

OpenMX Tutorial

- First trial
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- OpenMX Viewer

Taisuke Ozaki (ISSP, Univ. of Tokyo)

First trial

➤ Login

For the tutorial, we will use DeltaServer, please login the machine.

➤ A trial calculation

As a first step, please try to perform a SCF calculation for a methane molecule.

DeltaServer

Intel(R) Xeon(R) CPU E5520 @ 2.27GHz

192 cores (4 cores × 48 processors) are available in total.

Login

```
$ ssh -YC userid@10.105.17.42
```

**As for your account name and password,
please look at the word file you received.**

How to compile OpenMX

- (1) Go to <http://www.openmx-square.org/download.html>
- (2) Download `openmx3.8.tar.gz` and the patch
- (3) Follow the instruction at http://www.openmx-square.org/tech_notes/OpenMX-Compile.pdf

See also the page 11 in the manual.

Environment setting

Once you login DeltaServer, please do first the following:

```
$ make work
```

```
$ cp -rf /home/soft/openmx3.8/work/* ./work/
```

```
$ cd work
```

```
$ rm -rf *_rst
```

```
$ rm *.cube *.md *.md2 *.ene *.cif *.out *.xyz
```

Then, modify a line in 'job.pbs' as

```
$mpirun -machinefile $PBS_NODEFILE -np $ncpu $openmx Methane.dat -nt 2 > met.std
```

And, add a line in 'Methane.dat' as

```
DATA.PATH /home/soft/openmx3.8/DFT_DATA13
```

A test calculation: Methane.dat

As a first trial calculation, let's try to calculate a methane molecule by

```
$ qsub job.pbs
```

After finishing the calculations, you will obtain 11 files and 1 directory.

met.std	standard output of the SCF calculation
met.out	input file and standard output
met.xyz	final geometrical structure
met.ene	values computed at every MD step
met.md	geometrical structures at every MD step
met.md2	geometrical structure of the final MD step
met.cif	cif file of the initial structure for Material Studio
met.tden.cube	total electron density in the Gaussian cube format
met.v0.cube	Kohn-Sham potential in the Gaussian cube format
met.vhart.cube	Hartree potential in the Gaussian cube format
met.dden.cube	difference electron density measured from atomic density
met_rst/	directory storing restart files

See the 17 p in
the manual.

Methane.dat

```
xterm
[ozaki@vtppcc01 work]$ more Methane.dat
#
# SCF calculation of a methane molecule by the LDA
# and the cluster method
#
#
# File Name
#
System,CurrentDirectory      ./      # default=./
System,Name                  met
level.of.stdout              1      # default=1 (1-3)
level.of.fileout             1      # default=1 (0-2)
#
# Definition of Atomic Species
#
Species.Number               2
<Definition.of.Atomic.Species
H   H4.0-s1                  H_TM
C   C4.5-s1p1                C_TM_PCC
Definition.of.Atomic.Species>
#
# Atoms
#
Atoms.Number                 5
Atoms.SpeciesAndCoordinates.Unit Ang # Ang|AU
<Atoms.SpeciesAndCoordinates
1  C   0.000000   0.000000   0.000000   2.0  2.0
2  H   -0.889981  -0.629312   0.000000   0.5  0.5
3  H    0.000000   0.629312  -0.889981   0.5  0.5
4  H    0.000000   0.629312   0.889981   0.5  0.5
5  H    0.889981  -0.629312   0.000000   0.5  0.5
Atoms.SpeciesAndCoordinates>
```

- (1) Value **behind** keyword
- (2) The order is **arbitrarily**.
- (3) Put # to the head of line for comment

Selected keywords:

The name of system

The number of atoms

Definition of species

Atomic coordinates

met.out

SCF history

```
*****  
*****  
SCF history at MD= 1  
*****  
*****
```

SCF=	1	NormRD=	1.000000000000	Uele=	-3.523143659974
SCF=	2	NormRD=	0.567253699744	Uele=	-4.405605131921
SCF=	3	NormRD=	0.103433490729	Uele=	-3.982266241934
SCF=	4	NormRD=	0.024234990593	Uele=	-3.906896836134
SCF=	5	NormRD=	0.011006215697	Uele=	-3.893084558820
SCF=	6	NormRD=	0.006494145332	Uele=	-3.890357113476
SCF=	7	NormRD=	0.002722267527	Uele=	-3.891669816209
SCF=	8	NormRD=	0.000000672350	Uele=	-3.889285164733
SCF=	9	NormRD=	0.000000402419	Uele=	-3.889285102456
SCF=	10	NormRD=	0.000000346348	Uele=	-3.889285101128
SCF=	11	NormRD=	0.000000515395	Uele=	-3.889285101063

KS eigenvalues

```
*****  
*****  
Eigenvalues (Hartree) for SCF KS-eq.  
*****  
*****
```

Chemical Potential (Hartree) = 0.000000000000
Number of States = 8.000000000000
HOMO = 4
Eigenvalues

	Up-spin	Down-spin
1	-0.69897190537228	-0.69897190537228
2	-0.41522646150979	-0.41522646150979
3	-0.41522645534084	-0.41522645534084
4	-0.41521772830844	-0.41521772830844
5	0.21218282298348	0.21218282298348
6	0.21218282358344	0.21218282358344
7	0.21227055734372	0.21227055734372
8	0.24742493684297	0.24742493684297

Total energy

```
*****  
Total energy (Hartree) at MD = 1  
*****
```

Uele.	-3.889285101063
Ukin.	5.533754016241
UHO.	-14.855520072374
UH1.	0.041395625260
Una.	-5.040583803800
Unl.	-0.134640939010
Uxc0.	-1.564720823137
Uxc1.	-1.564720823137
Ucore.	9.551521413583
Uhub.	0.000000000000
Ucs.	0.000000000000
Uzs.	0.000000000000
Uzo.	0.000000000000
Uef.	0.000000000000
UvdW	0.000000000000
Utot.	-8.033515406373

Mulliken population

```
*****  
*****  
Mulliken populations  
*****  
*****
```

Total spin S = 0.000000000000

	Up spin	Down spin	Sum	Diff	
1	C	2.509755704	2.509755704	5.019511408	0.000000000
2	H	0.372561098	0.372561098	0.745122197	0.000000000
3	H	0.372561019	0.372561019	0.745122038	0.000000000
4	H	0.372561127	0.372561127	0.745122254	0.000000000
5	H	0.372561051	0.372561051	0.745122102	0.000000000

Sum of MulP: up = 4.00000 down = 4.00000
total= 8.00000 ideal(neutral)= 8.00000

Specification of the directory storing *.pao and *.vps

The directory *DFT_DATA* can be specified by the keyword in your input file as

```
DATA.PATH    /home/soft/openmx3.8/DFT_DATA13
```


Specification of PAO and VPS

PAO and VPS are specified by the following keyword:

```
<Definition.of.Atomic.Species  
  O   O7.0-s2p2d1      O_PBE13  
  H   H7.0-s2p1       H_PBE13  
>Definition.of.Atomic.Species>
```

- O7.0 means O7.0.pao.
- -s2p2d1 means 2, 2, and 1 radial functions are allocated to s-, p-, and d-orbitals.
- In this case, for oxygen atom, $2 \times 1 + 2 \times 3 + 1 \times 5 = 13$ basis functions are allocated.
- O_PBE13 means O_PBE13.vps.

The path for O7.0.pao and O_PBE13.vps is specified by

```
DATA.PATH /home/soft/openmx3.8/DFT_DATA13
```

Default value is ‘../DFT_DATA13’.

How to choose basis functions: H₂O case

By clicking H7.0.pao and O7.0.pao in the database(2013), you may find the following
http://www.jaist.ac.jp/~t-ozaki/vps_pao2013/H/index.html http://www.jaist.ac.jp/~t-ozaki/vps_pao2013/O/index.html

```
*****  
Eigen values(Hartree) of pseudo atomic orbitals  
*****
```

H7.0.pao

```
Eigenvalues  
Lmax= 3 Mul=15  
| mu 0 0 -0.23595211038442  
| mu 0 1 0.14109389991827  
| mu 0 2 0.61751730037441  
| mu 0 3 1.31890671598573  
| mu 0 4 2.24052765608302  
| mu 0 5 3.37954791544661  
| mu 0 6 4.73488369825610  
| mu 0 7 6.30608874470710  
| mu 0 8 8.09282718517299  
| mu 0 9 10.09464035732420  
| mu 0 10 12.31085267019158  
| mu 0 11 14.74057314485273  
| mu 0 12 17.38277845742691  
| mu 0 13 20.23645090753857  
| mu 0 14 23.30073926597344  
| mu 1 0 0.10914684890465  
| mu 1 1 0.47776040452236  
| mu 1 2 1.06988680483686  
| mu 1 3 1.88261331124981  
| mu 1 4 2.91175885838084  
| mu 1 5 4.15601184789448  
| mu 1 6 5.61454131060210  
| mu 1 7 7.28681796296307  
| mu 1 8 9.17254361476158  
| mu 1 9 11.27156766390586  
| mu 1 10 13.58381334569800  
| mu 1 11 16.10921637499960  
| mu 1 12 18.84767560575107  
| mu 1 13 21.79902110024685  
| mu 1 14 24.96300480798286  
| mu 2 0 0.27851528500170  
| mu 2 1 0.76970400958463  
| mu 2 2 1.47576814497054  
| mu 2 3 2.39881523735167  
| mu 2 4 3.537132246447754
```

3

6

5

```
*****  
Eigen values(Hartree) of pseudo atomic orbitals  
*****
```

O7.0.pao

```
Eigenvalues  
Lmax= 3 Mul=15  
| mu 0 0 -0.87913976280231  
| mu 0 1 0.06809061901229  
| mu 0 2 0.52709275865941  
| mu 0 3 1.24995140722317  
| mu 0 4 2.21829552402723  
| mu 0 5 3.41945468859267  
| mu 0 6 4.84509607059749  
| mu 0 7 6.48825090142865  
| mu 0 8 8.34207134316001  
| mu 0 9 10.39973244132192  
| mu 0 10 12.65513764955926  
| mu 0 11 15.10428688136984  
| mu 0 12 17.74677530362947  
| mu 0 13 20.58633940582683  
| mu 0 14 23.62957122674031  
| mu 1 0 -0.33075182895384  
| mu 1 1 0.16378499567753  
| mu 1 2 0.64129274864838  
| mu 1 3 1.35995471521821  
| mu 1 4 2.31377697411480  
| mu 1 5 3.50052618379026  
| mu 1 6 4.91841590421346  
| mu 1 7 6.56499793348396  
| mu 1 8 8.43634105410091  
| mu 1 9 10.52733540479795  
| mu 1 10 12.83276989780839  
| mu 1 11 15.34880171573570  
| mu 1 12 18.07414632417004  
| mu 1 13 21.01013110934429  
| mu 1 14 24.15933607166566  
| mu 2 0 0.26162948257116  
| mu 2 1 0.70705247436937  
| mu 2 2 1.34714706672243  
| mu 2 3 2.19799459356269  
| mu 2 4 3.26511989658328
```

1

4

2

7

8

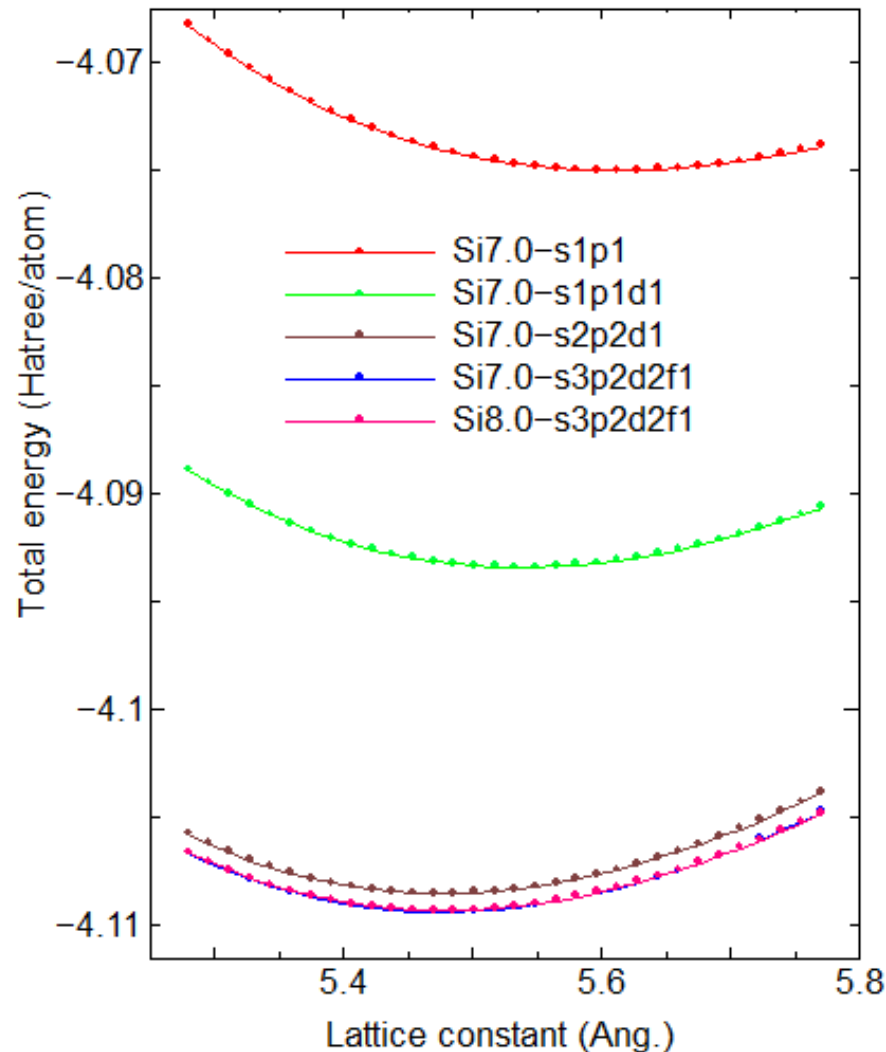
Choosing states with lower eigenvalues leads to H7.0-s2p1 and O7.0-s2p2d1.

How to choose basis functions: Si case(1)

Si7.0.pao

Eigenvalues		
Lmax=	Mul=	
0	0	-0.59320968623145
0	1	0.01654247265872
0	2	0.57915688461708
0	3	1.39915528592998
0	4	2.43106703909053
0	5	3.63498884739274
0	6	4.99885012002783
0	7	6.55485759476897
0	8	8.33541526908920
0	9	10.34055461939939
0	10	12.55818051232040
0	11	14.98369777887096
0	12	17.62014479571136
0	13	20.47011746372617
0	14	23.53245309073866
1	0	-0.31460013133101
1	1	0.12937640798489
1	2	0.71637380083701
1	3	1.54488697995006
1	4	2.59211686526084
1	5	3.84687324239366
1	6	5.31246180826158
1	7	6.99509799702661
1	8	8.89454075291723
1	9	11.00602102880752
1	10	13.32692005826443
1	11	15.85857455673626
1	12	18.60272353142246
1	13	21.55894843635971
1	14	24.72602323954447
2	0	0.01886411574821
2	1	0.35893514996065
2	2	0.94629918754692
2	3	1.76201765644983
2	4	2.81418723624388
2	5	4.10656012645961
2	6	5.63661971875011
2	7	7.39522693820483
2	8	9.37222098331867
2	9	11.56227937764802
2	10	13.96620568880690
2	11	16.58651812641581
2	12	19.42296574188659
2	13	22.47308081363278
2	14	25.73538272482773
3	0	0.28356769151846
3	1	0.79082836114569
3	2	1.52992065308726
3	3	2.52537261496132

Orbitals with lower eigenvalues in Si7.0.pao are taken into account step by step as the quality of basis set is improved.

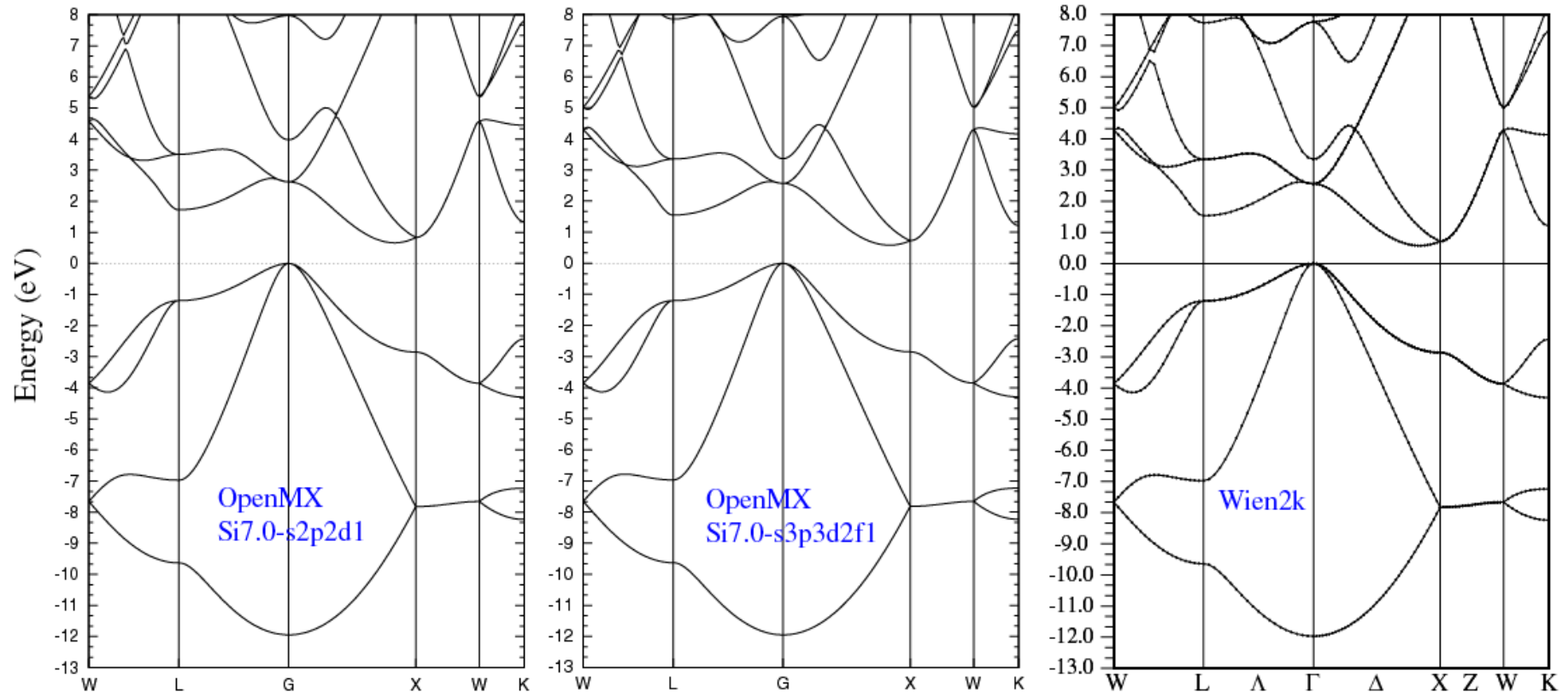


Si7.0-s2p2d1 is enough to discuss structural properties.

By comparing Si7.0-s3p2d2f1 with Si8.0-s3p2d2f1, it turns out that convergence is achieved at the cutoff of 7.0(a.u.).

How to choose basis functions: Si case(2)

With respect to band structure, one can confirm that Si7.0-s2p2d1 provides a nearly convergent result.



While the convergent result is achieved by use of Si7.0-s3p2d2f1(Si7.0-s3p3d2f1), Si7.0-s2p2d1 is a balanced basis functions compromising accuracy and efficiency to perform a vast range of materials exploration.

Choice of cutoff energy

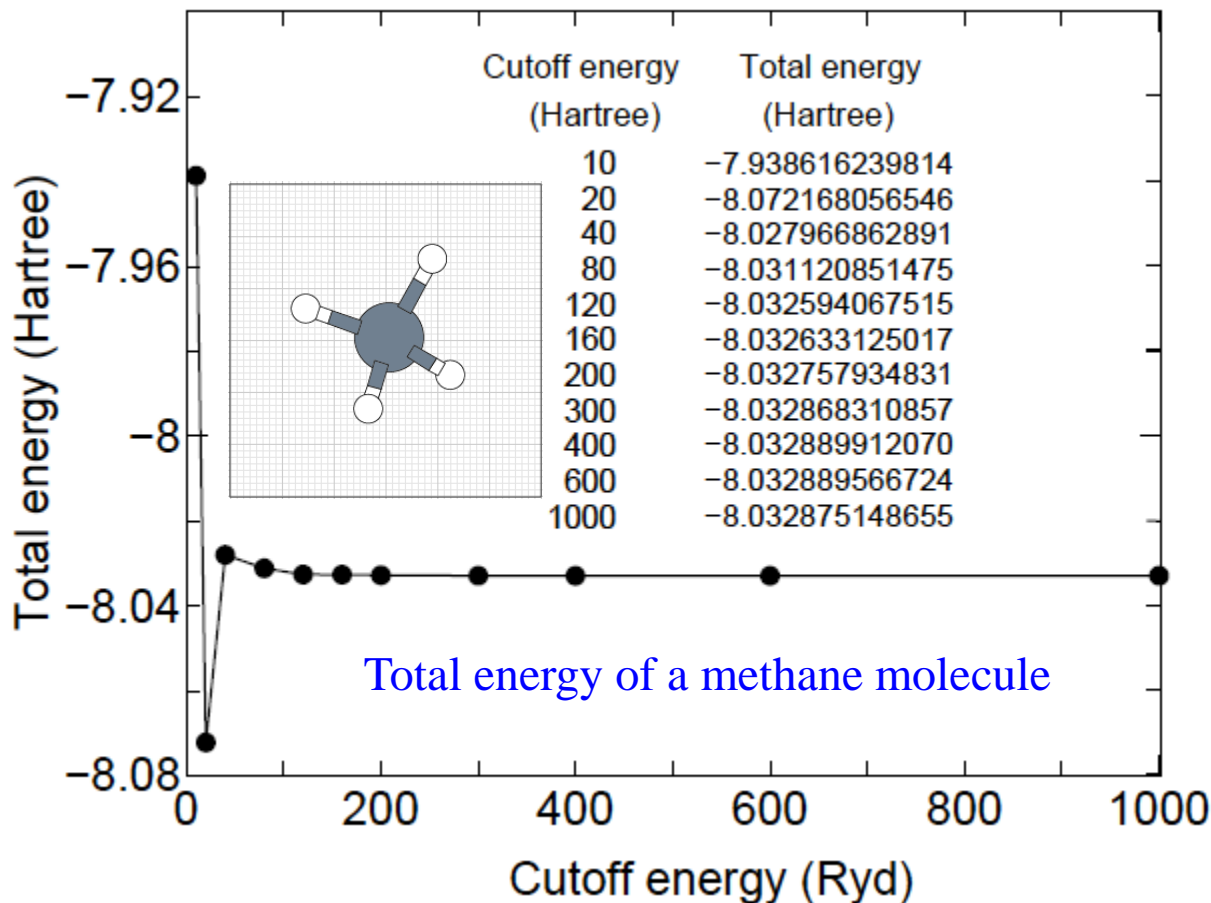
`scf.energycutoff` 200 # default=150 Ryd

The FFT grid is used to discretize real space and calculate $E_{\delta ee}$, E_{xc} , and can be specified by `scf.energycutoff`.

In most cases, 200 Ryd is enough to get convergence.

However, large cutoff energy (300-400 Ryd) has been used in cases such as the use of pseudopotentials with deep semi-core states.

Memory requirement
 $O(E^{3/2})$



Mixing methods

Available mixing methods:

Simple mixing (Simple)

Residual minimization method in the direct inversion iterative subspace (RMM-DIIS)

Guaranteed reduction Pulay method (GR-Pulay)

Kerker mixing (Kerker)

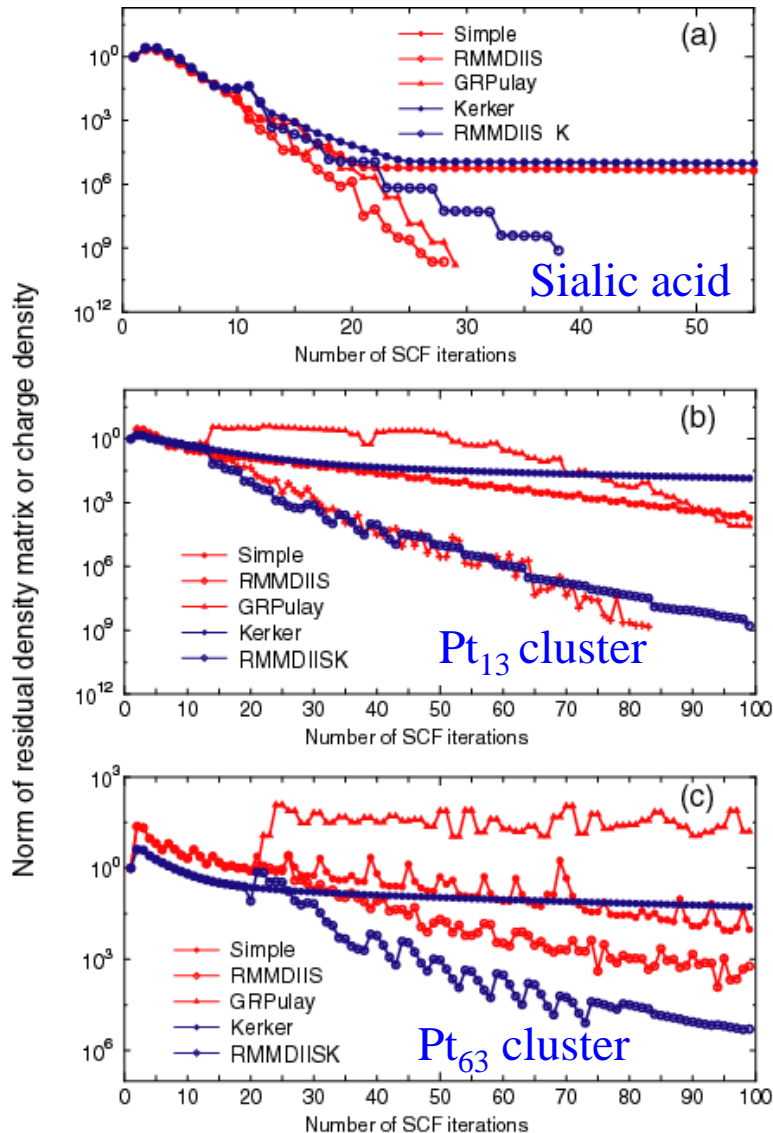
RMM-DIIS with Kerker metric (RMM-DIISK)

RMM-DIIS for Hamiltonian (RMM-DIISH)

Recommendation:

RMM-DIISK or RMM-DIISH

See also the page 56 in the manual.



Visualization

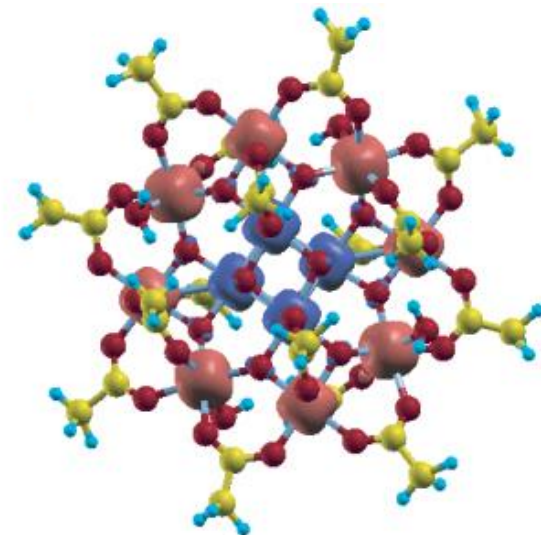
- Cube files such as *.tden.cube, *.sden.cube, *.dden.cube can be visualized by many software such as

XCrySDen
VESTA

- Also *.md file is stored in xyz format which can be visualized by XCrySDen and OpenMX Viewer.

On phi, XCrSDen is available.

- Data on DOS and band dispersion can be visualized by gnuplot.



On the manual

Please download the manual at

http://www.openmx-square.org/openmx_man3.8/openmx3.8.pdf

The manual is self-contained, the most of calculations explained in the manual are traceable by using the input file stored in the directory ‘work’.

Please try to perform those calculations one by one depending on your interests.

Recommended trials

1. Geometry optimization

Perform a geometry optimization using 'Methane2.dat'.
See the page 65 in the manual.

2. Density of states

Calculate DOS using 'Cdia.dat'.
See the page 79 in the manual.

All the input files can be
found in the directory 'work'.

3. Wannier functions

Calculate Wannier functions for Si bulk using 'work/wf_example/Si.dat', and perform the band interpolation. See the page 159 in the manual.

4. Reaction barrier by the nudged elastic band (NEB) method

Calculate a reaction barrier using 'C2H4_NEB.dat'.
See the page 182 in the manual.

5. Transmission of a carbon chain

Calculate an electric transmission of a carbon chain using 'Lead-Chain.dat', 'NEGF-Chain.dat'. See the page 136 in the manual.

6. Spin-orbit coupling

Calculate a band structure by taking account of SOC using 'GaAs.dat'.
See the page 117 in the manual.

Geometry optimization (Methane2.dat)

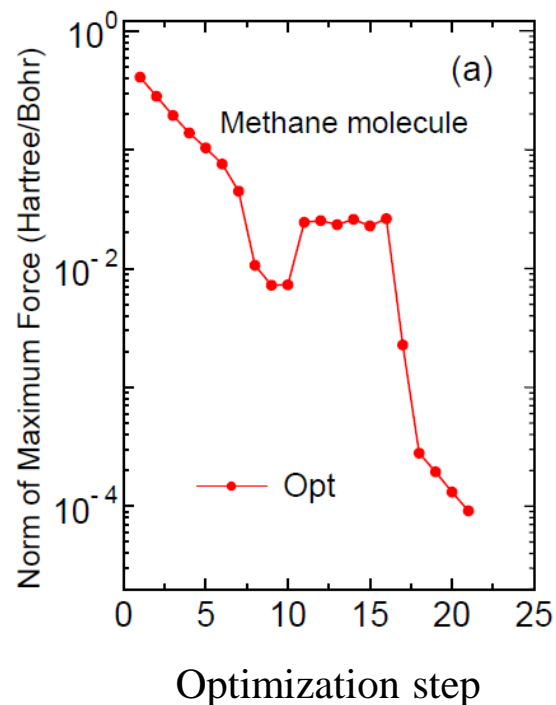
The x-coordinate of a methane molecule is displaced by 0.3 Å, and perform the geometry optimization.

```
<Atoms.SpeciesAndCoordinates
  1  C      0.300000  0.000000  0.000000  2.0  2.0
  2  H     -0.889981 -0.629312  0.000000  0.5  0.5
  3  H      0.000000  0.629312 -0.889981  0.5  0.5
  4  H      0.000000  0.629312  0.889981  0.5  0.5
  5  H      0.889981 -0.629312  0.000000  0.5  0.5
Atoms.SpeciesAndCoordinates>
```

Using the steepest decent (SD) method, the norm of maximum force converges as shown in the left figure:

met2.md can be visualized by OpenMX Viewer.

See also the page 65 in the manual.



Density of states (Cdia.dat)

Density of states (DOS) for Kohn-Sham eigenvalues can be calculated by the following two steps:

(1) SCF calculation

Perform the SCF calculation by 'Cdia.dat'.

(2) DOS calculation

(2.a) In openmx/source, compile DosMain

```
$ make DosMain
```

(2.b) mv work, and perform the DOS calculation using 'DosMain'.

```
$ ./DosMain cdia.Dos.Val cdia.Dos.Vec
```

See also the page 79 in the manual.

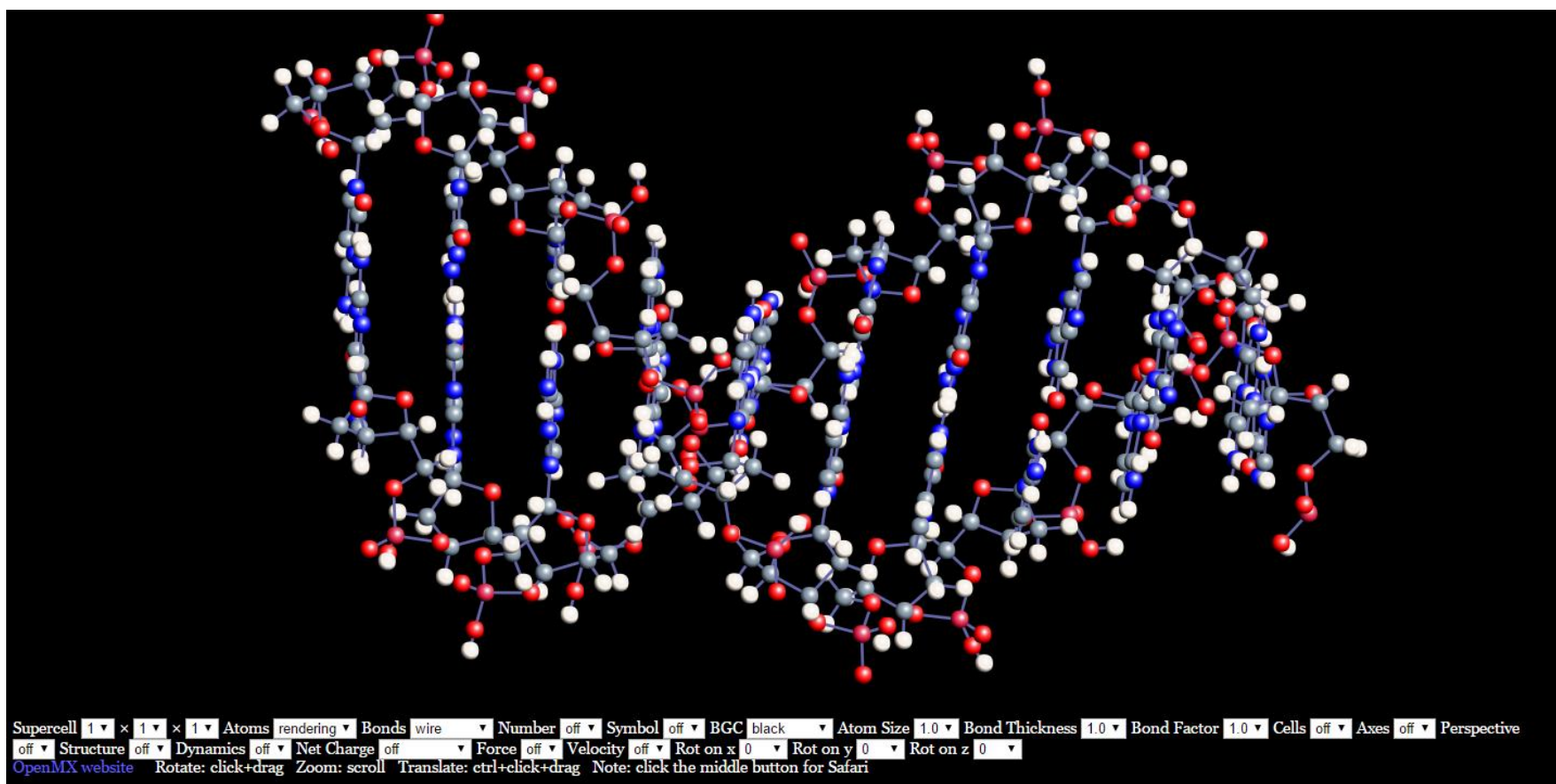
On OpenMX Viewer

<http://www.openmx-square.org/viewer/>

- 6611 lines written by html5, css3, js, and canvas2d
- dat, xyz, and md files are readable
- Rotation, zoom, and translation
- Easy change of atom size, bond thickness, and bond factor
- Atom index and atom symbol
- Supercell representation
- Structure analysis
- Analysis of geometry optimization
- Charge analysis
- Google Chrome recommended due to its performance

Easy check of a dat file

A dat file can be easily checked by drag and drop to any browser.



Analysis of structure optimization

A md file can be easily analyzed by drag and drop to any browser.

- Trace of optimization process
- Analysis of structure
- Check residual forces

Analysis of Structure

	x (Ang.)	y (Ang.)	z (Ang.)
A=N 33 (33,0,0,0)	+4.3650	+2.7331	+0.8831
B=La 6 (6,0,0,0)	+5.1220	+5.1220	+0.0000
C=N 32 (32,0,0,0)	+2.7331	+5.8790	+0.8831
D=			

Length: AB=2.6570 BC=2.6570 CD=
Angle: ABC=83.66 BCD=
Dihedral Angle: ABCD=
Length in Ang. Angle in Deg. Reset

Structural Change

time= 0.000 (fs) Energy= -530.83562 (Hartree)
Frame 1/21

First << < Stop > >> Last

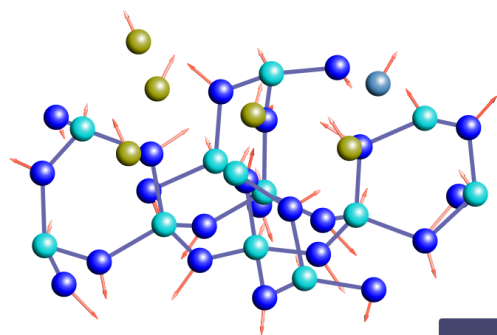
Make animation Every 1 frames
Delayed time (ms) 0

Start End File format not supp

Supercell 1 x 1 x 1 Atoms rendering Bonds wire Number off Symbol off BGC black Atom Size 1.0 Bond Thickness 1.0 Bond Factor 1.0 Cells off Axes off Perspective
off Structure on Dynamics on Net Charge off Force off Velocity off Rot on x 0 Rot on y 0 Rot on z 0
OpenMX website Rotate: click+drag Zoom: scroll Translate: ctrl+click+drag Note: click the middle button for Safari

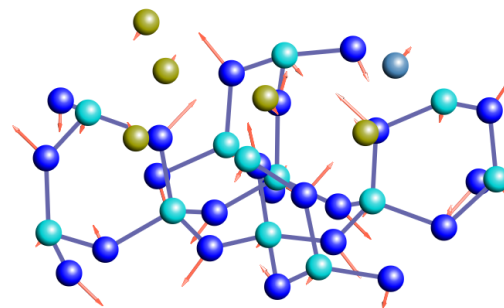
Analysis of structure optimization

Residual forces can be visualized by vectors.



1 step

Structural Change
time= 0.000 (fs) Energy= -530.83562 (Hartree)
Frame 1/21
First << < Stop > >> Last
Make animation Every 1 frames
Delayed time (ms) 0
Start End File format not supp

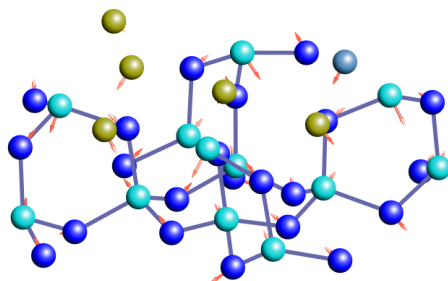


10 step

Structural Change
time= 9.000 (fs) Energy= -530.81794 (Hartree)
Frame 10/21
First << < Stop > >> Last
Make animation Every 1 frames
Delayed time (ms) 0
Start End File format not supp

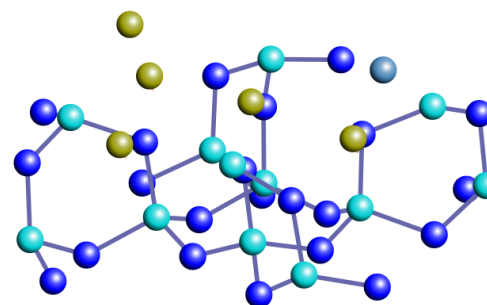
supercell 1 x 1 x 1 Atoms rendering Bonds wire Number off Symbol off BGC white Atom Size 1.0 Bond Thickness 1.0 Bond factor 1.0 Cells off Axes off Perspective off Structure off Dynamics on Net Charge off Force on Velocity off Rot on x 0 Rot on y 0 Rot on z 0
OpenMX website Rotate: click+drag Zoom: scroll Translate: ctrl+click+drag Note: click the middle button for Safari

supercell 1 x 1 x 1 Atoms rendering Bonds wire Number off Symbol off BGC white Atom Size 1.0 Bond Thickness 1.0 Bond factor 1.0 Cells off Axes off Perspective off Structure off Dynamics on Net Charge off Force on Velocity off Rot on x 0 Rot on y 0 Rot on z 0
OpenMX website Rotate: click+drag Zoom: scroll Translate: ctrl+click+drag Note: click the middle button for Safari



14 step

Structural Change
time= 13.000 (fs) Energy= -530.80467 (Hartree)
Frame 14/21
First << < Stop > >> Last
Make animation Every 1 frames
Delayed time (ms) 0
Start End File format not supp



21 step

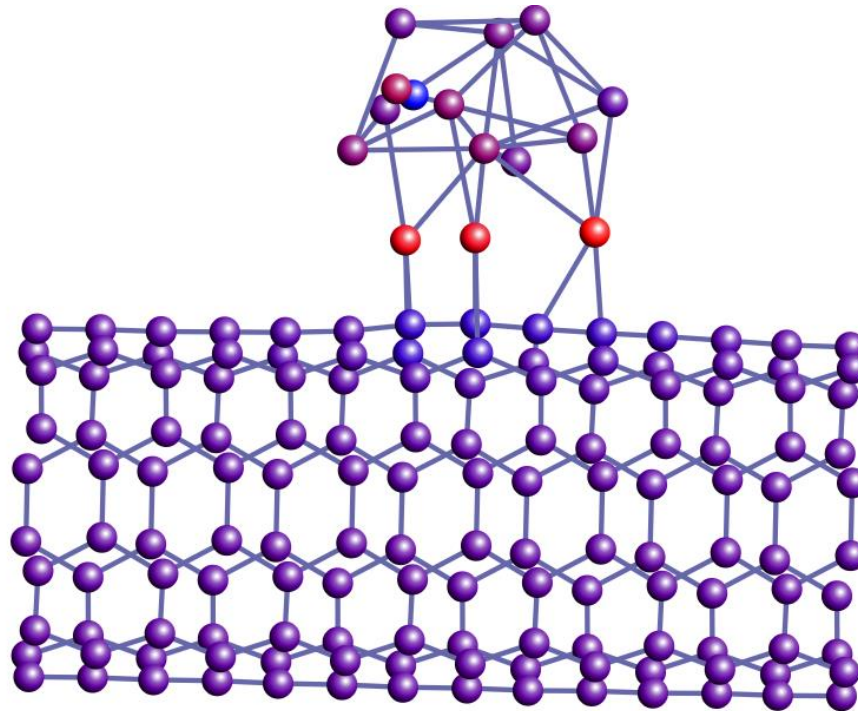
Structural Change
time= 20.000 (fs) Energy= -530.80623 (Hartree)
Frame 21/21
First << < Stop > >> Last
Make animation Every 1 frames
Delayed time (ms) 0
Start End File format not supp

supercell 1 x 1 x 1 Atoms rendering Bonds wire Number off Symbol off BGC white Atom Size 1.0 Bond Thickness 1.0 Bond factor 1.0 Cells off Axes off Perspective off Structure off Dynamics on Net Charge off Force on Velocity off Rot on x 0 Rot on y 0 Rot on z 0
OpenMX website Rotate: click+drag Zoom: scroll Translate: ctrl+click+drag Note: click the middle button for Safari

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Analysis of structure optimization

Using a md file, one can easily check spatial charge distribution.



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