

A first encounter with siestaTM

02/10/2023 - Federico Pedron

How do we run  siesta?
TM

Reminder: how to connect

https://siesta-project.org/siesta/events/SIESTA_School-2023/MN4.html



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SIESTA School 2023: MareNostrum 4

MareNostrum 4 is the largest operational supercomputer of the Barcelona Supercomputing Center. Each node contains 2 Intel Xeon Platinum processors (with 24 cores each) and 96 GB of RAM. They all run SUSE Linux 12. You should have received your BSC credentials in your institutional email account on September 30th. Your BSC account is already enabled. Please contact the organisers if you have not received this email.

Connecting to MN4

From your computer terminal, use your SSH client to connect to a MN4 login node. There are three login nodes, `m1.bsc.es`, `m2.bsc.es`, and `m3.bsc.es`. You can connect to any of them, they are used interchangeably in the examples below.

Via conventional SSH client (Mac, Linux, Windows with WSL/Cygwin)

From a terminal, run

```
$ ssh <USERNAME>@m1.bsc.es
```

where `<USERNAME>` is your BSC username (e.g., `ctn99999`). You will be prompted for the corresponding password.

If you want to enable visualization, you need to use the `-X` option:

```
$ ssh -X <USERNAME>@m1.bsc.es
```

Via PuTTY or mobaXterm

If you don't have any SSH client installed in your computer, please download and install [mobaXterm](#) (recommended) or PuTTY.

When you open the graphical interface define the server/host name (`m3.bsc.es`) and provide your username and password when prompted. For mobaXterm, there is a short generic demo video available [here](#).

For visualizations using PuTTY, please keep in mind the following:

- Under connection `→ SSH → X11`, make sure that the "Enable X11 forwarding" option is checked.
- You also need to install [xMing](#).

In mobaXterm this is enabled by default.

Reminder: how to connect

https://siesta-project.org/siesta/events/SIESTA_School-2023/MN4.html

ssh nct01YYY@mn1.bsc.es

Where **YYY** is a number unique to each of you (001, 008, 017, 035, 079).

Use also **mn2, mn3!** (only for data/password: **dt01.bsc.es**)

Then:

cd /gpfs/scratch/nct01/nct01YYY (we will run things here)

Reminder: tutorial files

Each day, you should copy the tutorial folder available at
`/gpfs/projects/nct00/nct00003/TUTORIALS/dayX`

For example:

```
cp -r /gpfs/projects/nct00/nct00003/TUTORIALS/day1 day1
```

Submitting a job

In each run folder, copy the sample run script from
`/gpfs/projects/nct00/nct00003/SCRIPTS/runmn.sh`

```
cp /gpfs/projects/nct00/nct00003/SCRIPTS/runmn.sh 1-FirstEncounter/CH4/.
```

```
cp /gpfs/projects/nct00/nct00003/SCRIPTS/runmn.sh 1-FirstEncounter/CH3/.
```

Submitting a job

Edit the run script!

```
#!/bin/bash
#SBATCH -J tutorialXX
#SBATCH -n 4
#SBATCH -t 0:30:00
#SBATCH -o %x-%j.out
#SBATCH -e %x-%j.err
#SBATCH -D .

# DO NOT CHANGE THIS LINE
source /gpfs/projects/nct00/nct00003/siestarc.sh

# EDIT THE CORRECT INPUT AND OUTPUT FILES.
srun -n 4 siesta < input.fdf > output.out
```

Submitting a job

Edit the run script!

```
#!/bin/bash
#SBATCH -J tutorialXX
#SBATCH -n 4
#SBATCH -t 0:30:00
#SBATCH -o %x-%j.out
#SBATCH -e %x-%j.err
#SBATCH -D .

# DO NOT CHANGE THIS LINE
source /gpfs/projects/nct00/nct00003/siestarc.sh

# EDIT THE CORRECT INPUT AND OUTPUT FILES.
srun -n 4 siesta < input.fdf > output.out
```


Submitting a job

Submit!

```
sbatch runmn.sh
```

To use reservations:

```
sbatch runmn.sh --reservation=SIESTA-DAY
```

```
sbatch runmn.sh --reservation=SIESTA-NIGHT
```

https://siesta-project.org/siesta/events/SIESTA_School-2023/MN4.html

A look at the inputs

What are the main ingredients?

For most basic SIESTA calculations, we need at least two inputs:

- Pseudo potential files (e.g. available in PSML format from <http://www.pseudo-dojo.org>, or a PSF created with ATOM).
- An fdf file with the input options.

What's in the FDF?

The fdf file contains all relevant input options for our simulation: geometry information, atomic species information, level of theory, basis set information, and a plethora of fine-tuning options.

Let's have a look at the first fdf for this tutorial...

What's in the FDF? System information

```
#General system specifications
SystemName      CH4 molecule
SystemLabel     ch4
NumberOfAtoms   5
NumberOfSpecies 2

%block ChemicalSpeciesLabel
 1  6  C  # Species index, atomic number, species label
 2  1  H  # Species index, atomic number, species label
%endblock ChemicalSpeciesLabel
```

All output filenames will begin with "ch4."

Total number of atoms in the simulation box.

Different "kinds" of atoms present.

Note that we have two types of inputs:
single variables, and blocks.

What's in the FDF? System geometry

```
#Unit cell for the calculation
LatticeConstant 15 Ang
%block LatticeVectors
  1.000 0.000 0.000
  0.000 1.000 0.000
  0.000 0.000 1.000
%endblock LatticeVectors

#Atomic coordinates
AtomicCoordinatesFormat Ang
%block AtomicCoordinatesAndAtomicSpecies
  0.000 0.000 0.000 1
  1.219 -0.284 -0.377 2
 -0.284 1.219 -0.377 2
 -0.140 -0.140 1.219 2
 -0.833 -0.833 -0.503 2
%endblock AtomicCoordinatesAndAtomicSpecies
```

Multiplies all lattice vectors by a constant. Note the units.

The lattice vectors themselves.

Unit for the atomic coordinates block. Can also be "fractional".

Atomic coordinates and species index (1 for C, 2 for H).

What's in the FDF? Other options

```
# Basis set definition
PAO.EnergyShift 250 meV
PAO.SplitNorm 0.15
PAO.BasisSize SZ
```

Basis Set Options

```
#Real space grid
MeshCutoff 125.0 Ry
```

Relates to the amount of points for grid-based operations.

```
# Convergence of SCF
MaxSCFIterations 50
DM.MixingWeight 0.4
DM.NumberPulay 2
```

Options for SCF acceleration.

```
# Type of solution
SolutionMethod diagon
```

Solver options.

Let's try it!

Reminders

TUTORIAL:

<https://docs.siesta-project.org/projects/siesta/en/latest/tutorials/basic/first-encounter/index.html> (last part is optional)

- 1) `ssh nct01YYY@mn2.bsc.es` ← (also mn1, mn3)
- 2) `cd /gpfs/scratch/nct01/nct01YYY`
- 3) `cp /gpfs/projects/nct00/nct00003/TUTORIALS/day1 .`
- 4) `cp /gpfs/projects/nct00/nct00003/SCRIPTS/runmn.sh .`
- 5) (move and edit runmn.sh to your run folders)
- 6) `sbatch runmn.sh`

Let's have a look at the outputs...

Installation and run info, Start Time

```
Architecture      : ----
Compiler version: GNU-11.3.0
Compiler flags    : -fallow-argument-mismatch;-O3 -march=native
PP flags          : ----
Libraries         : ----
Parallelisations: MPI
GEMM3M support
NetCDF support
NetCDF-4 support
Lua support

Runtime information:
* Directory : /home/fnpedron/siesta-docs/work-files/tutorials/basic/first-encounter/CH4
* Running on 4 nodes in parallel.
>> Start of run: 21-SEP-2023  9:39:43

*****
* WELCOME TO SIESTA *
*****
```

What are all of these files???

```
0_NORMAL_EXIT      H.ion.nc          ch4.FA
BASIS_ENTHALPY     H.ion.xml        ch4.KP
BASIS_HARRIS_ENTHALPY H.psf           ch4.ORB_INDX
C.gga.psf          INPUT_TMP.45433  ch4.STRUCT_OUT
C.ion              MESSAGES         ch4.XV
C.ion.nc           OUTVARS.yml      ch4.alloc
C.ion.xml          PARALLEL_DIST    ch4.bib
C.psf              TIMES            ch4.contrib.EPSIMG
CLOCK              ch4.BONDS        ch4.fdf
FORCE_STRESS       ch4.BONDS_FINAL  ch4.out
H.gga.psf          ch4.DM           ch4.times
H.ion              ch4.EIG          fdf.20230921T093943.534.log
```

What are all of these files???

Density Matrix Restart

Forces on atoms

Coordinate Restart

```
0_NORMAL_EXIT
BASIS_ENTHALPY
BASIS_HARRIS_ENTHALPY
C.gga.psf
C.ion
C.ion.nc
C.ion.xml
C.psf
CLOCK
FORCE_STRESS
H.gga.psf
H.ion
H.ion.nc
H.ion.xml
H.psf
INPUT_TMP.45433
MESSAGES
OUTVARS.yml
PARALLEL_DIST
TIMES
ch4.BONDS
ch4.BONDS_FINAL
ch4.DM
ch4.EIG
H.ion.nc
H.ion.xml
H.psf
INPUT_TMP.45433
MESSAGES
OUTVARS.yml
PARALLEL_DIST
TIMES
ch4.BONDS
ch4.BONDS_FINAL
ch4.DM
ch4.EIG
ch4.FA
ch4.KP
ch4.ORB_INDX
ch4.STRUCT_OUT
ch4.XV
ch4.alloc
ch4.bib
ch4.contrib.EPSIMG
ch4.fdf
ch4.out
ch4.times
fdf.20230921T093943.534.log
```

Forces and Stress

KS eigenvalues

Timing information

What are all of these files???

```
0_NORMAL_EXIT      H.ion.nc          ch4.FA
BASIS_ENTHALPY     H.ion.xml        ch4.KP
BASIS_HARRIS_ENTHALPY H.psf           ch4.ORB_INDX
C.gga.psf         INPUT_TMP.45433  ch4.STRUCT_OUT
C.ion            MESSAGES         ch4.XV
C.ion.nc         OUTVARS.yml      ch4.alloc
C.ion.xml       PARALLEL_DIST    ch4.bib
C.psf           TIMES           ch4.contrib.EPSIMG
CLOCK           ch4.BONDS       ch4.fdf
FORCE_STRESS    ch4.BONDS_FINAL ch4.out
H.gga.psf       ch4.DM          ch4.times
H.ion          ch4.EIG         fdf.20230921T093943.534.log
```

General Output file: log, out, **you** name it

Outputs

Things we have in our FDF file

```
***** Dump of input data file *****
#General system specifications
SystemName      CH4 molecule
SystemLabel     ch4
NumberOfAtoms   5
NumberOfSpecies 2
%block ChemicalSpeciesLabel
  1 6 C # Species index, atomic number, species label
  2 1 H # Species index, atomic number, species label
%endblock ChemicalSpeciesLabel
#Unit cell for the calculation
LatticeConstant 15 Ang
%block LatticeVectors
  1.000 0.000 0.000
  0.000 1.000 0.000
  0.000 0.000 1.000
%endblock LatticeVectors
#Atomic coordinates
AtomicCoordinatesFormat Ang
%block AtomicCoordinatesAndAtomicSpecies
  0.000 0.000 0.000 1
  1.219 -0.284 -0.377 2
 -0.284 1.219 -0.377 2
 -0.140 -0.140 1.219 2
 -0.833 -0.833 -0.503 2
%endblock AtomicCoordinatesAndAtomicSpecies
# Basis set definition
PAO.EnergyShift 250 meV
PAO.SplitNorm 0.15
PAO.BasisSize SZ
#Real space grid
MeshCutoff 125.0 Ry
# Convergence of SCF
MaxSCFIterations 50
DM.MixingWeight 0.4
DM.NumberPulay 2
# Type of solution
SolutionMethod diagon
***** End of input data file *****
```

Outputs

```
initatom: Reading input for the pseudopotentials and atomic orbitals -----
Species number: 1 Atomic number: 6 Label: C
Species number: 2 Atomic number: 1 Label: H

---- Processing specs for species: C
Ground state valence configuration: 2s02 2p02
Reading pseudopotential information in formatted form from:
  C.psf

---- Processing specs for species: H
Ground state valence configuration: 1s01
Reading pseudopotential information in formatted form from:
  H.psf

---- Pseudopotential check for C

Pseudized shells:
2s( 2.00) rc: 1.29
2p( 2.00) rc: 1.29
3d( 0.00) rc: 1.29
4f( 0.00) rc: 1.29
Valence configuration for ps generation: (assumed as above)

---- Pseudopotential check for H

Pseudized shells:
1s( 1.00) rc: 1.25
2p( 0.00) rc: 1.25
3d( 0.00) rc: 1.25
4f( 0.00) rc: 1.25
Valence configuration for ps generation: (assumed as above)
For C, standard SIESTA heuristics set lmxkb to 2
(one more than the basis l, including polarization orbitals).
Use PS.lmax or PS.KBprojectors blocks to override.
For H, standard SIESTA heuristics set lmxkb to 1
(one more than the basis l, including polarization orbitals).
Use PS.lmax or PS.KBprojectors blocks to override.
```

Species and pseudopotential information

Outputs

Basis set generation (next session!)

```
atom: -----
atom: SANKEY-TYPE ORBITALS:
SPLIT: Orbitals with angular momentum L= 0
SPLIT: Basis orbitals for state 2s
SPLIT: PAO cut-off radius determined from an
SPLIT: energy shift= 0.018374 Ry

  izeta = 1
          lambda = 1.000000
          rc = 4.191849
          energy = -0.983900
          kinetic = 0.912099
  potential(screened) = -1.895999
  potential(ionic) = -5.500930

SPLIT: Orbitals with angular momentum L= 1
SPLIT: Basis orbitals for state 2p
SPLIT: PAO cut-off radius determined from an
SPLIT: energy shift= 0.018374 Ry

  izeta = 1
          lambda = 1.000000
          rc = 4.993604
          energy = -0.381878
          kinetic = 2.577411
  potential(screened) = -2.959289
  potential(ionic) = -6.460511
atom: Total number of Sankey-type orbitals: 4

atm_pop: Valence configuration (for local Pseudopot. screening):
 2s( 2.00)
 2p( 2.00)
Vna: chval, zval: 4.00000 4.00000

Vna: Cut-off radius for the neutral-atom potential: 4.993604
```

Coordinates and selected options

```
coor: Atomic-coordinates input format = Cartesian coordinates
coor: (in Angstroms)

siesta: Atomic coordinates (Bohr) and species
siesta: 0.00000 0.00000 0.00000 1 1
siesta: 2.30358 -0.53668 -0.71243 2 2
siesta: -0.53668 2.30358 -0.71243 2 3
siesta: -0.26456 -0.26456 2.30358 2 4
siesta: -1.57414 -1.57414 -0.95053 2 5

siesta: System type = molecule

initatomlists: Number of atoms, orbitals, and projectors: 5 8 25

siesta: ***** Simulation parameters *****
siesta:
siesta: The following are some of the parameters of the simulation.
siesta: A complete list of the parameters used, including default values,
siesta: can be found in file out.fdf
siesta:
redata: Spin configuration = none
redata: Number of spin components = 1
redata: Time-Reversal Symmetry = T
redata: Spin spiral = F
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: Matel table size (NRTAB) = 1024
redata: Mesh Cutoff = 125.0000 Ry
redata: Net charge of the system = 0.0000 |e|
redata: Min. number of SCF Iter = 0
redata: Max. number of SCF Iter = 50
redata: SCF convergence failure will abort job =
redata: SCF mix quantity = Hamiltonian
redata: Mix DM or H after convergence = F
redata: Recompute H after scf cycle = F
redata: Mix DM in first SCF step = T
redata: Write Pulay info on disk = F
```

Outputs

```
=====
      Single-point calculation
=====

outcell: Unit cell vectors (Ang):
      15.000000   0.000000   0.000000
      0.000000   15.000000   0.000000
      0.000000   0.000000   15.000000

outcell: Cell vector modules (Ang)   :   15.000000   15.000000   15.000000
outcell: Cell angles (23,13,12) (deg):    90.0000    90.0000    90.0000
outcell: Cell volume (Ang**3)         :   3375.0000

<dSpData1D:S at geom step 0
  <sparsity:sparsity for geom step 0
    nrows_g=8 nrows=2 sparsity=.2500 nnzs=16, refcount: 7>
  <dData1D:(new from dSpData1D) n=16, refcount: 1>
refcount: 1>
new_DM -- step:      1
Initializing Density Matrix...
DM filled with atomic data:
<dSpData2D:DM initialized from atoms
  <sparsity:sparsity for geom step 0
    nrows_g=8 nrows=2 sparsity=.2500 nnzs=16, refcount: 8>
  <dData2D:DM n=16 m=1, refcount: 1>
refcount: 1>
No. of atoms with KB's overlapping orbs in proc 0. Max # of overlaps:      5      8

InitMesh: MESH = 108 x 108 x 108 = 1259712
InitMesh: Mesh cutoff (required, used) =   125.000   143.274 Ry
New grid distribution [1]: sub = 2
New grid distribution [2]: sub = 2
New grid distribution [3]: sub = 2
Setting up quadratic distribution...

stepf: Fermi-Dirac step function
```

Type of run, cell information.

Sparsity information.

Mesh information (tomorrow!)

Outputs

```
siesta: Program's energy decomposition (eV):  
siesta: Ebs      =      -86.773862  
siesta: Eions   =      383.324493  
siesta: Ena     =      115.426770  
siesta: Ekin    =      143.738590  
siesta: Enl     =     -16.728728  
siesta: Eso     =           0.000000  
siesta: Edftu   =           0.000000  
siesta: DEna    =           1.592579  
siesta: DUskf   =           0.349516  
siesta: DUext   =           0.000000  
siesta: Ex      =     -64.874822  
siesta: Ec      =     -10.703118  
siesta: Exc     =     -75.577940  
siesta: EbV     =           0.000000  
siesta: eta*DQ  =           0.000000  
siesta: Emadel  =           0.000000  
siesta: Emeta   =           0.000000  
siesta: Emolmec =           0.000000  
siesta: Ekinion =           0.000000  
siesta: Eharris =     -223.671697  
siesta: Etot    =     -214.523706  
siesta: FreeEng =     -214.523706
```

Initial, non-SCF energy
decomposition.

Outputs

```
iscf      Eharris(eV)      E_KS(eV)      FreeEng(eV)      dDmax      Ef(eV)      dHmax(eV)
scf:      1      -223.671697      -214.523706      -214.523706      1.090911      -7.083002      1.436999
timer: Routine, Calls, Time, % = IterSCF      1      0.133      29.48
scf:      2      -214.585551      -214.573147      -214.573147      0.040577      -6.647325      0.203018
scf:      3      -214.573456      -214.573477      -214.573477      0.004139      -6.585363      0.150120
scf:      4      -214.573442      -214.573493      -214.573493      0.002062      -6.424159      0.074339
scf:      5      -214.573514      -214.573506      -214.573506      0.000928      -6.476298      0.003034
scf:      6      -214.573506      -214.573506      -214.573506      0.000039      -6.474131      0.000389

SCF Convergence by DM+H criterion
max |DM_out - DM_in|      :      0.0000385344
max |H_out - H_in|      (eV) :      0.0003888059
SCF cycle converged after 6 iterations

Using DM_out to compute the final energy and forces
No. of atoms with KB's overlapping orbs in proc 0. Max # of overlaps:      5      8

siesta: E_KS(eV) =      -214.5735

siesta: E_KS - E_eggbox =      -214.5735

siesta: Atomic forces (eV/Ang):
-----
Tot      0.000066      0.000066      -0.001085
-----
Max      2.352006
Res      1.126956      sqrt( Sum f_i^2 / 3N )
-----
Max      2.352006      constrained

Stress tensor Voigt[x,y,z,yz,xz,xy] (kbar):      1.99      1.99      0.95      -0.20
(Free)E + p*V (eV/cell)      -218.0329
Target enthalpy (eV/cell)      -214.5735
```

SCF cycle information

Converged KS energy

Converged total forces and cell stress

Final energy decomposition

```
siesta: Program's energy decomposition (eV):
siesta: Ebs = -90.137390
siesta: Eions = 383.324493
siesta: Ena = 115.426770
siesta: Ekin = 141.310823
siesta: Enl = -16.669337
siesta: Eso = 0.000000
siesta: Edftu = 0.000000
siesta: DEna = 3.517376
siesta: DUskf = 0.257037
siesta: DUext = 0.000000
siesta: Ex = -64.416938
siesta: Ec = -10.674744
siesta: Exc = -75.091682
siesta: Ebv = 0.000000
siesta: eta*DQ = 0.000000
siesta: Emadel = 0.000000
siesta: Emeta = 0.000000
siesta: Emolmec = 0.000000
siesta: Ekinion = 0.000000
siesta: Eharris = -214.573506
siesta: Etot = -214.573506
siesta: FreeEng = -214.573506

siesta: Final energy (eV):
siesta: Band Struct. = -90.137390
siesta: Kinetic = 141.310823
siesta: Hartree = 282.193258
siesta: Edftu = 0.000000
siesta: Eso = 0.000000
siesta: Ext. field = 0.000000
siesta: Exch. = -64.416938
siesta: Corr. = -10.674744
siesta: Bulk bias = 0.000000
siesta: Exch.-corr. = -75.091682
siesta: Ion-electron = -697.792327
siesta: Ion-ion = 134.806422
siesta: Ekinion = 0.000000
siesta: D3 dispersion = 0.000000
siesta: Total = -214.573506
siesta: Fermi = -6.474131
```

Final forces

Final stress/pressure

Electric dipole

```
siesta: Atomic forces (eV/Ang):
siesta:   1   0.152980   0.152980  -1.053682
siesta:   2  -2.352006   0.483512   0.761553
siesta:   3   0.483512  -2.352006   0.761553
siesta:   4   0.342189   0.342189  -0.971719
siesta:   5   1.373392   1.373392   0.501211
siesta: -----
siesta:   Tot   0.000066   0.000066  -0.001085

siesta: Stress tensor (static) (eV/Ang**3):
siesta:   0.001241  -0.000019  -0.000128
siesta:  -0.000019   0.001241  -0.000128
siesta:  -0.000128  -0.000128   0.000593

siesta: Cell volume =          3375.000000 Ang**3

siesta: Pressure (static):
siesta:           Solid           Molecule  Units
siesta:          -0.00001116         0.00000003 Ry/Bohr**3
siesta:          -0.00102500         0.00000251 eV/Ang**3
siesta:          -1.64224685         0.00402704 kBar
(Free)E+ p_basis*V_orbitals =          -214.071102
(Free)Eharris+ p_basis*V_orbitals =          -214.071102

siesta: Electric dipole (a.u.) =  -0.011992  -0.011992   0.008053
siesta: Electric dipole (Debye) =  -0.030480  -0.030480   0.020469
```

Primary bibliography, and end-of-run time

```
cite: Please see "ch4.bib" for an exhaustive BiBTeX file.  
cite: Please clearly indicate Siesta version in published work:  
cite: This calculation has made use of the following articles  
cite: which are encouraged to be cited in a published work.  
      Primary SIESTA paper  
      DOI: www.doi.org/10.1088/0953-8984/14/11/302  
  
>> End of run: 21-SEP-2023 9:39:44  
Job| completed
```