

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

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*<https://github.com/Mtanaka77/>*

# *Settings and tests of simulations*

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

*Use Windows 11, VirtualBox 7 to login Linux OS  
Linux gfortran and pip3 packages*

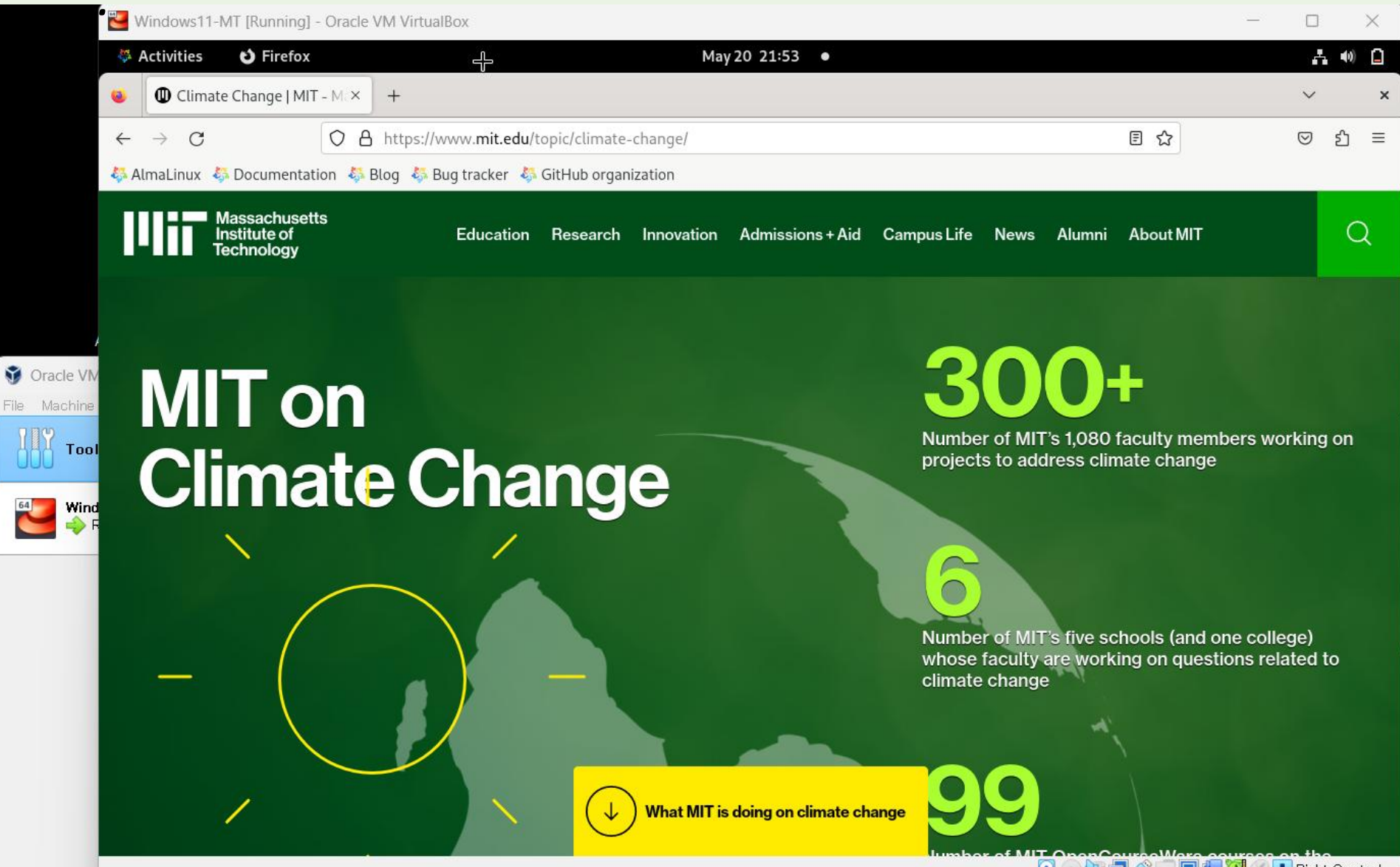
*Simulations, cf. <https://github.com/Mtanaka77/>*

*>> Three-dimensional electrostatic p3mtip5 code,  
with tip5p and Ewald sums*

*>> Siesta-4.1b, with mpich4 fft3w, OpenBLAS,  
and Scalapack*

*Firefox works for AlmaLinux and Debian OS*

*Debian can view all of internet, but AlmaLinux is quite limited*



# Windows to Linux terminal: Installation of mpich4, fftw3 Tests of p3mtip5, and Siesta-4.1b codes

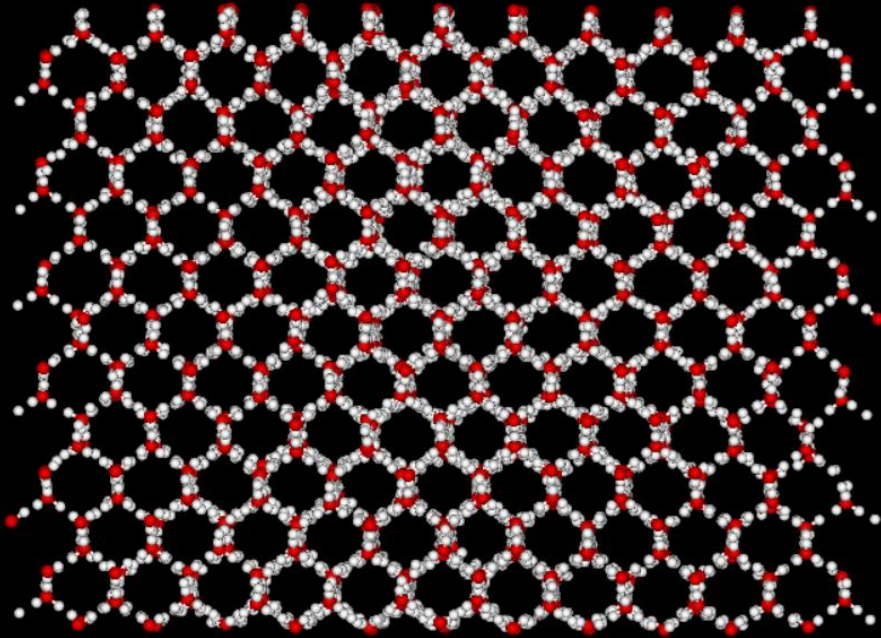
```
アクティビティ  端末  12月7日 10:32

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mtanaka@physique:~$ ls
C12H48-MD11      arch0bjja      mrg37          siesta7
Genece3-mh3exyz  arch0bjb      p3mtip5        ダウンロード
Genice2Log       bashrc-mtanaka3  sh_obj        テンプレート
Genice3          cnt3-para      siesta-4.1-b4gcc  デスクトップ
GeniceLog        conf-fftw3      siesta-4.1-b4gcc.tar.gz  ドキュメント
MPI_chginv       conf-mpich      siesta-4.1-b4gcc0.tar.gz  ビデオ
OperblasLog      make-BLACS-SRC  siesta-4.1-b4gccA  音楽
aaa-p3m.sh       make-PBLAS-SRC  siesta-4.1-b4gccA.tar.gz  画像
arch.make-MPIOMP  make-SRC        siesta4.1-MPI      公開
arch.make-OMP     make-TOOLS      siesta4.1-MPI-OMP

mtanaka@physique:~$ df
ファイルシス  1K-ブロック    使用    使用可  使用%  マウント位置
udev          1971352          0    1971352    0%  /dev
tmpfs         400732         1252    399480    1%  /run
/dev/sda1     30018340 10344732 18123428   37%  /
tmpfs         2003640          0    2003640    0%  /dev/shm
tmpfs          5120            8      5112    1%  /run/lock
tmpfs         400728         100    400628    1%  /run/user/1000
mtanaka@physique:~$
```

# Test of MD: @p3mtip5p07a.f03, by 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_c_r      e_lj      e_p3m
          e_tot      walltm      vm      exc      <ekin>      <eimg>      cpu
0
  cpu1      cpu2      cpu3
t=      20.0  1.7095E+00  1.9537E-01  0.0000E+00 -1.6974E+02  3.0997E+01  5.1888E
-04 -1.3684E+02      8.656D+02  1.353D-01  0.000D+00  9.893D-04  1.131D-04      1.1
15D+00  4.028D-04  1.106D+00  8.584D-03
t=      25.0  1.7269E+00  1.9599E-01  0.0000E+00 -1.6972E+02  3.0949E+01  5.3564E
-04 -1.3685E+02      1.076D+03  1.095D-01  0.000D+00  9.993D-04  1.134D-04      1.7
43D+00  3.641D-04  1.734D+00  8.680D-03
t=      30.0  1.7385E+00  2.0207E-01  0.0000E+00 -1.6976E+02  3.0940E+01  5.4725E
-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 configuration of water and hydrate are constructed (Dr. Matsumoto, <https://github.com/vitroid/>)  
\$ pip3 install genice2*

*Compilation goes OK in genice2 software of CentOS 7. However, it goes 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. The pip3 packages and ‘pip3 install genice2’ is successfully installed. The initial water configuration turns to be perfect.*

## *To compile Scalapack Version 2*

*“This is the inside story of Scalapack’s make.”*

*One downloads scalapack-2.2.0.tgz and expands it.*

*In BLACS, PBLAS, SRC, TOOLS, do \$ make (no option), except one difference in SRC.*

*Give -fallow-argument-mismatch at Makefile’s \$(FC) line in SRC, then type \$ make -k when one meets errors.*

*Scalapack is 10.7 MB for libscalapack.a*



# Test of *ab-initio* Siesta-4.1b code

*A keyword -fallow-argument-mismatch is added in the arch.make file of Siesta-4.1b for AlmaLinux-9 and Debian-12*

```
Siesta Version   : v4.1-b4
Architecture    : mpifort-MPI
Compiler version: GNU Fortran (Debian 12.2.0-14) 12.2.0
Compiler flags  : mpifort -O2 -ftree-vectorize -fprefetch-loop-arrays -march=native -fallow-argument-mismatch -fPIE
PP flags        : -DMPI -DFC_LAPACK -DSCF
Libraries       : -lgomp -L/...
calapack.a
PARALLEL version

>> Start of run: 4-DEC-2024

siesta: Cell volume = 720.000000 Ang**3

siesta: Pressure (static):
siesta: Solid Molecule Units
siesta: 0.00029221 0.00031048 Ry/Bohr**3
siesta: 0.02683002 0.02850685 eV/Ang**3
siesta: 42.98689824 45.67350469 kBar
**:(Free)E+ p_basis*V_orbitals = -2615.811581
* (Free)Eharris+ p_basis*V_orbitals = -2615.811581
**:
dhscf: Vacuum level (max, mean) = -0.569552 -0.682007 eV

reinit: Reading from c12h48.f
>> Start of run: 4-DEC-2024 17:38:16
>> End of run: 4-DEC-2024 17:39:58
1:42 min./10 cycles/6-MPI
Job completed
```



## *Overall Results of AlmaLinux and Debian OS*

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

*Many internet sites including FFTW3 failed by busy signal, the pip3 compilation of pairlist was wrong in AlmaLinux-9.*

*Debian 12 OS was installed, and gcc, make, mpich, fftw3 were set up on top. It was tested with MD and water initial cof pip3 (by Dr. Matsumoto) and Siesta-4.1b, all of which were quite fine on Debian.*