

Introduction to High Performance Computing **Software** and **Lmod**

Ali Kerrache

HPC Specialist

- Software distribution on HPC clusters.
- How to find software packages and modules?
- Software stacks on **GreX** / **Alliance** clusters.
- **Hands on:**
 - How to use modules to search for installed software?
- Software installation [**some examples**]

Operating system package managers / repos:

- **Ubuntu:** ~\$ *sudo apt-get install <package>*
- **CentOS:** ~\$ *sudo yum install <package>*
- **On HPC:** users do not have **sudo!** [**NO NEED TO ASK FOR IT**]

Centralized software stack:

- **Software distributed via CVMFS:** software stacks and **modules**, ...
- **Local software:** modules, restricted software (VASP, Gaussian, ORCA, ...etc)

Local installation: usually to \$HOME or \$PROJECT, ...

- **Download the code** [[sources/binaries](#)]: wget, git clone, curl, ... etc.
- **Settings:** load dependencies, set environment variables, ... etc.
- **Build:** *./configure {cmake ..} +opts; make; make test {check}; make install*



- **Home made:** programs, scripts and tools, ... etc.

Up to a user, ... Help is available!

→ **Free Software:** GNU Public License.

- ◆ Open Source, **Binaries**, **Libraries**, Compilers, Tools, ...

*Maintained by **analysts** and installed as **modules**.*

→ **Commercial Software:** restricted [VASP, STATA, ... etc]

- ◆ **Contact support** and provide more details about the license, ...
- ◆ We install the program & protect it with a **POSIX** group.

*Maintained by **analysts** and **admins**.*

What is a module?

- Configuration files with instructions to modify or initialize environment variables such as `PATH`, `LD_LIBRARY_PATH`. ... `etc` in order to use different installed programs.
- The modular architecture allows multiple versions of the same program to be installed without conflicts.

Why modules?

- Control different versions of the same program.
- Avoid conflicts between different versions and libraries.
- Set the right path to each program and library.



- module **list**; module **avail**
- module **spider** <soft>/<version>
- module **load** <deps> <soft>/<version>
- module **unload {rm}** <soft>/<version>

- module **show** <soft>/<version>
- module **help** <soft>/<version>
- module **whatis** <soft>/<version>

- module **purge**; module --force **purge**
- module **use** ~/modulefiles
- module **unuse** ~/modulefiles

```
[~@bison]$ module list
```

Currently Loaded Modules:

SBEnv (S)

Where:

S: Module is Sticky, requires --force to unload or purge

```
[~@rorqual3: ~]$ module list
```

Currently Loaded Modules:

```
1) CCconfig          4) gcc/12.3 (t)  7) libfabric/1.18.0
10) openmpi/4.1.5 (m) 13) aocl-lapack/5.1
2) gentoo/2023 (S)  5) hwloc/2.9.1   8) pmix/4.2.4
11) flexiblas/3.3.1 14) StdEnv/2023 (S)
3) gcccore/.12.3 (H) 6) ucx/1.14.1   9) ucc/1.2.0
12) aocl-blas/5.1
```



- A set of compilers and libraries:
 - ◆ **GCC, AOCC, Intel** compilers, ...
 - ◆ **Libraries:** hdf5, boost, netcdf, petsc, gsl, gdal, geos, proj, ... etc.
- Modules hierarchy:
 - ◆ **arch:** branch for a given architecture {~~sse3~~, avx2, **avx512**}
 - ◆ **CUDA:** any program using GPU acceleration under the same tree.
 - ◆ **Core modules:** java, perl, ... etc.
 - ◆ **Compiler:**
 - **GCC:** programs compiled with gcc
 - **AOCC:** programs compiled with gcc
 - **Intel:** compiled with Intel
 - ◆ **OpenMPI:** Compiler [**Intel, GCC, AOCC**] / OpenMPI
- Possibility to maintain one or more software stacks.

- Grex environment [default]: SBEnv
 - ◆ no module loaded by default, two architectures: avx2/avx512
 - ◆ use module spider <software name> to search for modules
 - ◆ Compilers {GCC, AOCC, Intel}, MKL, PETSc, GSL, NetCDF... etc.
 - ◆ Gaussian, ANSYS, MATLAB, ... etc.

- The Alliance (CC) environment [optional]: CCEnv
 - ◆ Switch to CCEnv; load a standard environment; choose the architecture[avx2/avx512], use module spider <software name>
 - module load CCEnv
 - module load arch/avx512
 - module load StdEnv/2023
 - module load gcc/12.3 geant4/11.3.0

Compilers/Libraries and more:

- **Compilers:** GCC [8.5 - 14.3]; Intel [2019, 2023], ... etc.
- **Libraries:** HDF5, PETSc, GSL, MKL, Libxc, Boost, NetCDF, ...
- **Gaussian, ANSYS, MATLAB, VASP, ORCA,** MCR, Java, Python, R, ... etc.
- LAMMPS, GROMACS, OpenMM, QE, OpenBABEL, ... etc.

Software maintenance on Grex and Alliance clusters:

- We install programs and update modules on request from users.
- Search for a program using “**module spider** <**name of your program**>”
- If not installed, ask support “support@tech.alliancecan.ca”
- We will install the module and/or update the version.
- **For commercial software, contact us before you purchase the code:**
 - ◆ to check license type.
 - ◆ see if it will run under Linux environment, ... etc.

Useful commands for working with modules

- module **list**
- module **avail**
- module **spider** <soft>/<version>
- module **load** <soft>/<version>
- module **unload {rm}** <soft>/<version>
- module **show** <soft>
- module **help** <soft>
- module **whatis** <soft>
- module **purge**; module --force **purge**
- module **use** ~/modulefiles; module **unuse** ~/modulefiles



List of modules: python, java, perl, hdf5, netcdf, **lammps**, **gromacs**, cp2k, ...

- Pick one module from the above list <myprogram>
- Run the command: **module spider** <myprogram>
- What to expect:
 - ◆ The module does not exist {ask for support if needed}
 - ◆ One or more versions of the module are available.
 - If many, pick one and run: **module spider** <myprogram>/<version>
 - Read the instructions, load the dependencies and the module
 - Experiment with other commands: **module list**, **module show myprogram**, **module help myprogram**, **module whatis**, **module rm**, ...
- Run “**module purge**” and repeat the exercise for another program.
- Switch to CCEnv, then load lammps-omp and gromacs.



Directory: quick-md-run

→ Have a look to the scripts using `cat {more or less}` command:

- Grex: `grex-runImp*.sh`
- CC/MC: `mc-runImp-*.sh`

Example: `cat grex-runImp-1cpu-serial.sh`

→ Run the scripts from the login node:

★ Grex:

- Serial test: `sh ./grex-runImp-1cpu-serial.sh`
- OpenMP test: `sh ./grex-runImp-4cpu-openmp.sh`
- MPI test: `sh ./grex-runImp-4cpu-mpi.sh`

★ CC/MC:

- Serial test: `sh ./mc-runImp-1cpu-serial.sh`
- OpenMP test: `sh ./mc-runImp-4cpu-openmp.sh`
- MPI test: `sh ./mc-runImp-4cpu-mpi.sh`



```
[~@bison ~]$ cvmfs_config probe  
Probing /cvmfs/cvmfs-config.computeCanada.ca... OK  
Probing /cvmfs/soft.computeCanada.ca... OK  
Probing /cvmfs/restricted.computeCanada.ca... OK
```

```
[~@bison ~]$ ls -l /cvmfs/  
cvmfs-config.computeCanada.ca [~@~]$ module load CCEnv  
restricted.computeCanada.ca [~@~]$ module load arch/avx512  
soft.computeCanada.ca [~@~]$ module load StdEnv/2023  
[~@~]$ module spider geant4  
[~@~]$ module spider geant4/11.3.0
```

```
[~@bison ~]$ ls /cvmfs/  
cvmfs-config.computeCanada.ca restricted.computeCanada.ca soft.computeCanada.ca  
[~@~]$ module load StdEnv/2023 gcc/12.3  
geant4/11.3.0
```



```
[~@yak ~]$ ls /cvmfs/  
cvmfs-config.computeCanada.ca restricted.computeCanada.ca soft.computeCanada.ca  
[~@yak ~]$ ls /cvmfs/neurodesk.ardc.edu.au  
[~@yak ~]$ ls /cvmfs  
  cvmfs-config.computeCanada.ca neurodesk.ardc.edu.au  
  restricted.computeCanada.ca soft.computeCanada.ca  
[~@yak ~]$ ls /cvmfs/neurodesk.ardc.edu.au/neurodesk-modules  
[~@yak ~]$ module use /cvmfs/neurodesk.ardc.edu.au/neurodesk-modules  
[~@yak ~]$ module spider fsl  
[~@yak ~]$ module spider functional_imaging/fsl/6.0.7.18  
[~@yak ~]$ module load functional_imaging/fsl/6.0.7.18  
[~@yak ~]$ module load singularity  
[~@yak ~]$ module list  
[~@yak ~]$ which fsl  
/cvmfs/neurodesk.ardc.edu.au/containers/fsl\_6.0.7.18\_20250928/fsl  
https://neurodesk.org/getting-started/neurocontainers/cvmfs/
```

GreX:

- <https://um-grex.github.io/docs/software/using-modules/>
- <https://um-grex.github.io/docs/software/software-list/>

Alliance clusters:

- https://docs.alliancecan.ca/wiki/Utiliser_des_modules/en
- https://docs.alliancecan.ca/wiki/Available_software

Modules:

- https://lmod.readthedocs.io/en/latest/010_user.html
- <https://modules.readthedocs.io/en/v4.1.3/index.html>
- https://lmod.readthedocs.io/en/stable/015_writing_modules.html

Building software



- **Local installation** [user's directory: home, project]:
 - ◆ R packages; Julia packages, Perl modules
 - ◆ Python packages: **virtual environment**
 - ◆ Home made programs and **commercial** software.

- **Installation with:**
 - ◆ make; make test {check}; make install
 - ◆ configure; make; make test {check}; make install
 - ◆ cmake; make; make test {check}; make install

- **Java applications:** jar files
- **Containers:** Singularity, Aptainer, Podman, Pixis {*separate talk*}
 - ◆ build the image and run your program using the container



What programs are installed locally?

- **R** packages: minimal installation
 - ◆ **R as a module**: users can install the packages in their home directory.
- **Python** as a module: python and scipy-stack or scipy-bundle
 - ◆ users can install the packages needed in their **home** or **project** directories.
- **Julia** as a module: **packages** installed locally
- **Perl** as a module:
 - ◆ users can install the packages needed in their home directory.
- Other software installed locally:
 - ◆ **Home made programs** {up to a user or a group}
 - ◆ **Restricted** and **licensed software** that can not be distributed
 - ◆ **Custom software**: patch from a user, changing parts of the code, ... etc.



- Download and unpack the code: `wget, ... gunzip, ... etc.`
- Load a compiler and dependencies: `module load gcc openmpi fftw`
- Configure the program
 - ◆ If configure not included, run: `autoreconf -fvi` [to generate it].
 - ◆ `./configure --help` [to see the different options].
 - ◆ `./configure --prefix=<path to install dir> {+other options}`
- Compile and test:
 - ◆ `make; make -j4`
 - ◆ `make check; make test`
- Install the program:
 - ◆ `make install`
- Set the path: `export PATH=${PATH}:<path to install dir>/bin`



- Download and unpack the code: `wget, ... gunzip, ... etc.`
- Load a compiler & dependencies: `module load gcc omp fftw cmake`
- Configure the program: `you may need to load cmake module`
 - ◆ `mkdir build && cd build`
 - ◆ `cmake .. --help` [to see the different options].
 - ◆ `cmake .. -DCMAKE_INSTALL_PREFIX=installdir {+options}`
- Compile and test:
 - ◆ `make; make -j8`
 - ◆ `make check; make test`
- Install the program:
 - ◆ `make install`



- Install **R** packages: `sp` and `dplyr`
- Install **python** packages: `cutadapt`
- Install **Perl** packages: `Hash::Merge`
- Install **Julia** packages: `JLD` (requires HDF5)
- Install **Java** applications: `Trimmomatic`
- Install **STAR**: uses `make` (Makefile included in the archive)
- Install **Treemix**: uses `configure` to generate the Makefile
`./configure {+opts} && make && make test {check} && make install`
- Install **DIAMOND**: uses `cmake` to generate the Makefile
`mkdir build && cd build && cmake .. {+opts} && make -j4 && make install`
- Install **Geant4**: uses `cmake` to generate the Makefile
`mkdir build && cd build && cmake .. {+opts} && make -j4 && make install`



- **R packages:** rgdal, adegenet, stats, rjags, dplyr, sf, ... etc.
- **Choose a module version:** module spider r
- **Load R and dependencies** (gdal, geos, jags, gsl, udunits... etc):
 - ◆ **module load gcc r <+other external modules>**
- **Launch R and install the packages:**

```
~$ R
```

```
> install.packages("dplyr")
```

```
'lib =/cvmfs/soft.computecanada.ca/easybuild/{..}/R/library' is not writable
```

```
Would you like to use a personal library instead? (yes/No/cancel) yes
```

```
Would you like to create a personal library '~/R/{...}' to install packages into? (yes/No/cancel) yes
```

```
--- Please select a CRAN mirror for use in this session ---
```

```
> install.packages("other packages")
```



- Load the modules:
 - ◆ `module load <deps> python <+ext>`
- Create a virtual environment
 - ◆ `virtualenv ~/my_venv`
- Activate the virtual environment
 - ◆ `source ~/my_venv/bin/activate`
- Update pip
 - ◆ `pip install --no-index --upgrade pip`
- Install the packages
 - ◆ `pip install pandas`
 - ◆ `pip install -r requirements.txt`
 - ◆ ~~`python setup.py install`~~

```
module load gcc python
virtualenv ~/my_venv
source ~/my_venv/bin/activate
pip install cutadapt
pip install <other package>
pip freeze > ~/requirements.txt
deactivate
```

```
source ~/my_venv/bin/activate
cutadapt [+options]
deactivate
```



- **Example:** Hash::Merge; Logger::Simple; MCE::Mutex; threads ...
- **Load Perl module:** module load perl
- **Install the the first package using cpan or cpanm:**
 - ◆ ~\$ cpan install YAML
 - Would you like to configure as much as possible automatically? [yes] **yes**
 - What approach do you want? (Choose 'local::lib', 'sudo' or 'manual') [local::lib] **local::lib**
 - Would you like me to append that to /home/\$USER/.bashrc now? [yes] **yes**
- **Install the rest of the packages using cpan or cpanm:**
 - ◆ ~\$ cpan install Hash::Merge
 - ◆ ~\$ cpan install Logger::Simple
 - ◆ ~\$ cpan install MCE::Mutex
- Update the ~/.bashrc [for the first time, it asks to update ~/.bashrc]

- Download and unpack the code using wget, curl, ... etc.
- Load java module [run module spider java]
- Run the code.

→ Example: Trimmomatic

wget <https://github.com/usadellab/Trimmomatic/releases/download/v0.40/Trimmomatic-0.40.zip>

unzip Trimmomatic-0.40.zip

- Run the code
 - Load java module: module load openjdk
 - java -jar <path to>/trimmomatic-0.40.jar {+options if any}

→ Download the code {wget; curl; git clone; ...}:

```
wget https://github.com/alexdobin/STAR/archive/refs/tags/2.7.11b.tar.gz
```

→ Unpack the code: `tar -xvf 2.7.11b.tar.gz`

→ Load GCC compiler: `module load gcc`

→ Compile the code:

```
cd STAR-2.7.11b/source  
make
```

→ Copy the binaries and set the path:

```
mkdir -p ~/software/star/2.7.11b/bin  
cp STAR ~/software/star/2.7.11b/bin  
export PATH=$PATH:${HOME}/software/star/2.7.11b/bin
```



→ Download the source files:

`wget https://bitbucket.org/nygcresearch/treemix/downloads/treemix-1.13.tar.gz`

→ Unpack the source files: `tar -xvf treemix-1.13.tar.gz`

→ Change the directory: `cd treemix-1.13/`

→ Load the modules: `module load gcc boost`

→ Configure: `./configure --prefix=/home/$USER/software/treemix/1.13`

→ Compile and install: `make && make test && make install`

→ Set a path: `export PATH=$PATH:$HOME/software/treemix/1.13/bin`

→ Usage in a job script:

```
module load gcc boost
```

```
export PATH=$PATH:$HOME/software/treemix/1.13/bin
```

```
treemix {+options if any}
```

→ Download the source files:

`wget https://github.com/bbuchfink/diamond/archive/refs/tags/v2.1.25.tar.gz`

→ Unpack the source files: `tar -xvf v2.1.25.tar.gz`

→ Change the directory: `cd diamond-2.1.25/`

→ Load the modules: `module load gcc cmake`

→ `mkdir build && cd build`

→ Configure: `cmake .. --DCMAKE_INSTALL_PREFIX=${HOME}/Softs/diamond/2.1.25`

→ Compile and install: `make && make test && make install`

→ Set a path: `export PATH=$PATH:$HOME/Softs/diamond/2.1.25/bin`

→ Usage in a job script:

`module load gcc boost`

`export PATH=$PATH:$HOME/Softs/diamond/2.1.25/bin`

`diamond {+options if any}`



- Download the source files:
`wget https://gitlab.cern.ch/geant4/geant4/-/archive/v11.3.1/geant4-v11.3.1.tar.gz`
- Unpack the source files: `tar -xvf geant4-v11.3.1.tar.gz`
- Change the directory: `cd geant4-v11.3.1/`
- Load the modules: `gcc clhep tbb qt cmake ...`
- `mkdir build && cd build`
- Create a `configure.cmd` file {Link}
- `source ./configure.cmd`
- `make`
- `make test`
- `make install`

```
cmake .. -DCMAKE_INSTALL_PREFIX=${HOME}/Softs/geant4/11.3.1 -DCMAKE_BUILD_TYPE="Release"  
-DGEANT4_BUILD_MULTITHREADED="ON" -DBUILD_STATIC_LIBS:BOOL="ON"  
-DGEANT4_USE_GDML:BOOL="ON" -DXercesC_LIBRARY_RELEASE="/usr/lib64/libxerces-c.so"  
-DEXPAT_LIBRARY="/usr/lib64/libexpat.so" -DEXPAT_INCLUDE_DIR="/usr/include"  
-DCLHEP_DIR="$MODULE_CLHEP_PREFIX/lib/CLHEP-${CLHEPVERSION}"  
-DPKG_CONFIG_EXECUTABLE="/usr/bin/pkg-config" -DGEANT4_BUILD_STORE_TRAJECTORY:BOOL="ON"  
-DGEANT4_BUILD_TESTS:BOOL="ON" -DGEANT4_USE_RAYTRACER_X11="ON"  
-DGEANT4_USE_SYSTEM_ZLIB="ON" -DGEANT4_USE_OPENGL_X11="ON"  
-DGEANT4_ENABLE_TESTING="ON" -DGEANT4_USE_TBB="ON" -DGEANT4_USE_QT:BOOL="OFF"  
-DQT_QMAKE_EXECUTABLE="$MODULE_QT_PREFIX/bin/qmake" -DGEANT4_USE_QT_QT6="ON"  
-DGEANT4_INSTALL_EXAMPLES:BOOL="ON" -DGEANT4_INSTALL_DATA:BOOL="ON"  
-DGEANT4_INSTALL_DATASETS_NUDEXLIB="ON" -DGEANT4_INSTALL_DATASETS_TENDL="ON"  
-DGEANT4_INSTALL_DATASETS_URRPT="ON" -DGEANT4_USE_HDF5="OFF" -DGEANT4_USE_VTK="OFF"  
-DGEANT4_USE_FREETYPE="ON" -DGEANT4_USE_G3TOG4="ON" -DGEANT4_USE_GDML="ON"  
-DGEANT4_USE_SMARTSTACK="ON" -DGEANT4_USE_INVENTOR="OFF" -DGEANT4_USE_PTL_LOCKS="ON"  
-DGEANT4_USE_SMARTSTACK="ON" -DGEANT4_USE_SYSTEM_PTL="OFF" -DGEANT4_USE_USOLIDS="OFF"  
-DGEANT4_USE_XM="ON"
```

```
cmake .. -DCMAKE_INSTALL_PREFIX=${HOME}/Softs/geant4/11.3.1 -DCMAKE_BUILD_TYPE="Release"  
-DGEANT4_BUILD_MULTITHREADED="ON" -DBUILD_STATIC_LIBS:BOOL="ON"  
-DGEANT4_USE_GDML:BOOL="ON" -DXercesC_LIBRARY_RELEASE="$EBROOTGENTOO/lib64/libxerces-c.so"  
-DEXPAT_LIBRARY="$EBROOTGENTOO/lib64/libexpat.so"  
-DEXPAT_INCLUDE_DIR="$EBROOTGENTOO/include"  
-DCLHEP_DIR="$EBROOTCLHEP/lib/CLHEP-${CLHEPVERSION}"  
-DPKG_CONFIG_EXECUTABLE="/usr/bin/pkg-config" -DGEANT4_BUILD_STORE_TRAJECTORY:BOOL="ON"  
-DGEANT4_BUILD_TESTS:BOOL="ON" -DGEANT4_USE_RAYTRACER_X11="ON"  
-DGEANT4_USE_SYSTEM_ZLIB="ON" -DGEANT4_USE_OPENGL_X11="ON"  
-DGEANT4_ENABLE_TESTING="ON" -DGEANT4_USE_TBB="ON" -DGEANT4_USE_QT:BOOL="OFF"  
-DQT_QMAKE_EXECUTABLE="$EBROOTQT/bin/qmake" -DGEANT4_USE_QT_QT6="ON"  
-DGEANT4_INSTALL_EXAMPLES:BOOL="ON" -DGEANT4_INSTALL_DATA:BOOL="ON"  
-DGEANT4_INSTALL_DATASETS_NUDEXLIB="ON" -DGEANT4_INSTALL_DATASETS_TENDL="ON"  
-DGEANT4_INSTALL_DATASETS_URRPT="ON" -DGEANT4_USE_HDF5="OFF" -DGEANT4_USE_VTK="OFF"  
-DGEANT4_USE_FREETYPE="ON" -DGEANT4_USE_G3TOG4="ON" -DGEANT4_USE_GDML="ON"  
-DGEANT4_USE_SMARTSTACK="ON" -DGEANT4_USE_INVENTOR="OFF"  
-DGEANT4_USE_PTL_LOCKS="ON" -DGEANT4_USE_SMARTSTACK="ON"  
-DGEANT4_USE_SYSTEM_PTL="OFF" -DGEANT4_USE_USOLIDS="OFF" -DGEANT4_USE_XM="ON"
```

Thank you for your attention

Any question?

module load intel openmpi gsl netcdf

instdir=<path to the installation directory>

```
../configure --prefix=${instdir} --enable-mpi --enable-mpi-io --with-fft-flavor=fftw3-mkl  
--with-linalg-flavor=mkl --with-math-flavor=gsl --enable-debug="no"  
--enable-optim="standard" --enable-64bit-flags  
--with-linalg-libs="-L$MKLROOT/lib/intel64 -lmkl_scalapack_lp64  
-lmkl_blacs_openmpi_lp64 -lmkl_intel_lp64 -lmkl_sequential -lmkl_core -lm"  
--with-fft-incs="-I$MKLROOT/include/fftw -I$MKLROOT/interfaces/fftw3xf"  
--with-fft-libs="-L$MKLROOT/interfaces/fftw3xf -lfftw3xf_intel_lp64"  
--with-dft-flavor="atompaw+libxc+wannier90" --with-trio-flavor="netcdf" --enable-lotf  
--enable-macroave --enable-gw-dpc CC=mpicc CXX=mpic++ FC=mpif90  
F77=mpif77 F90=mpif90
```

```
./configure --with-blas-lapack-dir=$MKLROOT/lib/intel64 --prefix=${instdir}  
--with-cxx-dialect=C++11 --download-scalapack=yes --download-blacs=yes  
--download-superlu_dist=yes --download-mumps=yes --download-parmetis=yes  
--download-metis=yes --download-spooles=yes --download-cproto=yes  
--download-prometheus=yes --with-mkl_pardiso=1 --with-mkl_pardiso-dir=$MKLROOT  
--with-mkl-sparse-optimize=1 --with-scalar-type=complex --with-debugging=0 --with-hdf5=yes  
--with-hdf5-dir=$HDF5HOME --download-suitesparse=yes --download-fftw=${fftsrc}  
--download-amd=yes --download-adifor=yes --download-superlu=yes --download-triangle=yes  
--download-generator=yes --with-64-bit-pointers=no --with-cc=mpicc --CFLAGS='-O2  
-I$MKLROOT/include -mkl -fPIC ' --with-cxx='mpicxx' --CXXFLAGS='-O2 -I$MKLROOT/include  
-mkl -std=c++11 -fPIC ' --with-fc='mpif90' --FFLAGS='-O2 -I$MKLROOT/include -mkl -fPIC '  
--with-single-library=yes --with-shared-libraries=yes --with-shared-ld=mpicc  
--sharedLibraryFlags="-fPIC -mkl -fPIC" --with-mpi=yes --with-mpi-shared=yes  
--with-mpirun=mpiexec --with-mpi-compilers=yes --with-x=yes {+other options}  
make && make install
```



- Download and unpack the source files
- Load modules:
module load intel openmpi fftw cmake
- configure; compile; install
cd gromacs-5.1.4; mkdir build; cd build
cmake -DCMAKE_INSTALL_PREFIX=<path to install dir> -DBUILD_SHARED_LIBS=off
-DBUILD_TESTING=off -DREGRESSIONTEST_DOWNLOAD=off
-DCMAKE_C_COMPILER=`which mpicc` -DCMAKE_CXX_COMPILER=`which mpicxx`
-DGMX_BUILD_OWN_FFTW=on -DGMX_SIMD=SSE4.1 -DGMX_DOUBLE=off
-DGMX_EXTERNAL_BLAS=on -DGMX_EXTERNAL_LAPACK=on
-DGMX_FFT_LIBRARY=fftw3 -DGMX_GPU=off -DGMX_MPI=on -DGMX_OPENMP=off
-DGMX_X11=on ../gromacs-5.1.4

make -j4; make install



- Use Lmod commands to search for the modules:
 - ◆ Compilers, OpenMPI, NetCDF, HDF5, PETSc, Gaussian, ANSYS, MATLAB, ORCA, MCR, Java, Python, R, ... etc.

- Some packages require a local installation:
 - ◆ Home made programs, Python, Perl, R, Julia packages, ...

- Software maintenance on Grex and Alliance clusters:
 - ◆ Search for a program using “module spider <name of your program>”
 - ◆ If not installed, ask for support “support@tech.alliancecan.ca”
 - ◆ We will install the module and/or update the version.
 - ◆ For commercial software, contact us before you purchase the code:
 - to check license type.
 - see if it will run under Linux environment, ... etc.

Additional Slides



```
[~@narval2: ~]$ module list
```

Currently Loaded Modules:

```
1) CCconfig      5) hwloc/2.9.1      9) ucc/1.2.0      13) StdEnv/2023 (S)
2) gentoo/2023 (S) 6) ucx/1.14.1      10) openmpi/4.1.5 (m) 14) mii/1.1.2
3) gcccore/.12.3 (H) 7) libfabric/1.18.0 11) flexiblas/3.3.1
4) gcc/12.3 (t)    8) pmix/4.2.4      12) blis/0.9.0
```

Where:

S: Module is Sticky, requires --force to unload or purge

m: MPI implementations / Implémentations MPI

t: Tools for development / Outils de développement

H: Hidden Module

```
[~@bison ~]$ module list
```

Currently Loaded Modules:

1) **S**Env (S)

Where:

S: Module is Sticky, requires --force to unload or purge

Note:

- Before starting, make sure you have the appropriate software stack and the compilers and libraries you need.
- Use “module spider” to search for the programs.

CC software stack:

```
[~@bison ~]$ module load CCEnv
```

```
[~@bison ~]$ module load arch/avx512
```

```
[~@bison ~]$ module load StdEnv/2023
```

```
[~@bison: ~]$ module spider python
```

```
python:
```

Description:

The Python programming language. Homepage: <https://www.python.org/>

Versions:

```
python/3.10.14
```

```
python/3.10.16
```

```
python/3.11.8
```

```
python/3.11.11
```

```
python/3.12.9
```

For detailed information about a specific "python" package (including how to load the modules) use the module's full name.

Note that names that have a trailing (E) are extensions provided by other modules.

For example:

```
$ module spider python/3.12.9
```

Module example: python/3.12.9

```
[~@bison: ~]$ module spider python/3.12.9
```

```
python: python/3.12.9
```

Description:

The Python programming language. Homepage: <https://www.python.org/>

You will need to load all module(s) on any one of the lines below before the "python/3.12.9" module is available to load.

```
arch/avx512 gcc/13.2.0
```

```
arch/avx512 gcc/14.3.0
```

```
arch/avx512 intel-one/2024.1
```

```
cuda/12.4.1 arch/avx2 gcc/13.2.0
```

Help:

The Python programming language.

Homepage: <https://www.python.org/>

Use one of the following to load the module:

```
module load arch/avx512 gcc/13.2.0 python/3.12.9
```

```
module load arch/avx512 gcc/14.3.0 python/3.12.9
```

```
module load arch/avx512 intel-one/2024.1 python/3.12.9
```

```
module load cuda/12.4.1 arch/avx2 gcc/13.2.0 python/3.12.9
```



```
[~@bison: ~]$ module show python/3.12.9
/global/software/alma8/sb/modules/base/python/3.11.8:
prepend_path("CMAKE_PREFIX_PATH", "/global/software/alma8/sb/opt/base/python/3.11.8")
prepend_path("PATH", "/global/software/alma8/sb/opt/base/python/3.11.8/bin")
prepend_path("LD_LIBRARY_PATH", "/global/software/alma8/sb/opt/base/python/3.11.8/lib")
prepend_path("MANPATH", "/global/software/alma8/sb/opt/base/python/3.11.8/share/man")
setenv("MODULE_PYTHON_PREFIX", "/global/software/alma8/sb/opt/base/python/3.11.8")
prepend_path("MODULEPATH", "/global/software/alma8/sb/modules/python-3.11.8")
whatis("Description: The Python programming language. Homepage: https://www.python.org/")
family("python")
setenv("PIP_DISABLE_PIP_VERSION_CHECK", "1")
setenv("PYDEVD_DISABLE_FILE_VALIDATION", "1")
prepend_path("PYTHONPATH", "/global/software/alma8/sb/python3/site-packages")
help([[
The Python programming language.
Homepage: https://www.python.org/
]])
```

```
[~@bison: ~]$ module spider boost
```

```
boost:
```

Description:

Boost provides free peer-reviewed portable C++ source libraries, emphasizing libraries that work well with the C++ Standard Library

Versions:

```
boost/1.78.0
```

```
boost/1.85.0
```

Other possible modules matches:

```
xgboost
```

To find other possible module matches execute:

```
$ module -r spider '.*boost.*'
```

For detailed information about a specific "boost" package (including how to load the modules) use the module's full name.

Note that names that have a trailing (E) are extensions provided by other modules.

For example:

```
$ module spider boost/1.85.0
```

```
[~@bison: ~]$ module spider boost/1.85.0
```

```
boost: boost/1.85.0
```

Description:

Boost provides free peer-reviewed portable C++ source libraries, emphasizing libraries that work well with the C++ Standard Library

You will need to load all module(s) on any one of the lines below before the "boost/1.85.0" module is available to load.

```
arch/avx512 gcc/13.2.0
```

- ```
cuda/12.4.1 arch/avx2 gcc/13.2.0
```
- ```
module load arch/avx512 gcc/13.2.0 boost/1.85.0
```
- ```
module load cuda/12.4.1 arch/avx2 gcc/13.2.0 boost/1.85.0
```

Help:

Boost provides free peer-reviewed portable C++ source libraries, emphasizing libraries that work well with the C++ Standard Library. Boost libraries are intended to be widely useful, and usable across a broad spectrum of applications. The Boost license encourages both commercial and non-commercial use.

Homepage: <http://www.boost.org>

```
[~@bison: ~]$ module load arch/avx512 gcc/13.2.0 boost/1.85.0
```

```
[~@bison: ~]$ module show boost
```

```
/global/software/alma8/sb/modules/arch-avx512-gcc-13.2.0/boost/1.85.0:
```

```
prepend_path("CMAKE_PREFIX_PATH", "/global/software/alma8/sb/opt/arch-avx512-gcc-13.2.0/boost/1.85.0")
```

```
prepend_path("CPATH", "/global/software/alma8/sb/opt/arch-avx512-gcc-13.2.0/boost/1.85.0/include")
```

```
prepend_path("LIBRARY_PATH", "/global/software/alma8/sb/opt/arch-avx512-gcc-13.2.0/boost/1.85.0/lib")
```

```
prepend_path("LD_LIBRARY_PATH", "/global/software/alma8/sb/opt/arch-avx512-gcc-13.2.0/boost/1.85.0/lib")
```

```
setenv("MODULE_BOOST_PREFIX", "/global/software/alma8/sb/opt/arch-avx512-gcc-13.2.0/boost/1.85.0")
```

```
whatis("Description: Boost provides free peer-reviewed portable C++ source libraries, emphasizing libraries
that work well with the C++ Standard Library")
```

```
whatis("Homepage: http://www.boost.org")
```

```
setenv("BOOST_ROOT", "/global/software/alma8/sb/opt/arch-avx512-gcc-13.2.0/boost/1.85.0")
```

```
help([[
```

```
 Boost provides free peer-reviewed portable C++ source libraries, emphasizing libraries that work well with the C++ Standard Library.
```

```
 Homepage: http://www.boost.org
```

```
]])
```



[~@bison: ~]\$ **module spider gromacs**

gromacs:

Description:

GROMACS (GRONingen MACHine for Chemical Simulations) is a molecular dynamics package primarily designed for simulations of proteins, lipids and nucleic acids

Versions:

gromacs/2022

gromacs/2023.3

gromacs/2024.1

gromacs/2025.2

gromacs/2025.3

For detailed information about a specific "gromacs" package (including how to load the modules) use the module's full name.

Note that names that have a trailing (E) are extensions provided by other modules.

For example:

\$ **module spider gromacs/2025.3**

```
[~@bison: ~]$ module spider gromacs/2025.3
```

```
gromacs: gromacs/2025.3
```

Description:

GROMACS (GRONingen MAchine for Chemical Simulations) is a molecular dynamics package primarily designed for simulations of proteins, lipids and nucleic acids

You will need to load all module(s) on any one of the lines below before the "gromacs/2025.3" module is available to load.

```
arch/avx512 gcc/13.2.0 openmpi/5.0.6
cuda/12.4.1 arch/avx2 gcc/13.2.0
```

Help:

GROMACS (GRONingen MAchine for Chemical Simulations) is a molecular dynamics package primarily designed for simulations of proteins, lipids and nucleic acids. It was originally developed in the Biophysical Chemistry department of University of Groningen, and is now maintained by contributors in universities and research centers across the world. Starting from version 4.6, GROMACS is released under the GNU Lesser General Public License.

Homepage: <http://www.gromacs.org>



**Gaussian:** restricted software; requires a registration

<https://um-grex.github.io/docs/specific-soft/gaussian/>

```
[~@bison ~]$ module spider gaussian
```

```
gaussian:
```

```
 Versions:
```

```
 gaussian/g16.b01
```

```
 gaussian/g16.c01
```

For detailed information about a specific "gaussian" package (including how to load the modules) use the module's full name. Note that names that have a trailing (E) are extensions provided by other modules.

For example:

```
$ module spider gaussian/g16.c01
```

- [~@bison ~]\$ module load gaussian
- [~@bison ~]\$ module load gaussian/g16.c01
- [~@bison ~]\$ module load gaussian/g16.b01



**ORCA:** restricted software; requires a registration

<https://um-grex.github.io/grex-docs/specific-soft/orca/>

```
[~@bison ~]$ module spider orca
```

```
orca:
```

```
 Versions:
```

```
 orca/5.0.4
```

```
 orca/6.0.1
```

For detailed information about a specific "orca" package (including how to load the modules) use the module's full name. Note that names that have a trailing (E) are extensions provided by other modules.

For example:

```
$ module spider orca/6.0.1
```

```
→ [~@bison ~]$ module load arch/avx512 intel/2023.2 openmpi/4.1.6 orca/5.0.4
```

```
→ [~@bison ~]$ module load arch/avx512 gcc/13.2.0 openmpi/4.1.6 orca/6.0.1
```

```
[~@bison ~]$ module load arch/avx512 gcc/13.2.0 openmpi/4.1.6 orca/6.0.1
```

To execute ORCA, run:

```
 ${MODULE_ORCA_PREFIX}/orca orca.inp > orca.out
```



```
[~@bison ~]$ module spider lammeps [~@bison ~]$ module spider lammeps/2024-08-29p1
lammeps:
 Versions:
 lammeps/2021-09-29
 lammeps/2024-08-29p1-nep
 lammeps/2024-08-29p1
 You will need to load all module(s) on any one of the
 lines below before the "lammeps/2024-08-29p1"
 module is available to load.
 ● arch/avx512 gcc/13.2.0 openmpi/4.1.6
 ● arch/avx512 intel-one/2024.1 openmpi/4.1.6
 ● cuda/12.4.1 arch/avx2 gcc/13.2.0 openmpi/4.1.6
```

```
[~@bison ~]$ module load arch/avx512 gcc/13.2.0 openmpi/4.1.6 lammeps/2024-08-29p1
```

```
[~@bison ~]$ module list
```

Currently Loaded Modules:

- |                |     |                  |                    |                              |                          |
|----------------|-----|------------------|--------------------|------------------------------|--------------------------|
| 1) SBEnv       | (S) | 4) openmpi/4.1.6 | 7) fftw/3.3.10     | 10) zstd/1.5.6               | 13) gsl/2.7              |
| 2) arch/avx512 |     | 5) kim/2.3.0     | 8) openblas/0.3.26 | 11) hdf5/1.14.2              | 14) lammeps/2024-08-29p1 |
| 3) gcc/13.2.0  |     | 6) ffmpeg/7.0.2  | 9) eigen/3.4.0     | 12) netcdf/4.9.2+hdf5-1.14.2 |                          |

```
[~@bison ~]$ which lmp
```

```
/global/software/alma8/sb/opt/arch-avx512-gcc-13.2.0-openmpi-4.1.6/lammeps/2024-08-29p1/bin/lmp
```



```
[~@bison ~]$ module spider espresso
```

```
espresso:
```

```
 Versions:
```

```
 espresso/7.3.1+aocl-4.2.0
```

```
 espresso/7.3.1
```

```
 espresso/7.4.1+aocl-4.2.0
```

```
 espresso/7.4.1
```

```
 espresso/7.5+aocl-4.2.0
```

```
 espresso/7.5
```

For detailed information about a specific "espresso" package (including how to load the modules) use the module's full name. Note that names that have a trailing (E) are extensions provided by other modules.

For example:

```
$ module spider espresso/7.4.1
```

```
→ [~@bison ~]$ module spider espresso/7.4.1
```

```
→ [~@bison ~]$ module load arch/avx512 intel/2023.2 openmpi/4.1.6 espresso/7.4.1
```



```
[~@bison ~]$ module spider matlab
```

```
matlab:
```

```
Versions:
```

```
matlab/R2020B2
```

```
matlab/R2022A
```

```
matlab/R2023B
```

```
matlab/R2024A
```

For detailed information about a specific "matlab" package (including how to load the modules) use the module's full name.

Note that names that have a trailing (E) are extensions provided by other modules.

For example:

```
$ module spider matlab/R2024A
```

- [~@bison ~]\$ module load matlab/R2024A
- [~@bison ~]\$ module load matlab/R2023B
- [~@bison ~]\$ module load matlab/R2022A
- [~@bison ~]\$ module load matlab/R2020B2

# Find and load MCR

```
[~@bison ~]$ module spider mcr
```

```
mcr:
```

```
 Versions:
```

```
 mcr/R2020b
```

```
 mcr/R2022a
```

```
 mcr/R2023b
```

```
 mcr/R2024a
```

For detailed information about a specific "mcr" package (including how to load the modules) use the module's full name.

For example:

```
$ module spider mcr/R2024a
```

- [~@bison ~]\$ module load mcr/R2020b
- [~@bison ~]\$ module load mcr/R2022a
- [~@bison ~]\$ module load mcr/R2023b
- [~@bison ~]\$ module load mcr/R2024a

```

----- /global/software/alma8/sb/modules/base -----
adf/2019.305-impi eigen/3.4.0 julia/1.10.3 nodejs/20.16.0 ramalama/0.12.0
adf/2021.106-impi expect/5.45.4 julia/1.11.3 nodejs/20.18.1 ratarmount/1.2.0
adf/2021.107-impi fastqc/0.12.1 kitops/1.6.0 nodejs/22.4.1 rclone/1.70.3
adf/2023.104-impi feko/2021.2 libaec/1.0.6 nodejs/22.5.1 rclone/1.71.0 (D)
adf/2024.105-impi-aocl fFmpeg/7.0.2 libvori/220621 nodejs/22.11.0 rs-server/2025.05.1-513
adf/2024.105-impi fftw/3.3.10 libxml2/2.11.9 nvtop/3.1.0 rs-server/2025.09.0-387 (D)
adf/2025.104-impi-aocl flex/2.6.4 mathematica/14.2 nvtop/3.2.0 rust/1.87.0 (D)
admixture/1.3.0 freeglut/3.4.0 matlab-proxy/0.26.0 omlmd/0.1.6 rust/1.89.0 (D)
ansys/21.1 gaussian/g16.b01 matlab-proxy/0.27.1 (D) openeye/2022.2.1 samtools/1.20
ansys/2023R2 gaussian/g16.c01 (D) matlab/R2020B2 openjdk/11.0.22 scons/3.1.2
ant/1.10.15 gcc/8.5.0 matlab/R2022A openjdk/17.0.11_9 singularity/4.1.2
arch/avx2 git-annex/10.20250828 matlab/R2023B openjdk/17.0.12_7 singularity/4.2.2 (D)
arch/avx512 git-lfs/3.7.0 matlab/R2024A openjdk/17.0.13_11 skopeo/1.20.0
autotools/2022a git/2.49.0 mcr/R2020b openjdk/21.0.2 snp-sites/2.5.1
bamtools/2.5.2 git/2.51.0 (D) mcr/R2022a openjdk/21.0.3_9 snpEff/5.2F
beagle/5.4-20241029 globus/3.35.2 mcr/R2023b openjdk/21.0.4_7 sparsehash/2.0.4
birch/3.90 globus/3.36.0 (D) mcr/R2024a openjdk/21.0.5_11 sqLite/3.35.5
birch/4.0 gmp/6.2.1 megahit/1.2.9 openjdk/21.0.6_7 stata/15.0-fagfs
buildah/1.41.0 gmp/6.3.0 (D) metashape-pro/2.1.4 openssl/3.4.0 stata/18.0-ffin (D)
busco/5.8.3 gnina/1.1 metashape-pro/2.2.1 openstack-client/8.2.0
cfitsio/4.4.1 gnina/2024 metashape-pro/2.2.2 (D) ovito/3.12.0
cfitsio/4.5.0 gnuplot/5.4.2 (D) micro/2.0.14 ovito/3.12.3
cmake/3.28.4 gnuplot/6.0.2 (D) mii/1.1.1 ovito/3.13.1 (D)
cmake/3.31.1 golang/1.24.4 minipae/2.28 pandoc/3.6.1
cmake/4.1.1 golang/1.25.1 (D) molder/7.3 paraview-offscreen/5.10.1
code-server/4.103.1 haploview/4.2 mpfr/4.2.1 picard/3.3.0
compleasm/0.2.6 hmmer/3.4 podman-compose/1.5.0 podman-tui/1.7.0
cppzmq/4.10.0 httgettoken/2.0-2 nbo/nbo7-2021 podman/5.6.0
cuda/11.8.0 intel-one/2023.2 nextflow/24.10.0 podman/5.6.1 (D)
cuda/12.2.2 intel-one/2024.1 (D) ninja/1.10.2 process-compose/1.75.1
cuda/12.4.1 intel/2023.2 nodejs/18.20.2 prodigal/2.6.3
cudnn/8.8.1.3+cuda-11.8.0 intelmpi/2019.8 nodejs/18.20.4 python/3.11.8
deepvariant/1.8.0-gpu intelmpi/2021.10 nodejs/18.20.5 qiime2/2024.10
deepvariant/1.8.0 (D) jellyfish/2.3.1 nodejs/20.12.2 quast/5.3.0
dejagnu/1.6.3 jemalloc/5.3.0 nodejs/20.15.1 r/4.3.3
diamond/2.1.10 jq/1.7
----- /opt/lmod/stacks -----
CCEnv (S) SBEnv (S,L)
----- This is a list of module extensions. Use "module --nx avail ..." to not show extensions. -----
autoconf (E) automake (E) boftools (E) gettext (E) htstlib (E) java (E) libtool (E)

These extensions cannot be loaded directly, use "module spider extension_name" for more information.

```

module avail

module spider python

module spider java

module spider gromacs

module load arch/avx512

gcc/13.2.0 openmpi/5.0.6

module avail

module spider <soft>

module spider <soft>/<ver>

module help <soft>

module show <soft>

module purge

```

----- /global/software/alma8/sb/modules/arch-avx512-gcc-13.2.0-openmpi-5.0.6 -----
arpack-ng/3.9.1+mkl-2024.1 (D) gromacs/2023.3 hdf5/1.14.6 (D) parmetis/4.0.3-shared scotch/7.0.5 (D) zoltan/3.901
asynch/git-a5d1d77 gromacs/2024.1 imb/2021.7 pnetcdf/1.14.0 sundials/7.4.0+mkl-2024.1
asynch/1.4.3 (D) gromacs/2025.2 openfoam/9 raxml-ng/1.2.1 taudem/5.3.8
espresso/7.5+aocl-4.2.0 gromacs/2025.3 (D) openmm/8.3.1 scalapack/2.2.2 valgrind/3.24.0
fftw/3.3.10 (D) hdf5/1.12.3 paraview/5.13.3 scotch/6.0.9 yast/0.11.3

----- /global/software/alma8/sb/modules/arch-avx512-gcc-13.2.0 -----
aocl/4.2.0 eccodes/2.31.0 hisat2/2.2.1 openbabel/3.11.1 qt/6.9.1 (D)
aocl/4.2.0-64 (D) eccodes/2.40.0 (D) homer/5.1 openblas/0.3.26 r/4.4.1+aocl-4.2.0
armadillo/11.4.3 fasttree/2.1.11 intelmpi/2019.8 openblas/0.3.28 (D) r/4.4.1+mkl-2019.5
armadillo/14.2.2 (D) fftw/3.3.10 intelmpi/2021.10 (D) openmpi/4.1.6 (L,D) r/4.4.1+mkl-2024.1
arpack-ng/3.9.1+mkl-2019.5 flexpart/11 jags/4.3.2+mkl-2019.5 openmpi/5.0.6 (D) r/4.5.0+mkl-2024.1
arpack-ng/3.9.1+mkl-2024.1 gate/9.4 jags/4.3.2+mkl-2024.1 (D) opennurbs/8.12 root/6.32.08 (D)
arrow/18.1.0 gatk/4.6.1.0 jasper/4.0.0 pandaseq/2.11 root/6.34.02 (D)
autodock-vina/1.2.7 gdal/3.9.1 kim/2.3.0 potraeconv/2.1.1 samtools/1.20 (D)
autodock/4.2.6 gdal/3.10.0 (D) libint-cp2k/2.6 proj/9.2.0 scipy-bundle/2023+python-3.10.14
blastplus/2.16.0 geant4/11.2.2 libkc/5.1.5 proj/9.5.0 (D) scipy-bundle/2023+python-3.10.16
blastplus/2.17.0 (D) geant4/11.3.0 (D) libkc/6.2.2 (D) python/3.10.14 stringtie/3.0.0
blat/3.7 geos/3.13.0 metis/5.1.0 python/3.11.8 (D) scipy-bundle/2023+python-3.11.8
bliis/0.9.0 gibbs2/1.0 metis32/5.1.0 python/3.11.11 superlu/5.3.0+mkl-2019.5
boost/1.78.0 gplp/5.0 mkl/2019.5 python/3.11.11 (D) tbb/2021.13.0
boost/1.85.0 (D) qmp/6.3.0 (D) mkl/2024.1 (D) python/3.12.9 (D) udunits/2.2.28
bwa/0.7.18 grace/5.99.0 mpfr/4.2.1 (D) qhull/2020.2 vt/9.4.0
cgald/5.5 gsl/2.7 mustang/3.2.4 qrupdate/1.1.2 wxwidgets/3.0.2
cgald/6.0.1 (D) hdf5/1.12.3 nco/5.3.1 qt/6.7.1 xfemm/4.0
cistem/1.0.0 hdf5/1.14.2 ncview/2.1.11 qt/6.8.1 xtb/6.7.1+mkl-2019.5
clhep/2.4.7.1 hdf5/1.14.6 netcdf/4.9.2+hdf5-1.14.2 qt/6.8.3

----- /global/software/alma8/sb/modules/arch-avx512 -----
aocc/4.2.0 bowtie2/2.5.4 gcc/9.5.0 gmp/6.3.0 openblas/0.3.26 r/4.3.3
autotools/2022a (D) circos/0.69-9 gcc/11.5.0 intel-one/2024.1 (D) perl/5.38.2 vcfTools/0.1.16
bedtools/2.31.1 cuda/12.2.2 gcc/13.2.0 (L) intel/2019.5 perl/5.40.1 (D)
binutils/2.42 eigen/3.4.0 (D) gcc/14.3.0 (D) intel/2023.2 (D) prsice/2.3.5

----- /global/software/alma8/sb/modules/base -----
adf/2019.305-imp1 eigen/3.4.0 julia/1.10.3 nodejs/20.16.0 ramalama/0.12.0
adf/2021.106-imp1 expect/5.45.4 julia/1.11.3 (D) nodejs/20.18.1 ratarmount/1.2.0
adf/2021.107-imp1 fastgo/0.12.1 ktopops/1.6.0 nodejs/22.4.1 rclone/1.70.3
adf/2023.104-imp1 feko/2021.2 libaec/1.0.6 nodejs/22.5.1 rclone/1.71.0 (D)
adf/2024.105-imp1-aocl ffmpeg/7.0.2 libvori/220621 nodejs/22.11.0 (D) rs-server/2025.05.1-513
adf/2024.105-imp1 fftw/3.3.10 libxml2/2.11.9 nvtop/3.1.0 rs-server/2025.09.0-387 (D)
adf/2025.104-imp1-aocl (D) flex/2.6.4 libxlm2/2.11.9 nvtop/3.2.0 (D) rust/1.87.0
admixture/1.3.0 freeglut/3.4.0 mathematica/14.2 omlmd/0.1.6 rust/1.89.0 (D)
ansys/21.1 gaussian/g16.b01 matlab-proxy/0.26.0 openeye/2022.2.1 samtools/1.20
lines 1-46

```

module load arch/avx512  
gcc/13.2.0 openmpi/5.0.6

module avail

module spider gromacs  
module spider cp2k

If not available:  
→ contact support  
[support@tech.alliancecan.ca](mailto:support@tech.alliancecan.ca)