with damped dynamics

# Structural optimization of a slab

We want now to optimize the geometry of an Al(001) slab.

- 1) Take a look at the provided input file for SCF calculation, `al001.scf.in`.  
 we have five atomic layers at the bulk-truncated positions (`alat` units):

Al	0.000000000	0.000000000	2.828426000	1	1	1
Al	0.500000000	0.500000000	2.121320000	1	1	1
Al	0.000000000	0.000000000	1.414213000	1	1	1
Al	0.500000000	0.500000000	0.707107000	0	0	0
Al	0.000000000	0.000000000	0.000000000	0	0	0

(The bottom two layers are fixed)

Visualize the structure with XCrySDen (show multiple unit cells, press `shift-n`)

- 2) Copy to a new file `cp al001.scf.in al001.bfgs.in`  
 and edit it by adding the variables for the structural optimization, as seen before.

- 4) Run the calculation `pw.x < al001.bfgs.in > al001.bfgs.out`

- 5) Look at the optimized geometries:

Al	-0.000000019	-0.000000008	2.757233372
Al	0.499999997	0.499999999	2.089582133
Al	-0.000000002	-0.000000001	1.397022977
Al	0.500000000	0.500000000	0.707107000
Al	0.000000000	0.000000000	0.000000000

- 6) Compare with experiments (LEED), but before check all the convergency parameters, which for this exercise are not accurate enough!