Simulations of Molecular Dynamics by AlmaLinux v.s. Debian-12 OS

> Motohiko Tanaka, PhD, Japan June-November, 2024

https://github.com/Mtanaka77/

Settings and tests for simulations

Installation of AlmaLinux-9, May 2024, and Debian 12, Nov. 2024

Use Windows 11, VirtualBox 7 Linux gfortran and pip packages

Simulations, https://github.com/Mtanaka77/ >> Three-dimensional electrostatic p3m code, with tip5p and Ewald sums >> Siesta-4.1b, with mpich, fft3w, OpenBLAS, Scalapack

Firefox works for AlmaLinux and Debian

Windows11-MT [Running] - Oracle VM VirtualBox									ב	\times		
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Wind

MIT on Climate Change

300+

9

Number of MIT's 1,080 faculty members working on projects to address climate change

Number of MIT's five schools (and one college) whose faculty are working on questions related to climate change

🌖 👽 Right Control

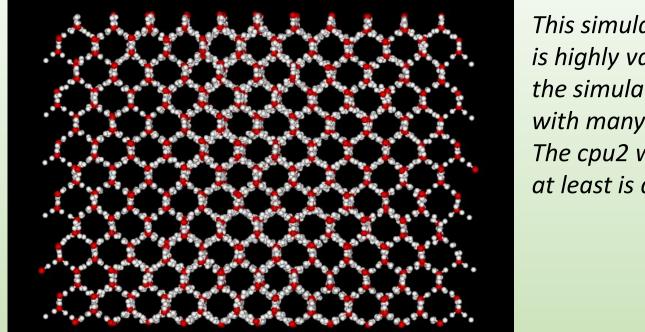
What MIT is doing on climate change

Linux terminal shows installation of mpich-4, fftw-3, and p3mtip5, Siesta-4.1b

Windows11-MT [Running] - Oracle VM VirtualBox

Windows11-MT [Running] - Oracle VM VirtualBox - C								
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[mtanaka@physique ~]\$ ls								
aaa.sh	Documents	old_atmfuncs.f						
aaa.txt	Downloads	old-SIESTA_atmfuncs.f						
a_bashrc	EnglishKey	OpenBLAS-0.3.27						
aclocal.m4		OpenBLAS-0.3.27.tar.gz						
a.f	fftw-3.3.10.tar.gz	openmpi-4.1.6.tar.gz						
a.f03	final_H_f_stress.F	p3mtip5						
a_mpich_how	final-SIESTA_H_f_stress.F	Pictures						
anaconda-ks.cfg		Public						
a_mpich_how anaconda-ks.cfg a.out	gpg-sign	scalapack-2.2.0						
arch.make	libopenblas	scalapack-2.2.0.tar.gz						
AUTOEXEC.BAT	LOCALE	sh_obj						
autogen.sh	log-fftw3	Siesta4						
autorun.inf	log-mpich	siesta-4.1-b4gcc						
bbb.txt	MPI_aggr	<pre>siesta-4.1-b4gcc.tar.gz</pre>						
C12H48-MD11	mpich-4.2.1	siesta-master.tar.gz						
COMMAND.COM	mpich-4.2.1.tar.gz	siesta.tar.gz						
conf-fftw3	MPI_chinv3	SLmake.inc_scalapack						
configure	MPI_expl	SUSE						
conf-mpich	MPI_nano	'System Volume Information						
conf-mpich-log	mrg37	Templates						
Desktop	Music	Videos						
[mtanaka@physique ~]\$								

Test of MD @p3mtip5p07a.f03, 5-points water model



This simulation run is OK, but timing is highly variable in time because the simulation in VirtualBox competes with many tasks of Windows 11. The cpu2 which should be 0.6 sec at least is different with the time steps.

time: e kin.W e img.W e kin(M) e lj e p3m ecr walltm e tot vm <ekin> <eimg> exc cpu cpu1 cpu2 cpu3 1.7095E+00 1.9537E-01 0.0000E+00 -1.6974E+02 3.0997E+01 5.1888E 20.0 T =-04 -1.3684E+02 8.656D+02 1.353D-01 9.893D-04 1.131D-04 0.000D+00 1.14.028D-04 1.106D+00 8.584D-03 15D+00 25.0 1.7269E+00 1.9599E-01 0.0000E+00 -1.6972E+02 3.0949E+01 5.3564E t= -04 -1.3685E+02 1.076D+03 1.095D-01 0.000D+00 9.993D-04 1.134D-04 1.743D+00 1.734D+00 8.680D-03 3.641D-04 0.0000E+00 -1.6976E+02 3.0940E+01 5.4725E 30.0 1.7385E+00 2.0207E-01 t =-04 -1.3688E+02 1.295D+03 1.117D-01 0.000D+00 1.006D-03 1.169D-04 5.6 95D-01 3.855D-04 5.607D-01 8.385D-03

Related pip3 packages

The initial states of water and hydrate are constructed (Dr. Matsumote, https://github.com/vitroid/). \$ pip3 install genice

Compilation goes all right for the genice software of CentOS 7. However, it goes the errors in the pairlist package and thus not go forward in AlmaLinux-9.

Debian 12

The Debian OS has been installed, and is tested by "mrg37" which is quite OK. Pip3 packages and 'pip3 install genice2' is successfully installed. The initial water configuration is perfect.

To compile Scalapack v.2

Download scalapack.2-2-0 and expand it. In BLACS, PBLAS, SRC, TOOLS, do \$ make (no option), except one difference is SRC.

Give—*fallow-argument-mismatch at Makefile's* \$ (*FC*) *line of SRC, and type* \$ *make*—*k when errors are shown.*

Scalapack is 10.7 MB for libscalapack.a

Test of ab-initio Siesta-4.1b code

A keyword -fallow-argument-mismatch in the arch.make file is added for AlmaLinux-9 and Debian-12

```
Architecture : gfortran-MPI
Compiler version: GNU Fortran (GCC) 11.4.1 20231218 (Red Hat 11.4.1-3)
Compiler flags : mpifort -O2 -fPIC -ftree-vectorize -march=native -fallow-argu
ment-mismatch
PP flags
Libraries
                : - DMPI - DFC HAVE ABORT
                : -lgomp -L/opt/openblas/lib -lopenblas -L/opt/scalapack/lib -l
scalapack
PARALLEL version
* Running on 6 nodes in parallel
>> Start of run: 2-JUN-2024 10:09:19
                          ***********************
                             WELCOME TO SIESTA
                          **********************
                                                                               •**२
reinit: Reading from c12h48.fdf
                                V.UZ683UU/
                                                      U.UZXCUXX
                                                                        ev/ang<sup>k</sup>*3
                       siesta:
                                        42,98698226 45,67350102
                                                                        kBar
                       siesta:
                       (Free)E+ p_basis*V orbitals = _____-2615.811579
                       (Free)Eharris+ p basis*V orbitals =
                                                                 -2615.811579
                       dhscf: Vacuum level (max, mean) = -0.569553
                                                                      -0.682007 eV
                       >> Start of run: 2-JUN-2024 10:09:19
                       >> End of run: 2-JUN-2024 10:11:55
                       Job completed
```

<mark>Overall results</mark>

The tests of classic and ab-initio molecular dynamics on AlmaLinux-9 OS are successful. Some alterations must be necessary on this specific operating system.

However, other sites including FFTW3 software fail by busy signal. The scalapack software is on unresolved errors. The pip3 of pairlist goes wrong in AlmaLinux-9.

Debian 12 OS is installed, and gcc, make, mpich, fftw3 are set up. It is tested with MD and water initial configuration of pip3 (by Dr. Matsumoto) and Siesta-4.1b, all of which are quite fine