A first encounter with siesta



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How do we run siesta?

Reminder: how to connect

https://siesta-project.org/siesta/events/SIESTA_School-2024/MN5.html

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https://siesta-project.org/siesta/events/SIESTA School-2024/MN5.html

ssh nct01YYY@glogin1.bsc.es

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

Use also glogin2! (only for data/password: transfer1.bsc.es)

Reminder: tutorial files

Each day, you should copy the tutorial folder available at /gpfs/projects/nct_315/TUTORIALS/dayX

For example:

cp -r /gpfs/projects/nct_315/TUTORIALS/day1 day1

In each run folder, copy the sample run script from /gpfs/projects/nct_315/SCRIPTS/runmn.sh

cp /gpfs/projects/nct_315/SCRIPTS/runmn.sh 1-FirstEncounter/.

```
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 -q gp_training
            #SBATCH -A nct_315
           #SBATCH -D .
            ## Uncomment the following line, change X by the current day of the school (1-5)
            ##SBATCH --reservation=SIESTA24-DAY1X
           # DO NOT CHANGE THIS LINE
            source /gpfs/projects/nct_315/SIESTA/siestarc.sh
            # FOTT TO ADD THE CORRECT INDUT AND OUTPUT FILES.
```

srun -n 4 siesta input.fdf output.out

```
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 -q gp_training
         #SBATCH -A nct_315
         #SBATCH -D .
         ## Uncomment the following line, change X by the current day of the school (1-5)
         ##SBATCH --reservation=SIESTA24-DAY1X
         # DO NOT CHANGE THIS LINE
         source /gpfs/projects/nct_315/SIESTA/siestarc.sh
         # EDIT TO ADD THE CORRECT INDUT AND OUTPUT FILES.
          srun -n 4 siesta 4 input.fdf > output.out
```

Submit!

sbatch runmn.sh

To use reservations (from 12:00 to 19:00 CET):

sbatch runmn.sh --reservation=SIESTA24-DAY1X (X=1,2,3,4,5)

https://siesta-project.org/siesta/events/SIESTA_School-2024/MN5.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).

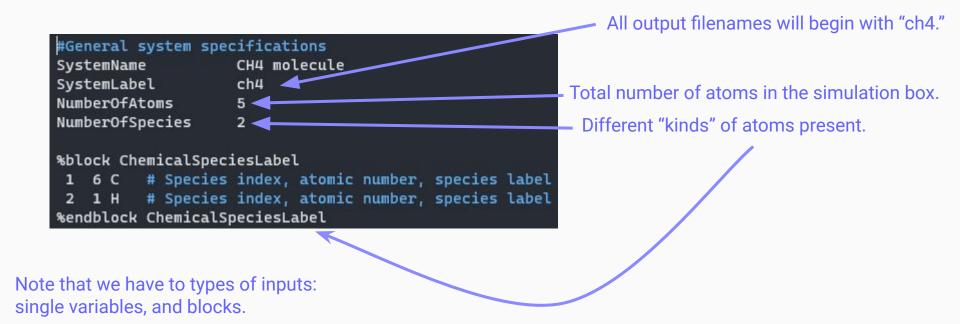
An fdf file with the input options.

What's in the FDF?

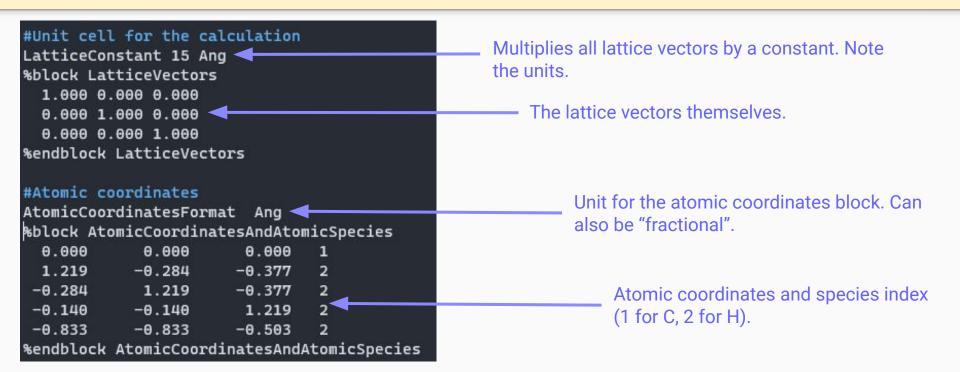
The fdf file contains all relevant input options for our simulation: geometry information, atomic species information, level of theory, basis set information, and a plethora of fine-tuning options.

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

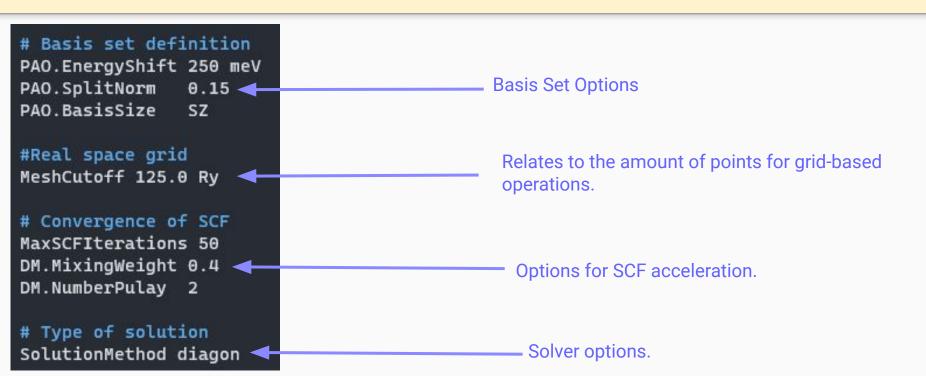
What's in the FDF? System information



What's in the FDF? System geometry



What's in the FDF? Other options



Let's try it!

Reminders

TUTORIAL:

https://docs.siesta-project.org/projects/siesta/en/latest/tutorials/basic/first-encounter/index.html (skip the last part)

- 1) $ssh \underline{nct01YYY@glogin2.bsc.es} \leftarrow (also glogin1)$
- 2) cp -r /gpfs/projects/nct_315/TUTORIALS/day1.
- 3) cd day1/01.FirstEncounter
- 4) cp /gpfs/projects/nct_315/SCRIPTS/runmn.sh.
- 5) (move and edit runmn.sh to your run folders)
- 6) sbatch runmn.sh

A First Encounter - Part 1: Running SIESTA

In this exercise we will get a first acquaintance with SIESTA by studying two simple molecules, CH4 and CH3. We will cover quite a lot of features and concepts, without worrying too much about issues of convergence.

Have you set up the local environment?

If not, do that now before proceeding.

The input files - fdf



Enter the directory 'first-encounter'

You will find an input file named ch4.fdf, along with the files c.psf and H.psf which contain the information about the pseudopotentials. The FDF (Flexible Data Format) file is the core input in

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

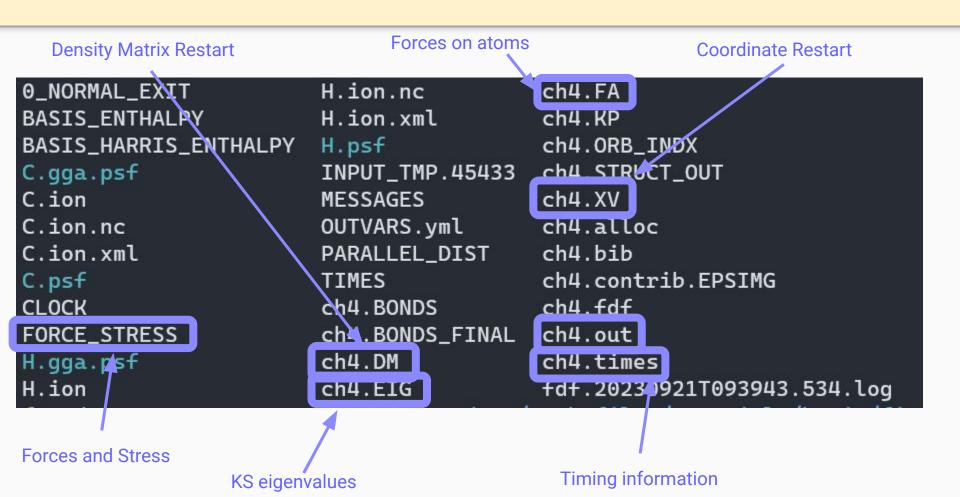
Installation and run info, Start Time

```
Architecture
Compiler version: GNU-11.3.0
Compiler flags : -fallow-argument-mismatch;-03 -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
                        H.ion.xml
BASIS_ENTHALPY
                                         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???



What are all of these files???

```
0_NORMAL_EXIT
                       H.ion.nc
                                         ch4.FA
BASIS_ENTHALPY
                       H.ion.xml
                                         ch4.KP
BASIS_HARRIS_ENTHALPY
                                         ch4.ORB_INDX
                       H.psf
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
                       ch4.BONDS
                                         ch4.fdf
CLOCK
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

Things we have in our FDF file

```
#General system specifications
SystemName
                CH4 molecule
SystemLabel
                ch4
NumberOfAtoms
NumberOfSpecies
%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
         -0.284
 1.219
                   -0.377
 -0.284
        1.219
                   -0.377 2
         -0.140
 -0.140
                   1.219
 -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
```

```
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

```
atom: SANKEY-TYPE ORBITALS:
SPLIT: Orbitals with angular momentum L= Θ
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
                            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
```

Basis set generation (next session!)

Coordinates and selected options

```
Atomic-coordinates input format =
                                              Cartesian coordinates
                                              (in Angstroms)
coor:
siesta: Atomic coordinates (Bohr) and species
siesta:
            0.00000 0.00000 0.00000 1
siesta:
            2.30358 -0.53668 -0.71243 2
siesta:
           -0.53668
                    2.30358 -0.71243 2
siesta:
           -0.26456 -0.26456 2.30358 2
                                                  4
siesta:
           -1.57414 -1.57414 -0.95053 2
siesta: System type = molecule
initatomlists: Number of atoms, orbitals, and projectors:
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
redata: Number of Atomic Species
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 lel
redata: Min. number of SCF Iter
                                                            Θ
redata: Max. number of SCF Iter
                                                           50
redata: SCF convergence failure will abort job
                                                   = Hamiltonian
redata: SCF mix quantity
redata: Mix DM or H after convergence
redata: Recompute H after scf cycle
redata: Mix DM in first SCF step
redata: Write Pulay info on disk
```

```
_____
                       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 (Ana)
                                       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</pre>
    nrows_g=8 nrows=2 sparsity=.2500 nnzs=16, refcount: 7>
  <dData1D:(new from dSpData1D) n=16, refcount: 1>
refcount: 1>
new_DM -- step:
Initializing Density Matrix...
DM filled with atomic data:
<dSpData2D:DM initialized from atoms
  <sparsity:sparsity for geom step 0</pre>
    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 overlaping orbs in proc 0. Max # of overlaps:
                                                                                8
TnitMesh: MESH = 108 \times 108 \times 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!)

```
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: DUscf
                          0.349516
siesta: DUext
                          0.000000
siesta: Ex
                        -64.874822
siesta: Ec
                        -10.703118
siesta: Exc
                        -75.577940
siesta: EbV
                          0.000000
siesta: eta*D0 =
                          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.

```
Eharris(eV)
                                                 FreeEng(eV)
                                                                           Ef(eV) dHmax(eV)
        iscf
                                    E KS(eV)
                                                                  dDmax
                 -223.671697
   scf:
                                 -214.523706
                                                             1.090911 -7.083002 1.436999
timer: Routine, Calls, Time, % = IterSCF
                                                     0.133 29.48
   scf:
                                                 -214.573147 0.040577 -6.647325 0.203018
                 -214.585551
                                 -214.573147
   scf:
                 -214.573456
                                 -214.573477
                                                 -214.573477 0.004139 -6.585363 0.150120
   scf:
                 -214.573442
                                 -214.573493
                                                 -214.573493 0.002062 -6.424159 0.074339
   scf:
                 -214.573514
                                 -214.573506
                                                 -214.573506 0.000928 -6.476298 0.003034
   scf:
                 -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 overlaping orbs in proc 0. Max # of overlaps:
siesta: E_KS(eV) =
                               -214.5735
siesta: E_KS - E_eggbox =
                               -214.5735
siesta: Atomic forces (eV/Ang):
          0.000066
   Tot
                      0.000066
          2.352006
   Max
                      sqrt( Sum f_i^2 / 3N )
          1.126956
          2.352006
                      constrained
   Max
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
                          0.00000
siesta: Edftu
                          0.000000
siesta: DEna
                          3.517376
siesta: DUscf
                          0.257037
siesta: DUext
                          0.00000
siesta: Ex
                        -64.416938
siesta: Ec
                        -10.674744
siesta: Exc
                        -75.091682
siesta: EbV
                          0.00000
siesta: eta*DQ =
                          0.00000
siesta: Emadel =
                          0.00000
siesta: Emeta =
                          0.000000
siesta: Emolmec =
                          0.00000
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:
                            282.193258
              Hartree =
siesta:
                Edftu =
                              0.00000
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 =
                              0.00000
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:
                  0.152980
                              0.152980
                                         -1.053682
siesta:
                 -2.352006
                              0.483512
                                          0.761553
siesta:
                  0.483512
                             -2.352006
                                          0.761553
siesta:
                  0.342189
                              0.342189
                                         -0.971719
siesta:
                  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:
                                        Molecule Units
                       Solid
siesta:
                                      0.00000003 Rv/Bohr**3
                 -0.00001116
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
```

What else is there?

Utilities and others

A look at the SIESTA suite:

Eig2DOS cdf2dm cdf2grid cdf2xsf cdf_diff cdf_fft cdf_fft cdf_get_cell cdf_laplacian countJobs denchar dm2cdf dmUnblock	dm_creator dm_noncol_sign_flip4 dmbs2dm dmfilter eig2bxsf eigfat2plot f2fmaster f2fslave fat fcbuild fdf2grimme fmixmd-driver	fmpdos fractional g2c_ng gen-basis getResults gnubands grid1d grid2cdf grid2cube grid2d grid2val grid_rotate	grid_supercell horizontal hs2hsx hsx2hs ioncat ionplot.sh lwf2cdf macroave mctc-convert md2axsf mixps mpi_driver	mprop ol-stm optical optical_input orbmol_proj pdosxml permute phonons phtrans pipes_parallel pipes_serial plstm	plsts protoNEB psml2psf psop pvtsp read_spin_texture readwf readwfx rho2xsf runJobs s-dftd3 sies2arc	siesta simplex sockets_parallel sockets_serial spin_texture swarm tbtrans ts2ts tscontour tshs2tshs unfold v_info	vib2xsf vibra wfs2wfsx wfsnc2wfsx wfsx2wfs xv2xsf
--	---	---	--	--	--	---	--

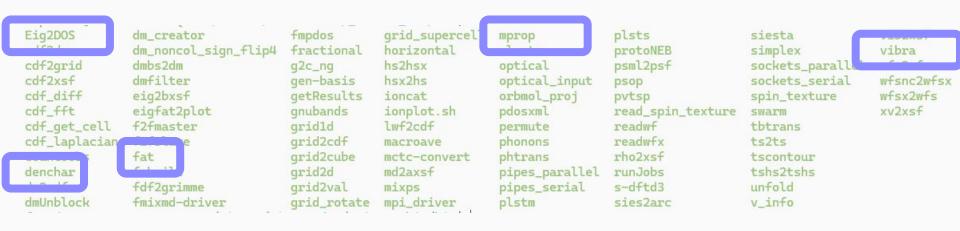
Utilities and others

A look at the SIESTA suite:

Eig2DOS cdf2dm	dm_creator dm_noncol_sign_flip4	fmpdos fractional	grid_supercell	mprop ol-stm	plsts	siesta	vib2xsf vibra
cdf2grid	dmbs2dm	g2c_ng	hs2hsx	optical	psml2psf	sockets_parallel	wfs2wfsx
cdf2xsf	dmfilter	gen-basis	hsx2hs	optical_input	psop	sockets_serial	wfsnc2wfsx
cdf_diff	eig2bxsf	getResults	ioncat	orbmol_proj	pvtsp	spin_texture	wfsx2wfs
cdf_fft	eigfat2plot	gnubands	ionplot.sh	pdosxml	read_spin_texture	swarm	xv2xsf
cdf_get_cell	f2fmaster	grid1d	lwf2cdf	permute	readwf	tbtrans	
cdf_laplacian	f2fslave	grid2cdf	macroave	phonons	readwfx	ts2ts	
countJobs	fat	grid2cube	mctc-convert	phtrans	rho2xsf	tscontour	
denchar	fcbuild	grid2d	md2axsf	pipes_parallel	runJobs	tshs2tshs	
dm2cdf	fdf2grimme	grid2val	mixps	pipes_serial	s-dftd3	unfold	
dmUnblock	fmixmd-driver	grid_rotate	mpi_driver	plstm	sies2arc	v_info	

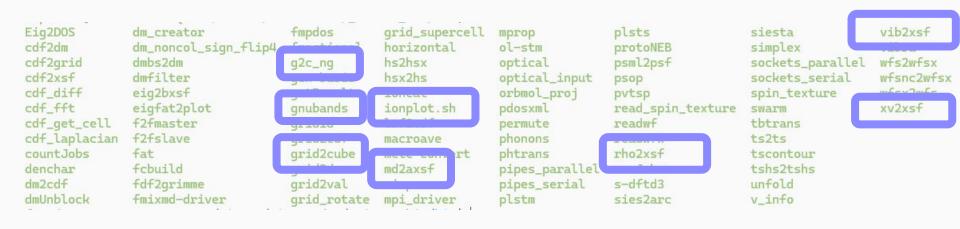
Utilities and others

Some utilities we will use later...



Utilities and others

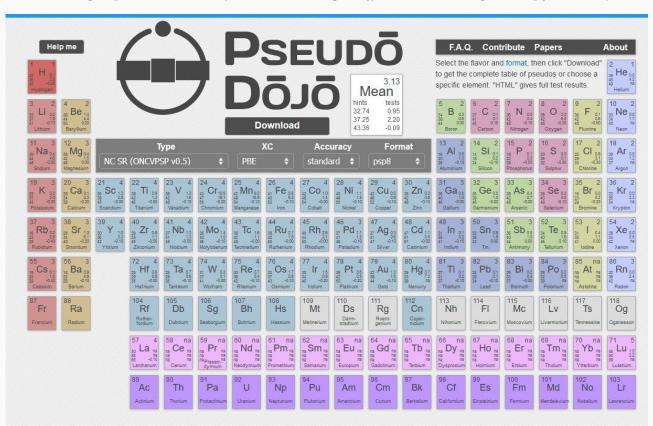
Format converters to help with visualization



Choosing a pseudo

Choosing a pseudo

Recommended way: get it from pseudo-dojo (pseudo-dojo.org) as a psml file.



Choosing a pseudo

 If you want more control, you can generate your own pseudopotentials using ATOM (https://docs.siesta-project.org/projects/atom/en/latest/tutorial/ps-generation/index.html)

 Remember to always read the literature carefully and test your pseudos accordingly.

Choosing functional

Choosing a DFT functional

SIESTA offers different families of DFT functionals:

- LDA (CA, PW91)
- GGA (BLYP, PBE, PBESol, RevPBE)
- Van der Waals functionals (DRSLL, VV)

Choosing a DFT functional

In the tutorials for day1 go to **02-LevelOfTheory**, remember to copy the run script from /gpfs/projects/nct_315/SCRIPTS/runmn.sh.

Follow the steps in the DFT Functional section of:

https://docs.siesta-project.org/projects/siesta/en/latest/tutorials/basic/first-encounter-theorylevel/index.html

A First Encounter - Part 2: Choosing your level of theory

Choosing a Pseudopotential

An important thing that determines the quality of the calculation is the choice of the pseudopotential. As mentioned in A First Encounter - Part 1: Running SIESTA, pseudopotentials in SIESTA are provided as external input files, usually in psf or psml formats.

Have you set up the local environment?

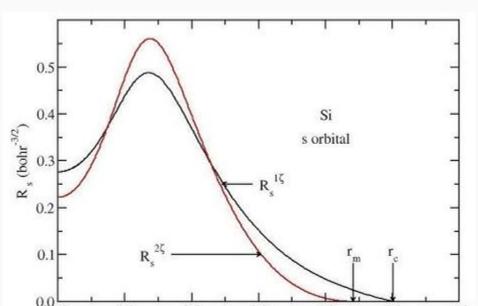
If not, do that now before proceeding.

Generating a new pseudopotential from scratch is outside the scope of this tutorial, but more information can be seen in Pseudopotentials. For most cases, you can instead download pre-

A peek into basis sets

SIESTA's basis sets are at the core of its efficiency.

For now, suffice to say that SIESTA has atomic-centered basis functions, that become zero after a certain cut-off radius.



A peek into basis sets

For now, we will only concern ourselves with:

- Exploring the basis set cardinality (SZ, SZP, DZP), i.e. the amount of basis functions per atom. In principle, more functions imply a better quality, but also an increase in computational costs.
- Playing with the energy shift, which essentially modifies the cut-off radius of the basis set. The lower the energy shift, the larger the cut-off radius of the orbitals.

We will cover this more in-depth in the upcoming sessions!

First tests with water (geometry optimization)

Follow the steps in the Playing with CH4 section of:

https://docs.siesta-project.org/projects/siesta/en/latest/tutorials/basic/first-encount er-theorylevel/index.html

Take note of how the total energy (from output), bond lengths (ch4.BONDS_FINAL file), and total time (from ch4.times) change in these cases:

1) When changing the basis set between **SZ, SZP, and DZP**. Use an energy shift of **250 meV**.

PAO.BasisSize DZP

2) For **DZP**, changing the energy shift between **5 meV**, **100 meV**, and **500 meV**.

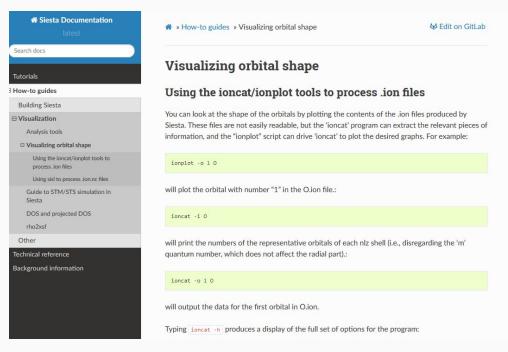
PAO.EnergyShift 0.01 Ry

TIP: Visualizing orbital shapes

How-to -> Visualization -> Visualizing Orbital Shapes

https://docs.siesta-project.org/projects/siesta/en/latest/how-to/visualization/

orbitals.html



Questions?