How to run SIESTA

Introduction to input & output files

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

- Born-Oppenheimer
- DFT
- Pseudopotentials
- Numerical atomic orbitals
- Numerical evaluation of matrix elements

- *relaxations, MD, phonons.*
- *LDA, GGA*
- *norm conserving, factorised.*
- *finite range*

siesta

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)

What do we need?

- Access to the executable file: siesta
- An input file
  - Flexible Data Format (fdf) *(A. García & J.M. Soler).*
- A psedopotential file for each kind of element in the input file.
  - Unformatted binary (vps)
  - Formatted ASCII (.psf): more transportable and readable

Siesta package: (under license)

- Src: Sources of the Siesta code
- Docs: Documentation and user conditions
  - User's Guide (siesta.tex)
- Pseudo: ATOM program to generate and test pseudopotentials
- Examples: fdf and pseudopotentials input files for simple systems
- Utils: Programs or scripts to analyze the results

The input file

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

FDF characteristics (II)

- Data can be given in any order
- Data can be omitted in favour of *default* values.
- Labels are case insensitive and characters `-´ , `_´, `.´ are ignored:

  LatticeConstant is equivalent to lattice_constant
- Text following # are comments
- You may 'include' other FDF files or redirect the search to another file

FDF characteristics (II)

- 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

- Physical magnitudes: followed by their units.

Many units are valid for the same magnitude (m, cm, nm, Ang, Bohr).
There is an automatic conversion to the units required internally.

- Character strings, NOT in apostrophes

FDF characteristics (III)

- Logical values: T / .true. / true / yes
  F / .false. / false / no

- Complex data structures: *blocks*

```
%block label
...
%endblock label
```

Basic input variables

a) General system descriptors

b) Structural and geometrical variables

c) Functional and solution mehod

d) Convergence of the results

e) Self-consistency

f) Basis set generation related variables.

General System description

SystemName: descriptive name of the system

SystemName	Si bulk, diamond structure

SystemLabel: nickname to label output files

SystemLabel	silicon

After a succesful run, you should have files like

silicon.DM : *Density matrix*

silicon.XV : *Final positions and velocities*

silicon.EIG : *Eigen-energies*

...

Geometrical and structural 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!!

Lattice Vectors

LatticeConstant: length to define the scale of the lattice vectors

```
LatticeConstant                 5.43 Ang
```

LatticeParameters: Crystallograhic way

```
%block LatticeParameters
  1.0   1.0   1.0   60.0   60.0   60.0
%endblock LatticeParameters
```

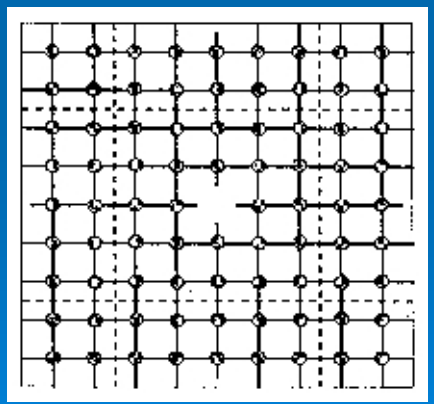
LatticeVectors: read as a matrix, each vector being a line

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

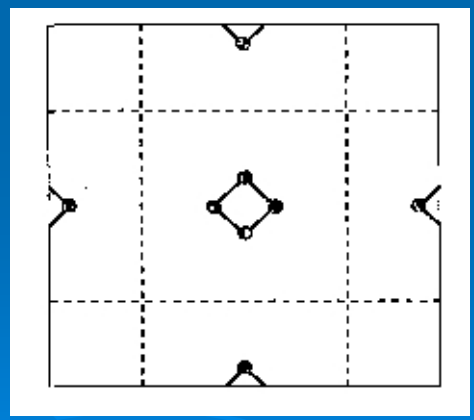
Periodic Boundary Conditions (PBC)

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

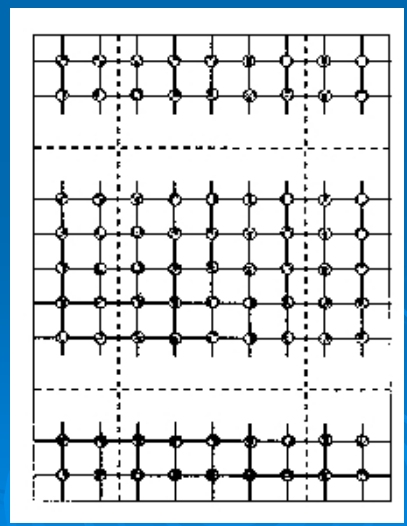
- Periodic systems and crystalline solids: √
- Aperiodic systems: *Supercell approximation*



Defects



Molecules



Surfaces

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:

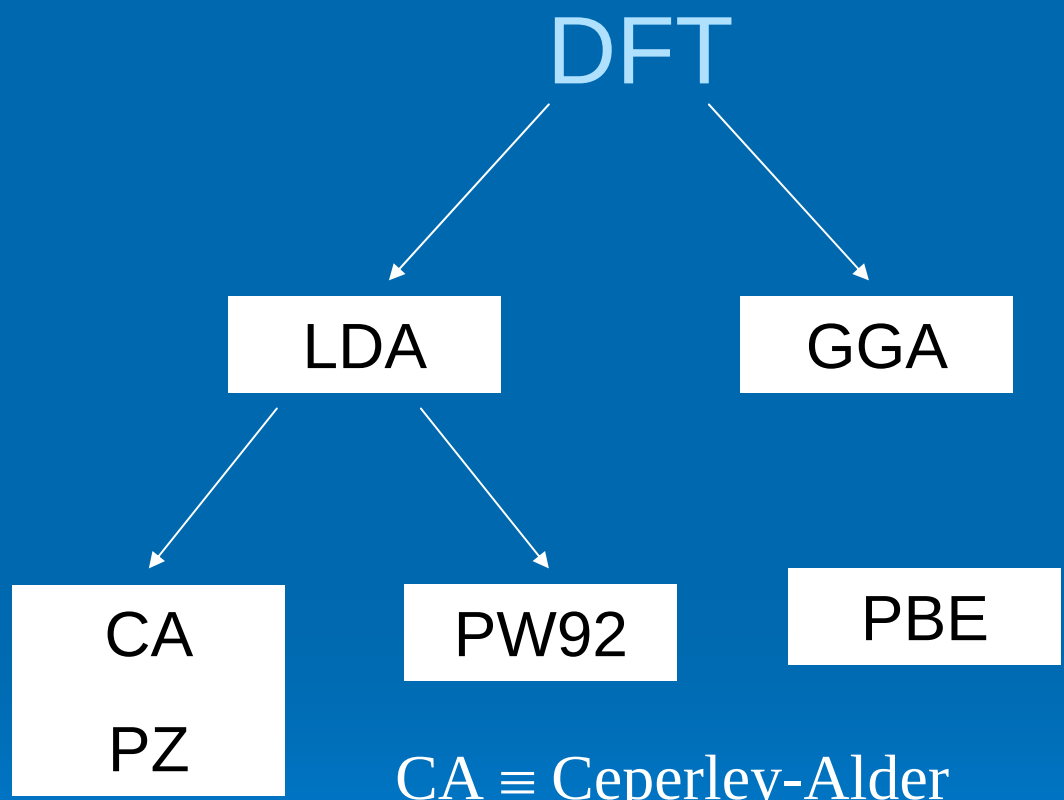
```
%block AtomicCoordinatesAndAtomicSpecies
        0.00    0.00    0.00    1       Si
        0.25    0.25    0.25    1       Si
%endblock AtomicCoordinatesAndAtomicSpecies
```

XC-functional

XC.Functional

XC.authors

SpinPolarized



CA ≡ Ceperley-Alder

PZ ≡ Perdew-Zunger

PW92 ≡ Perdew-Wang-92

PBE ≡ Perdew-Burke-Ernzerhof

Solution method

Two options to solve the electronic problem (Kohn-Sham equations)

From the atomic coordinates and the unit cell $\{\vec{R}, \vec{a}\}$

Order *N* operations

$$(H - \varepsilon S)C = 0$$

Hamiltonian, ***H***, and Overlap, ***S***, matrices

SolutionMethod

diagon

ordern

K-point sampling

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

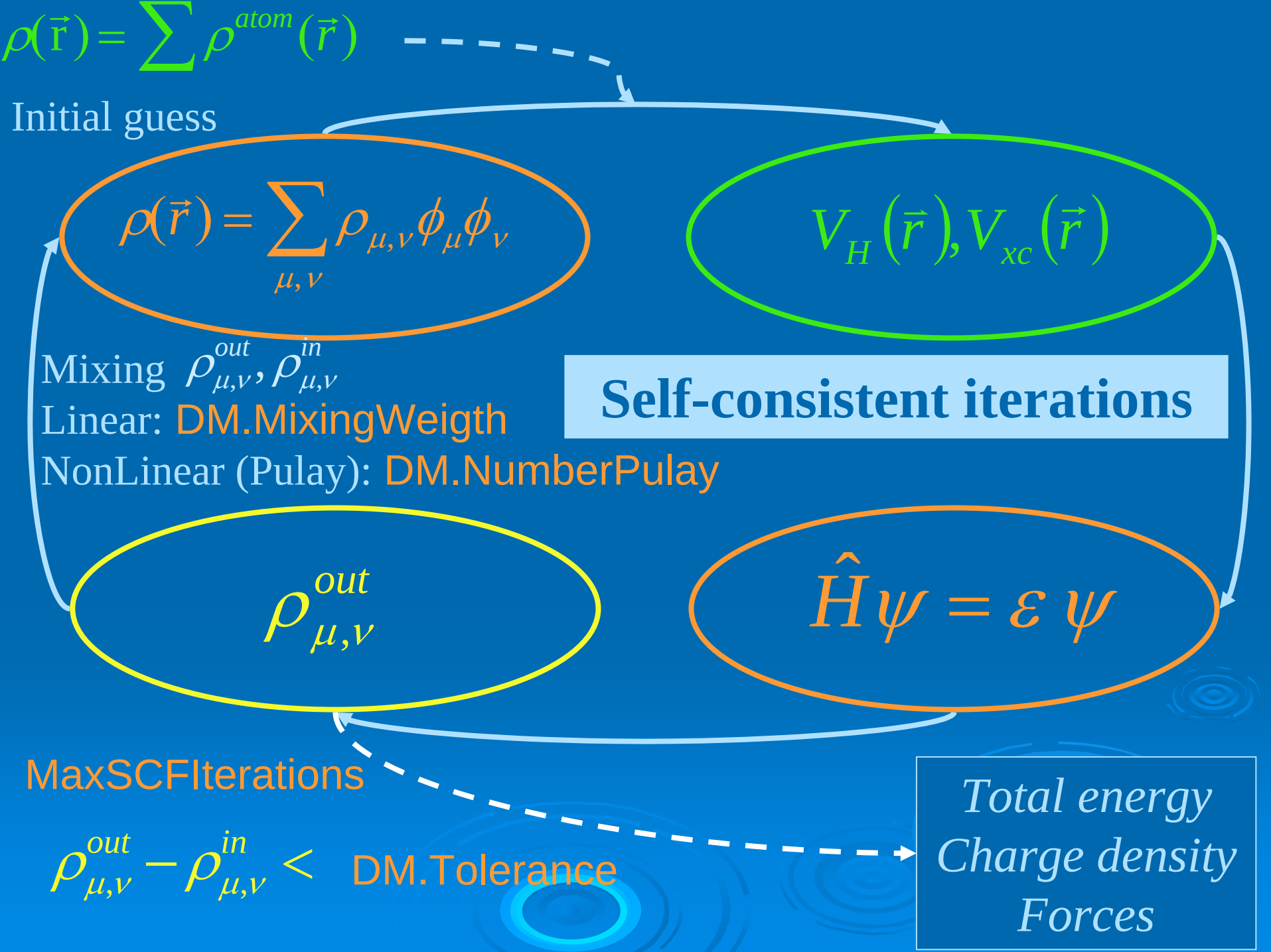
Baldereschi, Chadi-Cohen, Monkhorst-Pack

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
$\rho(\vec{r}) = \sum_{\mu,\nu} \rho_{\mu,\nu}\phi_\mu\phi_\nu$
$V_H(\vec{r}), V_{xc}(\vec{r})$
Mixing $\rho^{out}_{\mu,\nu}, \rho^{in}_{\mu,\nu}$
Linear: DM.MixingWeigth
NonLinear (Pulay): DM.NumberPulay
Self-consistent iterations
$\rho^{out}_{\mu,\nu}$
$\hat{H}\psi = \varepsilon\psi$
MaxSCFIterations
$\rho^{out}_{\mu,\nu} - \rho^{in}_{\mu,\nu} <$ DM.Tolerance
Total energy
Charge density
Forces

How to run siesta

- To run the serial version:

```
[path]/siesta < input.fdf > output &
```

- To see the information dumbed in the output file while it runs

```
tail –f output
```

- Alternatively:

```
[path]/siesta < input.fdf | tee ouput
```

Analysing the output (I)

The output header

Siesta version

Fortran compiler

```
SIESTA 1.2.3 -- [iorho parallel fix/O(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
```

Analysing the output (II)

dumping the input file

```
*: ******************** Dump of input data file ************************
SystemName          Water molecule
SystemLabel         h2o
NumberOfAtoms       3
NumberOfSpecies     2
%block ChemicalSpeciesLabel
 1  8  O      # 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 *************************
```

```
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: ***********************************************************
```

processing the input

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: DUscf   =         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
```

```
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: DUscf    =       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: Total energy

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:  diagon        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 & reading information (I)

Some information is stored by Siesta and can be used to restart the simulations from previous a run:

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

All of them are logical variables *(and save lots of time!!)*

Saving & 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 symetry lines of the BZ
  - BandLineScale: scale of the k vectors in BandLines

```
BandLinesScale      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

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

- ProjectedDensityOfStates:

```
%block ProjectedDensityOfStates
    -20.00   10.00   0.200   500   eV
%endblock ProjectedDensityOfStates
```

Lower energy — Higher energy — width — # points

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	0 = None 1 = Atomic and orbitals charges 2 = 1 + atomic overlap pop. 3 = 2 + orbital overlap pop.

Utilities (I)

Various post-processing tools:

- PHONONS:
  - Finite differences: VIBRA (P. Ordejón)
  - Linear response: LINRES (M. Pruneda et al.)
  - Interphase with Phonon program (Parlinsky)

- Visualization of the CHARGE DENSITY and POTENTIALS
  - -3D: PLRHO (J. M. Soler), grid2cube, grid2xfs, …
  - -2D: CONTOUR (E. Artacho)
  - -2D: DENCHAR (J. Junquera)

Utilities (II)

- TRANSPORT PROPERTIES:
  - TRANSIESTA (M. Brandbydge *et al.*),
  - SMEAGOL (S. Sanvito *et. al.*)
- PSEUDOPOTENTIAL and BASIS information:
  - PyAtom (A. García).
- XML output:
  - Visualization of the ouput (J. Wakelin & A. García).
  - PDOS-xml tool (A. García)
- ATOMIC COORDINATES:
  - Sies2arc (J. Gale)