

Basic introduction to running 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)

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Implemented in the SIESTA method:

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

Siesta resources (I)

- Web page: <http://www.uam.es/siesta>
 - Pseudos and basis database
 - Mailing list
 - Usage manual
- Soon:
 - Issue tracker (for bugs, etc)
 - Mailing list archives
 - Wiki

Siesta resources (2)

- Andrei Postnikov Siesta utils page:
[http://www.home.uni-osnabrueck.de/apostnik/
download.html](http://www.home.uni-osnabrueck.de/apostnik/download.html)
- Lev Kantorovich Siesta utils page:
[http://www.cmmmp.ucl.ac.uk/~lev/codes/lev00/
index.html](http://www.cmmmp.ucl.ac.uk/~lev/codes/lev00/
index.html)

Siesta software package:

- **Src**: Sources of the Siesta code.
- **Src/Sys**: makefiles for the compilation
- **Src/Tests**: A collection of tests.
- **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 11:10)
- **Examples**: fdf and pseudos input files for simple systems.
- **Tutorials**: Tutorials for basis and pseudo generation.
- **Utils**: Programs or scripts to analyze the results.

To run Siesta you need:

- 1.- Access to the **executable file**: T. White: “*And now that you are back at home ... what?*” Friday 13:00.
- 2.- An **input file**: written in ascii (plain text) using:
Flexible Data Format (FDF) (A. García and J. M. Soler)
- 3.- A **pseudopotential file** for each kind of element in the input file. Two differen't formats:
Unformatted binary (.vps)
Formatted ASCII (.psf) (more transportable and easy to look at)

Running siesta

Siesta has no windows, it is run from a UNIX terminal or from a MSDOS console.

Main input file: “name”.fdf

- Contents:

- Physical data of the system
- Variables to control the approximations

- Format:

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

FDF (I)

- Data can be given in **any order**
- Data can be **omitted** in favor 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.** , **yes**, **F** , **.false.** , **no**

By default logicals are true: **DM.UseSaveDM**

- **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, so for example in the main fdf it’s possible:

AtomicCoordinatesFormat < system_xyz.fdf

AtomicCoordinatesAndAtomicSpecies < system_xyz.fdf

Basic input variables

- 1.- General system descriptors
- 2.- Structural and geometrical variables
- 3.- Functional and solution method (Order-N/diagonalization)
- 4.- Convergence of the results
- 5.- Self-consistency
- 6.- Basis set generation related variables:

“How to test and generate basis sets”, Tu 12:00

General system descriptor: output

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

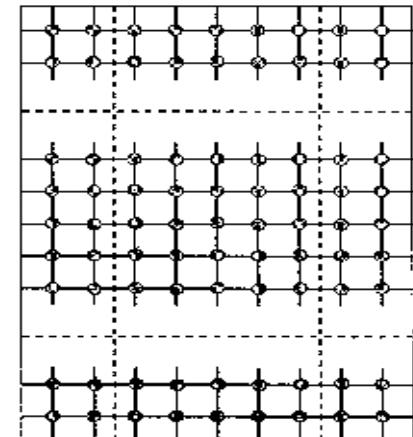
%endblock ChemicalSpeciesLabel

ALL THESE VARIABLES ARE MANDATORY

Lattice Vectors

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

Surfaces



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 on it's own 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 scaled to the lattice constant

Fractional: referred to the lattice vectors

AtomicCoordinatesFormat Fractional

AtomicCoordinatesAndAtomicSpecies:

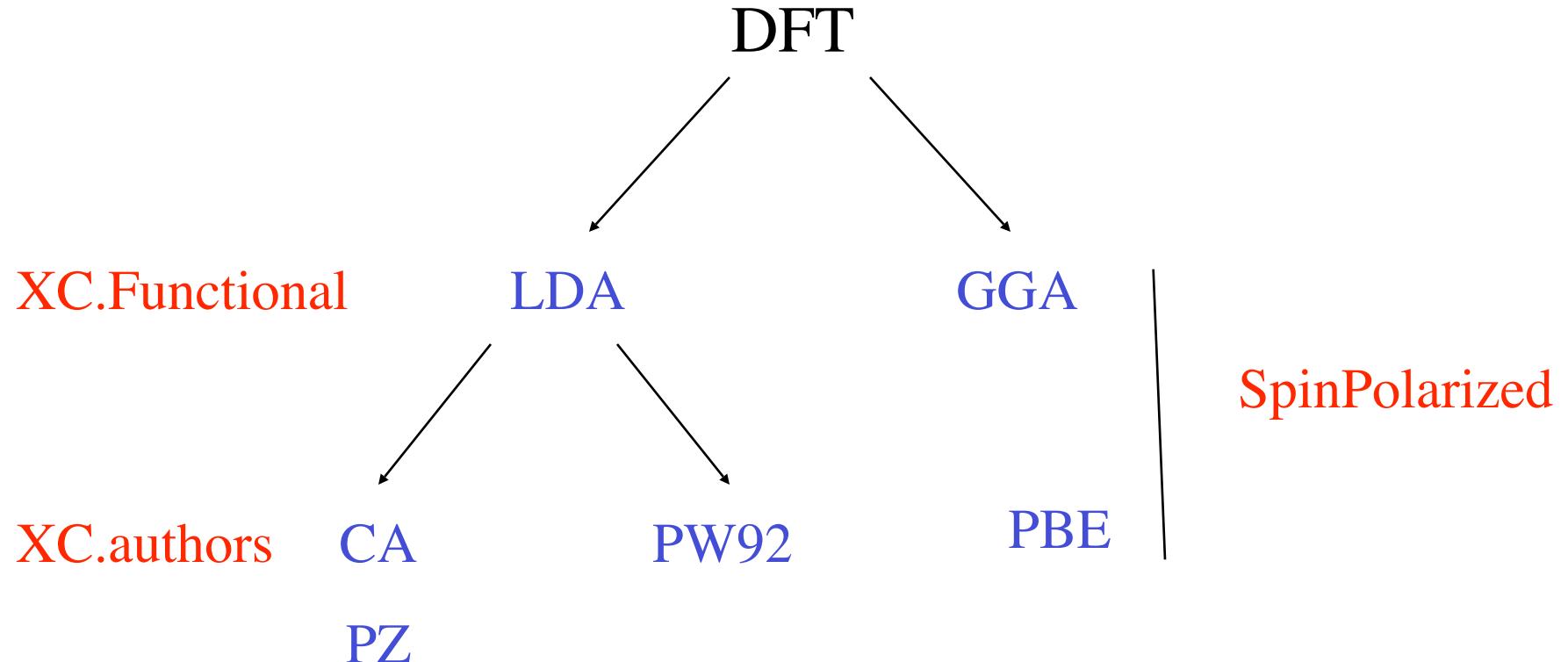
%block AtomicCoordinatesAndAtomicSpecies

 0.00 0.00 0.00 1

 0.25 0.25 0.25 1

%endblock AtomicCoordinatesAndAtomicSpecies

Functional



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

0: Start from the atomic coordinates and the unit cell

$$\{\vec{R}\}_N \{\vec{a}\}$$

1: Compute H,S (Order N):



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

2: SolutionMethod

diagon

Order-N

Execution time

P. Ordejón, “*How to run with linear-scaling solvers*”, Wed 11:10

k-sampling

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

$$\rho(\vec{r}) = \sum_i \int_{BZ} d\vec{k} n(\vec{k}) |\psi_i(\vec{k})|^2$$

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

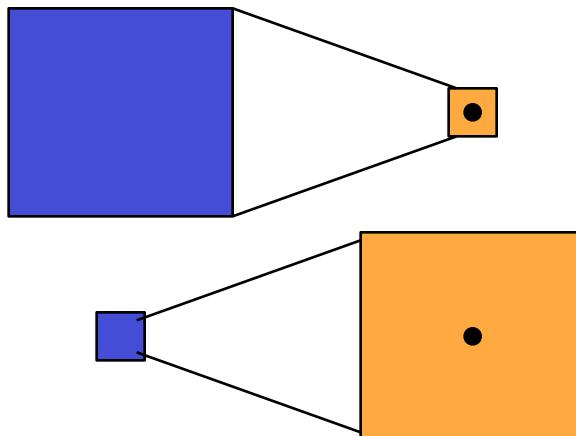
Essential for:

Small periodic systems

Metals

Magnetic systems

Real space \leftrightarrow Reciprocal space



Good description of the Bloch states at the Fermi level

Even in same insulators:

Perovskite oxides

k-sampling

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

kgrid_cutoff (1): kgrid_cutoff 10.0 Ang

kgrid_Monkhorst_Pack (2):

```
%block kgrid_Monkhorst_Pack
```

```
    4  0  0  0.5
```

```
    0  4  0  0.5
```

```
    0  0  4  0.5
```

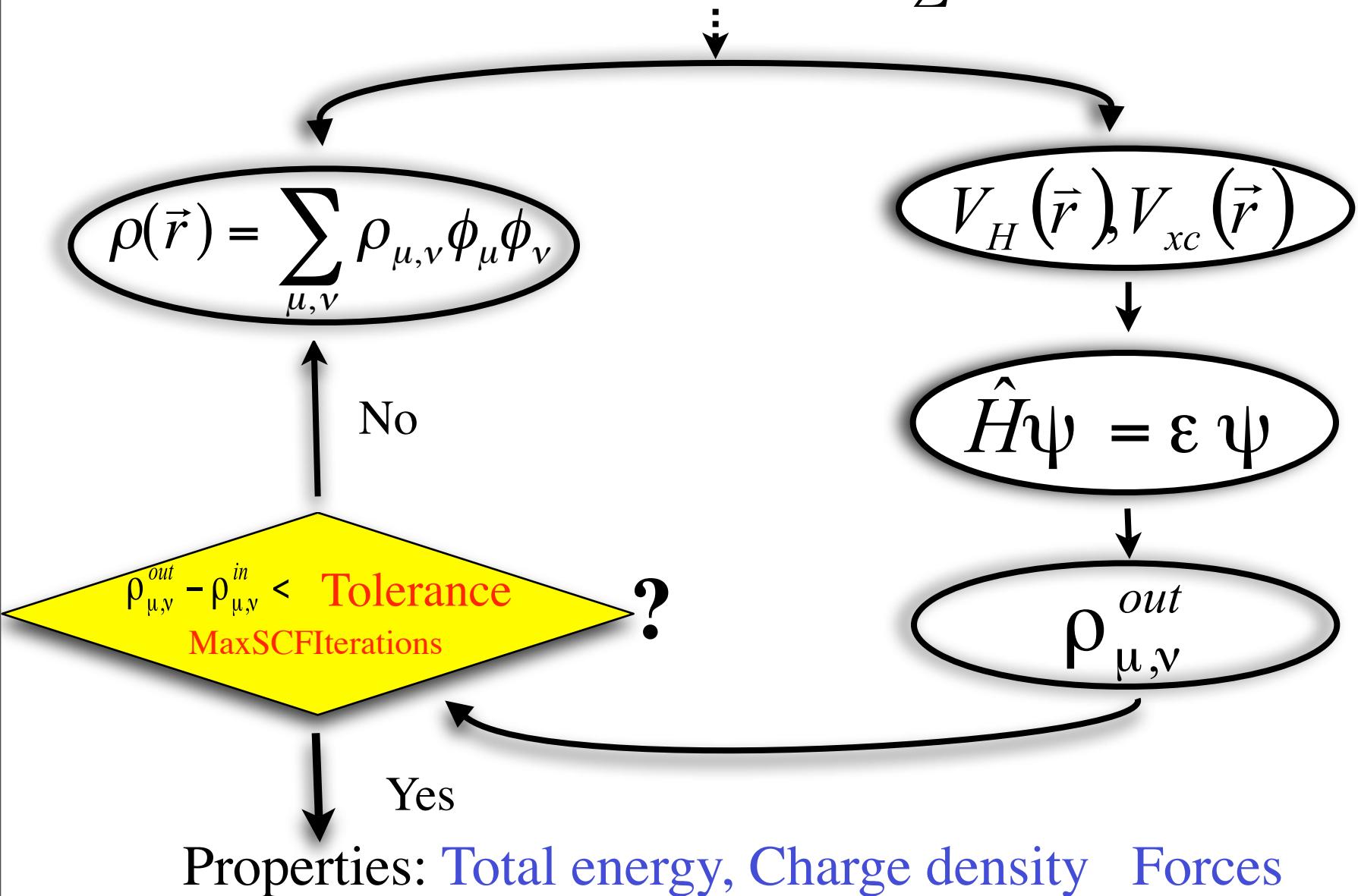
```
%endblock kgrid_Monkhorst_Pack
```

1 Moreno and Soler, PRB 45, 13891 (1992).

2 Monkhorst and Pack, PRB 13, 5188 (1997)

Selfconsistency (SCF)

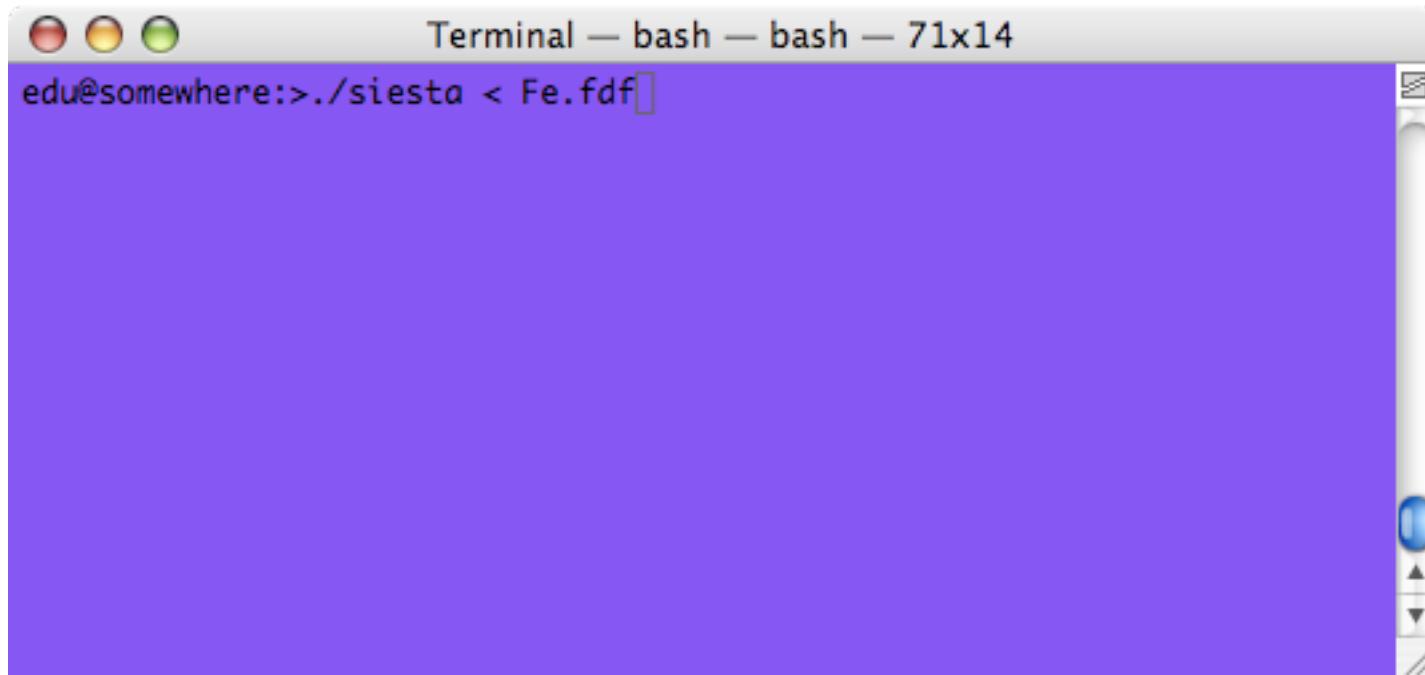
Initial guess: $\rho(\vec{r}) = \sum \rho^{atom}(\vec{r})$



How to run Siesta

To run the **serial** version, from a unix/terminal:

```
edu@somewhere:>./siesta < Fe.fdf
```



To see the output and save it at the same:

```
edu@somewhere: ./siesta < Fe.fdf | tee Fe.out
```

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: 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

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: DUscf = 0.727284
siesta: DUext = 0.000000
siesta: Exc = -112.912881
siesta: eta*DQ = 0.000000
siesta: Emadel = 0.000000
siesta: Ekinition = 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: Ekinition = 0.000000
siesta: Total = -465.837551

Output: timer (real and cpu times)

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:	CELLAC	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 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**

EXTREMLY 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 π/a

BandLines: lines were 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:

ProjectedDensityOfState

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

```
%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)
- 3D: **sies2xsf (Xcrysden)** (A. Postnikov: Friday 11:15)
- 3D: **grid2cube (Gaussian)** (P. Ordejón)
- 3D: **rho2grd (Materials Studio)** (O. Paz)

Tools (II)

-TRANSPORT PROPERTIES:

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

-PSEUDOPOTENTIAL and BASIS information:

-**PyAtom** (A. García)

-ATOMIC COORDINATES:

-**Sies2arc** (J. Gale)

- DOS, PDOS, Bands:

-**PlotUtils** (O. Paz)