



Université de Liège

Introduction to run Siesta

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Our method



Linear-scaling DFT based on NAOs (Numerical Atomic Orbitals)

P. Ordejon, E. Artacho & J. M. Soler, Phys. Rev. B 53, R10441 (1996)

J. M. Soler *et al*, J. Phys.: Condens. Matter **14**, 2745 (2002)

- *Born-Oppenheimer* (relaxations, mol.dynamics)
- *DFT* (LDA, GGA)
- *Pseudopotentials* (norm conserving, factorised)
- *Numerical atomic orbitals as basis* (finite range)
- *Numerical evaluation of matrix elements* (3Dgrid)

*Implemented in the **SIESTA** program*

D. Sanchez-Portal, P. Ordejon, E. Artacho & J. M. Soler
Int. J. Quantum Chem. 65, 453 (1997)

To run **Siesta** you need:

1.- Access to the **executable file**

2.- An **input file**

Flexible Data Format (FDF) (A. García and J. M. Soler)

3.- A **pseudopotential file** for each kind of element in the input file

Unformatted binary (**.vps**)

Formatted ASCII (**.psf**) (more transportable and easy to look at)

Siesta package:

- **Src:** Sources of the Siesta code
- **Docs:** Documentation and user conditions
User's Guide (siesta.tex)
- **Pseudo:** ATOM program to generate and test pseudos
(A. García; *Pseudopotential and basis generation*, Tu 12:00)
- **Examples:** fdf and pseudopotentials input files for simple systems
- **Utils:** Programs or scripts to analyze the results

The input file

Main input file:

- Physical data of the system
- Variables to control the approximations
- Flexible Data Format (FDF)

developped by A. García and J. M. Soler

FDF (I)

- Data can be given in **any order**
- Data can be **omitted** in favour of **default values**
- Syntax: 'data label' followed by its value

Character string:	SystemLabel	h2o
Integer:	NumberOfAtoms	3
Real:	PAO.SplitNorm	0.15
Logical:	SpinPolarized	.false.
Physical magnitudes	LatticeConstant	5.43 Ang

FDF (II)

- Labels are **case insensitive** and characters `-_.` are **ignored**

`LatticeConstant` is equivalent to `lattice_constant`

- Text following `#` are **comments**
- **Logical** values: `T` , `.true.` , `true` , `yes`

`F` , `.false.` , `false` , `no`

- **Character** strings, **NOT** in apostrophes
- **Complex** data structures: **blocks**

`%block label`

`...`

`%endblock label`

FDF (III)

- **Physical magnitudes:** followed by its **units**.

Many physical units are recognized for each magnitude

(Length: m, cm, nm, Ang, bohr)

Automatic conversion to the ones internally required.

- You may **'include'** other FDF files or **redirect** the search to another file

Basic input variables

- 1.- General system descriptors
- 2.- Structural and geometrical variables
- 3.- Functional and solution method
- 4.- Convergence of the results
- 5.- Self-consistency

(Basis set generation related variables:

A. García; *Pseudopotential and basis generation*, Tu 12:00)

General system descriptor

SystemName: descriptive name of the system

SystemName Si bulk, diamond structure

SystemLabel: nickname of the system to name output files

SystemLabel Si

(After a successful run, you should have files like

Si.DM : Density matrix

Si.XV: Final positions and velocities

...)

Structural and geometrical variables

NumberOfAtoms: number of atoms in the simulation

```
NumberOfAtoms 2
```

NumberOfSpecies: number of different atomic species

```
NumberOfSpecies 1
```

ChemicalSpeciesLabel: specify the different chemical species.

```
%block ChemicalSpeciesLabel
```

```
1 14 Si
```

```
%endblock ChemicalSpeciesLabel
```

ALL THESE VARIABLES ARE MANDATORY

Periodic Boundary Conditions (PBC)

Atoms in the unit cell are **periodically repeated throughout space** along the lattice vectors

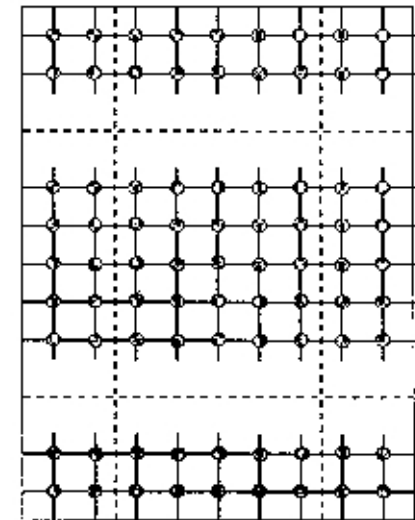
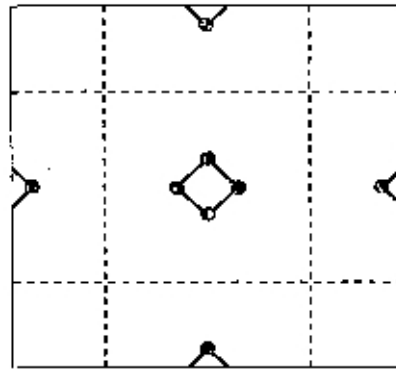
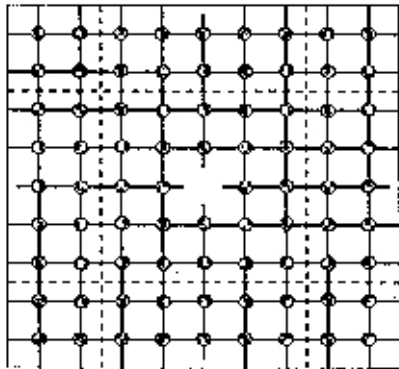
Periodic systems and crystalline solids:

Aperiodic systems: Supercell approximation

Defects

Molecules

Surfaces



M. C. Payne *et al*, Rev. Mod. Phys., **64**, 1045 (92)

Lattice Vectors

LatticeConstant: real length to define the scale of the lattice vectors

```
LatticeConstant    5.43 Ang
```

LatticeParameters: Crystallographic way

```
%block LatticeParameters  
    1.0  1.0  1.0  60.  60.  60.  
%endblock LatticeParameters
```

LatticeVectors: read as a matrix, each vector being a line

```
%block LatticeVectors  
    0.0  0.5  0.5  
    0.5  0.0  0.5  
    0.5  0.5  0.0  
%endblock LatticeVectors
```

Atomic Coordinates

AtomicCoordinatesFormat: format of the atomic positions in input:

Bohr: cartesian coordinates, in bohrs

Ang: cartesian coordinates, in Angstroms

ScaledCartesian: cartesian coordinates, units of the lattice constant

Fractional: referred to the lattice vectors

AtomicCoordinatesFormat Fractional

AtomicCoordinatesAndAtomicSpecies:

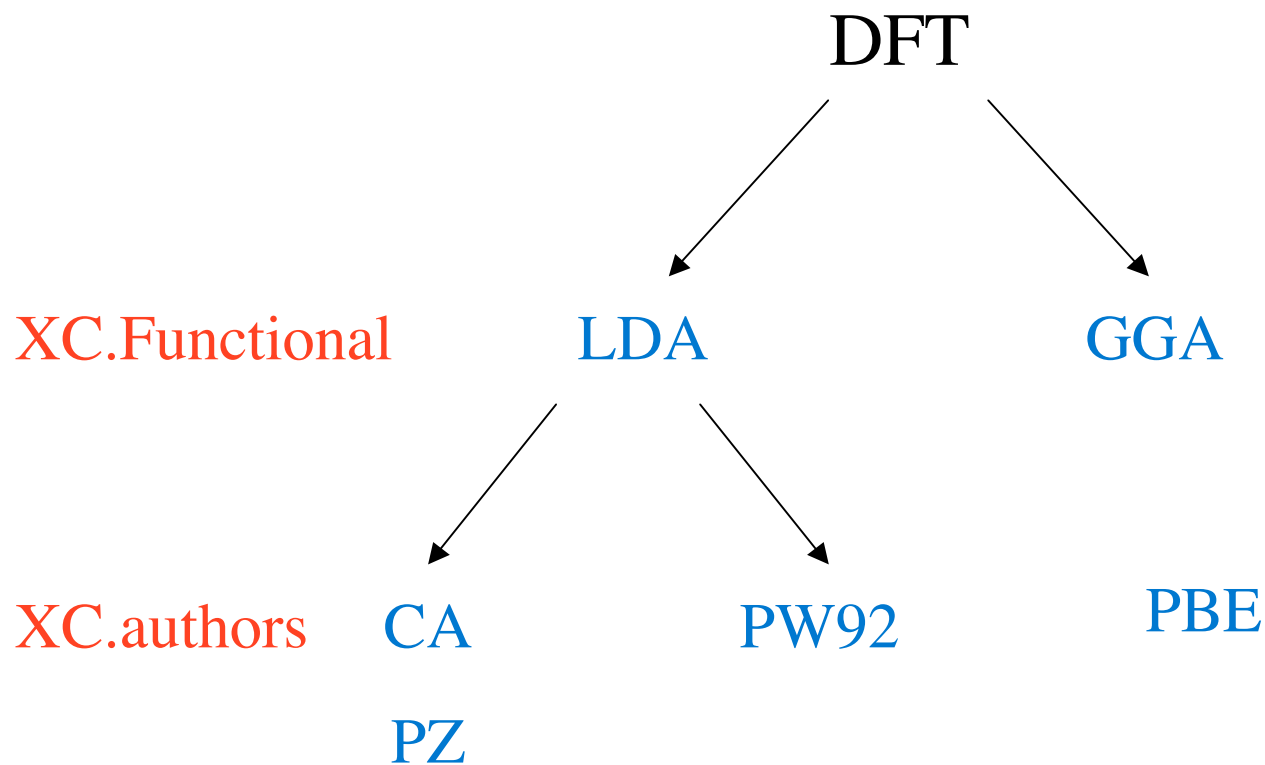
```
%block AtomicCoordinatesAndAtomicSpecies
```

```
0.00 0.00 0.00 1
```

```
0.25 0.25 0.25 1
```

```
%endblock AtomicCoordinatesAndAtomicSpecies
```

Functional



SpinPolarized

DFT ≡ Density Functional Theory

LDA ≡ Local Density Approximation

GGA ≡ Generalized Gradient Approximation

CA ≡ Ceperley-Alder

PZ ≡ Perdew-Zunger

PW92 ≡ Perdew-Wang-92

PBE ≡ Perdew-Burke-Ernzerhof

Solution method

From the atomic coordinates and the unit cell

$$\{\vec{R}, \vec{a}\}$$

Order N operations

Hamiltonian, H , and Overlap, S , matrices

$$(H \square \square S)C = 0$$

SolutionMethod

diagon

Order-N

E. Artacho, *Running with Order-N*,
Wed 11:40

k-sampling

Many magnitudes require integration of Bloch functions over Brillouin zone (BZ)

$$\chi(\vec{r}) = \int_i \int_{BZ} d\vec{k} n(\vec{k}) |\chi_i(\vec{k})|^2$$

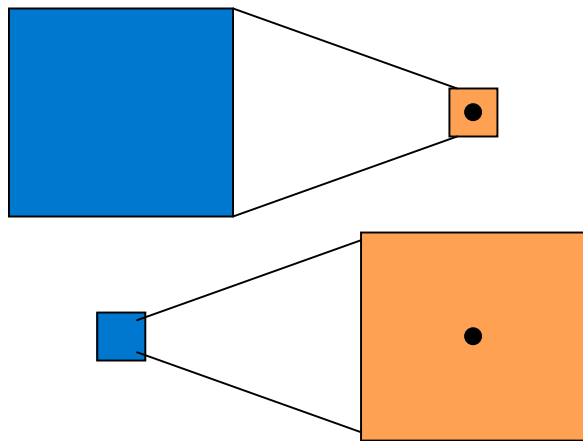
In practice: integral \int \int sum over a finite uniform grid

Essential for:

Small systems

Metals

Magnetic systems



Good description of the Bloch states at the Fermi level

Even in same insulators:

Perovskite oxides

Real space \square Reciprocal space

k-sampling

Spetial set of k-points: **Accurate results** for a **small #** k-points:

Baldereschi, Chadi-Cohen, **Monkhorst-Pack**

kgrid_cutoff:

```
kgrid_cutoff    10.0 Ang
```

kgrid_Monkhorst_Pack:

```
%block kgrid_Monkhorst_Pack
```

```
4  0  0  0.5
```

```
0  4  0  0.5
```

```
0  0  4  0.5
```

```
%endblock kgrid_Monkhorst_Pack
```

$$\rho(\vec{r}) = \sum \rho^{atom}(\vec{r})$$

Initial guess

$$\rho(\vec{r}) = \sum_{\alpha, \beta} \rho_{\alpha, \beta}$$

$$V_H(\vec{r}), V_{xc}(\vec{r})$$

Self-consistent iterations

Mixing $\rho_{\alpha, \beta}^{out}, \rho_{\alpha, \beta}^{in}$

Linear: **DM.MixingWeight**

NonLinear (Pulay): **DM.NumberPulay**

$$\rho_{\alpha, \beta}^{out}$$

$$\hat{H}\psi = \epsilon\psi$$

MaxSCFIterations

$$\rho_{\alpha, \beta}^{out} - \rho_{\alpha, \beta}^{in} <$$

DM.Tolerance

Total energy
Charge density
Forces

How to run Siesta

To run the **serial** version:

```
[path]siesta < myinput.fdf > myoutput &
```

To see the information dumped in the
output file during the run:

```
tail -f myoutput
```

Output: the header

```
SIESTA 1.2.3 -- [iorho parallel fix/0(N)] (Nov 20, 2001)
Architecture : lahey
Compiler flags: lf95 -O --warn --quiet --tpp --ntrace
SERIAL version
```

```
* Running in serial mode
```

```
>> Start of run: 3-JUL-2002 17:06:18
```

Output: dumping the input file

```
*: ***** Dump of input data file *****
SystemName          Water molecule
SystemLabel         h2o
NumberOfAtoms       3
NumberOfSpecies     2
%block ChemicalSpeciesLabel
  1  8  0      # Species index, atomic number, species label
  2  1  H
%endblock ChemicalSpeciesLabel
AtomicCoordinatesFormat Ang
%block AtomicCoordinatesAndAtomicSpecies
  0.000  0.000  0.000  1
  0.757  0.586  0.000  2
 -0.757  0.586  0.000  2
%endblock AtomicCoordinatesAndAtomicSpecies
*: ***** End of input data file *****
```

Output: processing the input

```
prinput: *****
coor: Atomic-coordinates input format = Cartesian coordinates
      coor: (in Angstroms)
redata: Number of spin components = 1
redata: Long output = F
redata: Number of Atomic Species = 2
redata: Charge density info will appear in .RHO file
redata: Write Mulliken Pop. = NO
redata: Mesh Cutoff = 50.0000 Ry
redata: Net charge of the system = 0.0000 |e|
redata: Max. number of SCF Iter = 50
redata: Mixing is linear
redata: Mix DM in first SCF step ? = F
redata: Write Pulay info on disk? = F
redata: New DM Mixing Weight = 0.2500
redata: No kicks to SCF
redata: DM Mixing Weight for Kicks = 0.5000
redata: DM Tolerance for SCF = 0.000100
redata: Use continuation files for DM = F
redata: Neglect nonoverlap interactions = F
redata: Method of Calculation = Diagonalization
redata: Electronic Temperature = 0.0019 Ry
redata: Fix the spin of the system = F
redata: Dynamics option = Verlet MD run
redata: Initial MD time step = 1
redata: Final MD time step = 1
redata: Length of MD time step = 1.0000 fs
redata: Length of MD time step = 1.0000 fs
redata: Initial Temperature of MD run = 0.0000 K
redata: Perform a MD quench = F
redata: *****:
```

Output: coordinates and k-sampling

```
siesta: Atomic coordinates (Bohr) and species
siesta:      0.00000  0.00000  0.00000  1      1
siesta:      1.43052  1.10738  0.00000  2      2
siesta:     -1.43052  1.10738  0.00000  2      3

siesta: Automatic unit cell vectors (Ang):
siesta:      7.286412  0.000000  0.000000
siesta:      0.000000  5.746952  0.000000
siesta:      0.000000  0.000000  5.621012

siesta: System type = molecule
...

siesta: System type = bulk

siesta: k-grid: Number of k-points = 196
siesta: k-grid: Cutoff = 14.021 Ang
siesta: k-grid: Supercell and displacements
siesta: k-grid:   7  0  0  0.000
siesta: k-grid:   0  7  0  0.000
siesta: k-grid:   0  0  7  0.000
```


Output: First MD step

```
siesta: =====  
siesta:          Begin MD step =          1  
siesta: =====  
  
InitMesh: MESH =      32 x      30 x      24 =      23040  
InitMesh: Mesh cutoff (required, used) =      50.000      50.384 Ry  
  
* Maximum dynamic memory allocated =      3 MB  
  
siesta: Program's energy decomposition (eV):  
siesta: Eions      =      815.854478  
siesta: Ena        =      175.154399  
siesta: Ekin       =      341.667405  
siesta: Enl        =      -52.736793  
siesta: DEna       =      -0.000001  
siesta: DUscaf     =      0.000000  
siesta: DUext      =      0.000000  
siesta: Exc        =     -109.951257  
siesta: eta*DQ     =      0.000000  
siesta: Emadel     =      0.000000  
siesta: Eharris    =     -466.430254  
siesta: Etot       =     -461.720725  
siesta: FreeEng    =     -461.720725
```

Output: Self-consistency

```
siesta: iscf Eharris(eV)    E_KS(eV) FreeEng(eV)    dDmax  Ef(eV)
siesta:   1  -466.4303    -461.7207  -461.7207  1.4383 -4.2475
timer: Routine,Calls,Time,% = IterSCF           1      7.930 72.22
siesta:   2  -466.8703    -465.2425  -465.2425  0.1755 -0.1474
siesta:   3  -465.9264    -465.4655  -465.4655  0.0515 -1.5862
siesta:   4  -465.8472    -465.5656  -465.5656  0.0176 -1.9935
siesta:   5  -465.8397    -465.6346  -465.6346  0.0087 -2.1116
siesta:   6  -465.8388    -465.6857  -465.6857  0.0083 -2.1448
siesta:   7  -465.8387    -465.7240  -465.7240  0.0067 -2.1531
siesta:   8  -465.8387    -465.7527  -465.7527  0.0051 -2.1545
siesta:   9  -465.8387    -465.7742  -465.7742  0.0038 -2.1543
siesta:  10  -465.8387    -465.7903  -465.7903  0.0028 -2.1539
siesta:  11  -465.8387    -465.8024  -465.8024  0.0021 -2.1535
siesta:  12  -465.8387    -465.8115  -465.8115  0.0016 -2.1533
siesta:  13  -465.8387    -465.8183  -465.8183  0.0012 -2.1531
siesta:  14  -465.8387    -465.8234  -465.8234  0.0009 -2.1530
siesta:  15  -465.8387    -465.8272  -465.8272  0.0006 -2.1530
siesta:  16  -465.8387    -465.8301  -465.8301  0.0005 -2.1530
siesta:  17  -465.8387    -465.8322  -465.8322  0.0004 -2.1530
siesta:  18  -465.8387    -465.8338  -465.8338  0.0003 -2.1530
siesta:  19  -465.8387    -465.8351  -465.8351  0.0002 -2.1530
siesta:  20  -465.8387    -465.8360  -465.8360  0.0001 -2.1530
siesta:  21  -465.8387    -465.8367  -465.8367  0.0001 -2.1530
siesta:  22  -465.8387    -465.8372  -465.8372  0.0001 -2.1530
```

Output: Eigenvalues, forces, stress

siesta: Eigenvalues (eV):

ik	is	eps					
1	1	-24.74	-12.70	-8.71	-6.23	1.68	4.09
		14.68	21.97	24.22	27.21	28.65	32.19
		49.89	70.65	96.18			

siesta: Atomic forces (eV/Ang):

siesta:	1	0.000001	-0.504870	0.000000
siesta:	2	0.719664	0.279830	0.000000
siesta:	3	-0.719663	0.279829	0.000000
siesta:	-----			
siesta:	Tot	0.000002	0.054788	0.000000

siesta: Stress tensor (eV/Ang**3):

siesta:	-0.012622	0.000000	0.000000
siesta:	0.000000	-0.002309	0.000000
siesta:	0.000000	0.000000	0.014000

Output: Total energy

siesta: Fermi energy = -2.152975 eV

siesta: Program's energy decomposition (eV):

siesta:-Eions	=	-815.854478
siesta: Ena	=	175.154399
siesta: Ekin	=	350.784945
siesta: Enl	=	-61.958840
siesta: DEna	=	-1.777979
siesta: DUsf	=	0.727284
siesta: DUext	=	0.000000
siesta: Exc	=	-112.912881
siesta: eta*DQ	=	0.000000
siesta: Emadel	=	0.000000
siesta: Ekinion	=	0.000000
siesta: Eharris	=	-465.839084
siesta: Etot	=	-465.837551
siesta: FreeEng	=	-465.837551

siesta: Final energy (eV):

siesta: Kinetic	=	350.784945
siesta: Hartree	=	382.616610
siesta: Ext. field	=	0.000000
siesta: Exch.-corr.	=	-112.912881
siesta: Ion-electron	=	-1072.820417
siesta: Ion-ion	=	-13.505807
siesta: Ekinion	=	0.000000
siesta: Total	=	-465.837551

Output: timer

```
timer: CPU execution times:
timer: Routine   Calls Time/call  Tot.time   %
timer: siesta    1    13.660   13.660 100.00
timer: Setup     1     0.850    0.850  6.22
timer: bands     1     0.000    0.000  0.00
timer: KSV_init  1     0.000    0.000  0.00
timer: IterMD    1    12.800   12.800 93.70
timer: hsparse   2     0.005    0.010  0.07
timer: overfsm   2     1.095    2.190 16.03
timer: IterSCF   23    0.461   10.600 77.60
timer: kinefsm   2     1.010    2.020 14.79
timer: nlefsm    2     2.780    5.560 40.70
timer: DHSCF     23    0.128    2.950 21.60
timer: DHSCF1    1     0.060    0.060  0.44
timer: DHSCF2    1     0.190    0.190  1.39
timer: REORD     186   0.001    0.130  0.95
timer: POISON    24    0.020    0.480  3.51
timer: DHSCF3    23    0.110    2.520 18.45
timer: rhoofd    23    0.030    0.690  5.05
timer: CELLXC    23    0.027    0.610  4.47
timer: vmat      23    0.018    0.410  3.00
timer: diagonal  22    0.002    0.050  0.37
timer: rdiag     22    0.002    0.040  0.29
timer: DHSCF4    1     0.180    0.180  1.32
timer: dfscf     1     0.150    0.150  1.10
```

```
>> End of run: 3-JUL-2002 17:06:32
```

Saving and reading information (I)

Some information is stored by Siesta to restart simulations from:

- Density matrix: **DM.UseSaveDM**
- Localized wave functions (Order-N): **ON.UseSaveLWF**
- Atomic positions and velocities: **MD.UseSaveXV**
- Conjugent gradient history (minimizations): **MD.UseSaveCG**

All of them are **logical variables**

EXTREMELY USEFUL TO SAVE LOT OF TIME!

Saving and reading information (II)

Information needed as input for various post-processing programs,

for example, to visualize:

- Total charge density: `SaveRho`
- Deformation charge density: `SaveDeltaRho`
- Electrostatic potential: `SaveElectrostaticPotential`
- Total potential: `SaveTotalPotential`
- Local density of states: `LocalDensityOfStates`
- Charge density contours: `WriteDenchar`
- Atomic coordinates: `WriteCoorXmol` and `WriteCoorCerius`

All of them are `logical variables`

Analyzing the electronic structure (I)

- **Band structure** along the high symmetry lines of the BZ

BandLineScale: scale of the k vectors in BandLines

```
BandLineScale    pi/a
```

BandLines: lines along with band energies are calculated.

```
%block BandLines
```

```
1    1.000    1.000    1.000    L
```

```
20   0.000    0.000    0.000    \Gamma
```

```
25   2.000    0.000    0.000    X
```

```
30   2.000    2.000    2.000    \Gamma
```

```
%endblock BandLines
```


Analyzing the electronic structure (II)

• **Density of states:** total and projected on the atomic orbitals

- Compare with experimental spectroscopy

- Bond formation

- Defined as:

$$g(\epsilon) = \sum_i \sum_{\mathbf{k}} \delta(\epsilon - \epsilon_i(\mathbf{k}))$$
$$\simeq \sum_i \sum_{\mathbf{k}} \frac{1}{\sigma\sqrt{\pi}} \exp\left(-\frac{(\epsilon - \epsilon_i(\mathbf{k}))^2}{\sigma^2}\right)$$

ProjectedDensityOfStates:

```
%block ProjectedDensityOfStates
```

```
-20.00 10.00 0.200 500 eV
```

```
%endblock ProjectedDensityOfStates
```

Analyzing the electronic structure (III)

- **Population analysis: Mulliken** prescription

- Amounts of charge on an atom or in an orbital inside the atom
- Bond formation
- Be careful, very dependent on the basis functions

WriteMullikenPop

WriteMullikenPop 0 = None

1 = Atomic and orbitals charges

2 = 1 + atomic overlap pop.

3 = 2 + orbital overlap pop.

Tools (I)

- Various **post-processing programs**:

- **PHONONS**:

- Finite differences: **VIBRA** (P. Ordejón)

- Linear response: **LINRES** (J. M. Alons-Pruneda et al.)

- **Interphase** with **Phonon** program (Parlinsky)

- Visualize of the **CHARGE DENSITY** and **POTENTIALS**

- 3D: **PLRHO** (J. M. Soler)

- 2D: **CONTOUR** (E. Artacho)

- 2D: **DENCHAR** (J. Junquera)

Tools (II)

-TRANSPORT PROPERTIES:

-**TRANSIESTA** (M. Brandbydge *et al.*)

-PSEUDOPOTENTIAL and BASIS information:

-**PyAtom** (A. García)

-ATOMIC COORDINATES:

-**Sies2arc** (J. Gale)