



## Software on HPC clusters

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### **Outline**

- ★ Software distribution on HPC clusters
  - Env. modules, Why modules? How to find software?
  - Local and CC Software stacks on Grex, CVMFS
- ★ How to build software from sources
  - Autotools and CMake
  - Compiling best practices, optimizations for CPUs
- ★ How to deal with interpreted languages/libraries
  - R packages
  - Python modules
- ★ Containers in HPC, : Singularity and others



## **Software distribution**

### Operating system package managers / repos:

- ★ Ubuntu: ~\$ sudo apt-get install <package>
- ★ CentOS: ~\$ sudo yum install <package>
- ★ On HPC: users do not have sudo!

#### **Using a centralized HPC software stack:**

- ★ Software distributed via CVMFS: CC software stack (CC clusters), ...
- ★ Local software: legally restricted software (VASP, Gaussian, ...)

#### Local installation: usually to \$HOME or \$PROJECT

- ★ Get the code: download the sources/binaries: wget, git clone, ... etc.
- ★ Settings: load dependencies, set environment variables, ... etc.
- ★ Build: ./configure {cmake ..} +opts; make; make test {check}; make install



# Why modules?

- ★ Modules were initially developed at OSC, in TCL, then Lua modules (Lmod) at TACC.
  - Most HPC centres use either TCL Modules or Lmod. CC & Grex are Lmod
- **★** Why modules?
  - Control different versions of the same program.
  - Avoid conflicts between different versions and libraries.
  - Set the right path to each program or library.
- **★** How it works?
  - module commands dynamically change the Environment
  - Variables like PATH, LD LIBRARY PATH are appended or prepended
  - Variables like GAUSS EXEDIR or ANSYS HOME are set and unset



# Why modules?



### **Most frequently used Lmod module commands:**

- module list; module avail # shows loaded and available items
- o module spider; module spider <soft>/<version> # deep search
- module show soft/version; # shows what module does
- module help <soft>/<version>; module whatis <soft>/<version>
- module load soft/version; module unload {rm} <soft>/<version>
- module purge; module --force purge # mass unload
- module use ~/modulefiles; module unuse ~/modulefiles



# Why Lmod modules?



### Lmod supports a "hierarchical module structure"

- As modules grow in number, complexity rises. So one can end up having conflicts between modules and their dependencies..
- In the Hierarchical structure, module form a dynamic tree structure based on their dependencies (toolchains like Compiler, MPI, CUDA kinds)



#### How it works?

- module commands dynamically change the MODULEPATH Environment
- Each toolchain has its own module path, preventing cross-toolchain loads
- o module avail only uses the current toolchain's MODULEPATH.
- A new command module spider was added to Lmod to search across all of the hierarchy
- Switching toolchain components causes automatic module reloads, if works



### **Modules on Grex**

- ★ About 450 modules:
  - o GCC [5,7,9]; Intel [2014 2020].
  - o Libraries: HDF5, PETSc, GSL, MKL, Libxc, Boost, ...
  - Gaussian, ANSYS, MATLAB, VASP, MCR, Java, ... etc.
  - LAMMPS, GROMACS, ABINIT, QE, VMD, Molden, ... etc.
- ★ Software maintenance on Grex:
  - We install programs on request from users.
  - Search for a program using "module spider <name of your program>"
  - If not installed, ask for support "support@computecanada.ca"
  - We will install the module 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.



# Modules and slurm jobs

- **sbatch**, **salloc**, **srun** would propagate the environment from the submitting shell. Including loaded modules.
- But we recommend to add module commands to the job scripts explicitly, for it will be easier to track changes and troubleshoot.



## **Software stacks on Grex**

- ★ Grex environment [default]: GrexEnv
  - no module loaded by default, except the GrexEnv
  - use module spider to search for modules
  - Compilers {GCC, Intel}, MKL, PETSc, ... etc.
  - Commercial UM-licensed soft: Gaussian, ANSYS, MATLAB, ... etc.
- ★ Compute Canada environment [optional]: CCEnv
  - Switch to CCEnv; load a standard environment; choose the architecture[sse3, avx2, avx512], use module spider <soft>

module load CCEnv module load StdEnv/2016 module load arch/sse3 module load nixpkgs/16.09 gcc/5.4.0 geant4/10.05.p01

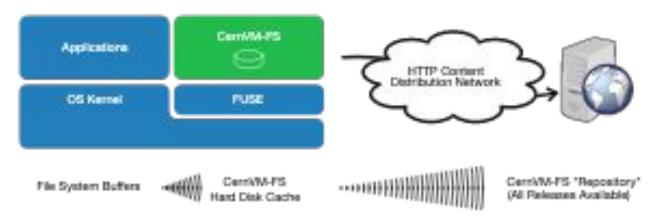


### **CVMFS Software stacks on Grex**

- ★ Compute Canada environment [optional]: CCEnv
  - Tries to be independent of base OS, self contained
  - Encapsulates low level OS libs, own LMOD, and has several HPC toolchains. (GCC, Intel)
  - Commercial software on "restricted" repo
  - Innovative, a model for european stack etc.
- ★ OpenScience Grid
  - Singularity containers, or software distros
- ★ MUGIC project for Genomics



## **CC CVMFS Software Stack**



Picture from https://cvmfs.readthedocs.io/en/stable/cpt-overview.html

- CVMFS filesystem for distribution (CC maintains Stratum 0, Stratum 1)
- A base-OS layer included in the distribution and makes the stack almost self-contained software
- HPC software stack is managed by Lmod and Easybuild



## **CC CVMFS Software Stack**



User layer: Python packages, Perl and R modules, home made codes, ...

**Easybuild layer:** modules for Intel, PGI, OpenMPI, CUDA, MKL, high-level applications. Multiple architectures (sse3, avx, avx2, avx512)

Nix or gentoo layers: GNU libc, autotools, make, bash, cat, ls, awk, grep, etc.

**Gray area:** Slurm, Lustre client libraries, IB/OmniPath/InfiniPath client libraries (all dependencies of OpenMPI) in Nix {or gentoo} layer, but can be overridden using PATH & LD\_LIBRARY\_PATH.

OS: kernel, daemons, drivers, libcuda, anything privileged (e.g. the sudo command): always local.



## **CC CVMFS Local installations**

- ★ R packages: ComputeCanada provide a minimal installation of:
  - o R as modules: users can install the packages in their home directory.
- ★ Python as modules: python and scipy-stack
  - users can install the packages needed in their home directory.
  - Wheels for many popular packages are provided on CC CVMFS.
- ★ Perl and bioperl as modules:
  - users can install the packages needed in their home directory.
- ★ Other software installed locally:
  - Home made programs
  - Restricted and licensed software that can not be distributed via CVMFS.
  - Custom software: patch from a user, changing parts of the code, ... etc.
     <a href="https://docs.computecanada.ca/wiki/Installing">https://docs.computecanada.ca/wiki/Installing</a> software in your home directory



### Local software installations: when?

- Local installation (user's directory):
  - R packages
  - Python packages: virtual environment, conda
  - Perl modules
- Installation from sources with:
  - make; make test {check}; make install
  - Autotools: configure; make test {check}; make install
  - CMake: ccmake; make test {check}; make install
- Java applications: jar files
- Singularity:
  - build the image and run the program from the container



## The question for local installs:

- ★ Are parts/dependences already present on the systems?
- ★ Before building every dependency, check if they already are present
  - Especially low-level libraries like MPI, BLAS/LAPACK/FFT
  - MPI might have non-obvious choices for particular hardware
  - Follow the stack: HPC machine provides base toolchain (Compilers, MPI, CUDA, HDF5/NetCDF, etc.);
  - Dont try re-doing everything; instead build just the top level (application, specific dependencies that are missing).
- ★ Decisions might require knowledge of the particular HPC system
  - If unsure, don't hesitate to contact an HPC analyst



## **Notes on GrexEnv Toolchains**

- ★ Systems compilers and interpreters like CC, CXX, F90, PYTHON etc. not necessarily are the compilers to be used.
- ★ Toolchains: we support mainly Intel (for speed) and GCC (for stable, standard compliant for C++). Intel 2017-2020; GCC 4.8, 7.4, 9.2
- ★ CPU architecture: -msse4.2 (old nodes), -mavx2 (new nodes)
  - For Intel, can do dispatch: -axCORE-AVX512,CORE-AVX2,SSE4.2
  - -xHost easy but dangerous (depends on which node you build!)
- ★ MessagePassingInterface (MPI) libraries:
  - OpenMPI 3.1, 4.0 are recommended (there are older vers. Like 1.6.5, we keep them for compatibilit).
  - Intel MPI 2017-2019 are also provided.



# Java applications

- Download and unpack the code
- ★ Load java module: module load java
- ★ Run the code
- **★** Example: Trimmomatic
  - wget <a href="http://www.usadellab.org/cms/uploads/supplementary/Trimmomatic/Trimmomatic-0.39.zip">http://www.usadellab.org/cms/uploads/supplementary/Trimmomatic/Trimmomatic-0.39.zip</a>
  - unzip Trimmomatic-0.39.zip
- ★ Run the code module load java java -jar <path to>/trimmomatic-0.39.jar {+options if any}



## Local installation: Python/Virtualenv

Example: install Numba package that requires LLVM which is not on GrexEnv Switch to CCEnv and load Pyton module:

```
$> module load CCEnv arch/avx2 # note that this needs skylake!
$> module load StdEnv/2016.4
```

\$> module load python

#### Create a virtual environment "numbaeny" and install Numba:

```
$> virtualenv numbaenv
$> source numbaenv/bin/activate
(numbaenv) $> pip install numba
(numbaenv) $> python -c "import numba"
```

Needs the same virtual environment activated in job scripts!



# Local installation: R packages

```
R packages: rgdal, adegenet, stats, rjags, dplyr, ... etc.
Choose your module: module spider r
Load R and dependencies (mkl, gdal, jags, gsl, udunits... etc):
    module load gcc/7.3.0 r/3.6.0 gdal udunits...
Launch R and install the packages:
    ~$ R
    > install.packages("sp")
    '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 \sim /R/\{...\} to install packages into? (yes/No/cancel) yes
    --- Please select a CRAN mirror for use in this session ---
    > install.packages("dplyr")
```



## Local installation: perl

Example: Hash::Merge; Logger::Simple; MCE::Mutex; threads ...

Load Perl module: module load perl

Install the the first package using cpan:

~\$ 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:

~\$ cpan install Hash::Merge

~\$ cpan install Logger::Simple

~\$ cpan install MCE::Mutex



### Installation with make: STAR

- ★ Download the code:
  - wget https://github.com/alexdobin/STAR/archive/refs/tags/2.7.8a.tar.gz
- ★ Unpack the code: tar -xvf 2.7.8a.tar.gz
- ★ Load GCC compiler: module load gcc
- Compile the code:
  cd STAR-2.7.8a/source
  make
- ★ Copy the binaries and set the path:
  - mkdir -p ~/software/star/2.7.8a/bin cp STAR ~/software/star/2.7.8a/bin
  - export PATH=\$PATH:~/software/star/2.7.8a/bin



# Adding a local module

Create a file **star-2.7.8a** under **\$HOME/modulefiles** with the following:

#%Module1.0

module-whatis "STAR version 2.7.8a"

prepend-path PATH /home/username/software/star/2.7.8a/bin

- \$> module use \$HOME/modulefiles
- \$> module avail # should list /home/username/modulefiles and star-2.7.8a
- \$> module load star-2.7.8a # loads the local module
- \$> which STAR



### **Installation with GNU Autotools**

- ★ Download and unpack the code: wget, ... gunzip, ... etc.
- ★ Load the modules and dependencies: module load gcc ompi fftw
- ★ Configure the program
  - If configure not included, run: autoreconf -fvi [to generate it].
  - ./configure --help [to see the different options].
  - ./configure --prefix=installdir {+other options}
- ★ Compile and test:
  - make
  - make check
- ★ Install the program:
  - make install



# Notes on build dependencies

- ★ Most of the time they are modules already present on the systems
- ★ Can be often autodetected in CPATH, FPATH, INCLUDE, LIBRARY\_PATH. HDF5\_HOME; If not, manually add the options.
- ★ Names: BLAS/LAPACK/FFTW are part of Intel MKL libraries (now OneAPI). Just load the mkl module! Intel has calculator for Include options if not autodetected.
- ★ CPU architecture: -msse4.2 (old nodes), -mavx2 (new nodes)
  - For Intel, can do dispatch: -axCORE-AVX512,CORE-AVX2,SSE4.2
  - -xHost easy but dangerous (depends on which node you build!)
  - Compatibility vs. speed. CCEnv has "modules" for arch/, GrexEnv doesn't.



# **Example: options for PETSc**

```
./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=${fftsrc} --download-amd=yes --download-adifor=yes --download-superlu=yes
--download-triangle=yes --download-generator=yes --with-64-bit-pointers=no --with-cc=mpicc
--CFLAGS='-O2 -msse4.2 -xSSE4.2 -mp1 -I$MKLROOT/include -mkl -fPIC ' --with-cxx='mpicxx'
--CXXFLAGS='-O2 -msse4.2 -xSSE4.2 -mp1 -I$MKLROOT/include -mkl -std=c++11 -fPIC '
--with-fc='mpif90' --FFLAGS='-O2 -msse4.2 -xSSE4.2 -mp1 -I$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
```



## **Example with Cmake**

- ★ Download and unpack the code: wget, ... gunzip, ... etc.
- ★ Load the modules and dependencies: module load gcc ompi fftw
- ★ Configure the program: you may need to load cmake module
  - mkdir build && cd build
  - cmake .. --help [to see the different options]. Or ccmake ..
  - cmake .. -DCMAKE\_INSTALL\_PREFIX=installdir {+other options}
- ★ Compile and test:
  - make
  - make check; make test
- ★ Install the program:
  - make install



## **Cmake options for GROMACS**

module load intel/2019.5 module load ompi/3.1.4 fftw module load cmake

```
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
```



# Notes on installing binaries

- ★ Usually, use tarballs rather that packages (yum, deb, etc.) as there is no sudo access.
- ★ Dependencies might be there as modules, or parts of modules (libstdc++ comes from newer GCC, etc.).
- ★ Install under \$HOME, modify path
  - One standard way is to create a local module
  - Module use ~/modulefiles
  - Module avail; module load ..
- ★ GLIBC dependency missing: no chance, recompile the code from sources or use containerized deployment.
- ★ CPU architecture is different: no chance at all. Esp. for old compute



# **Singularity**

Resources: Github, DockerHub, SingularityHub.

Singularity examples: <a href="https://github.com/singularityware/singularity/tree/master/examples">https://github.com/singularityware/singularity/tree/master/examples</a>

- Documentation: <a href="http://singularity.lbl.gov/user-guide">http://singularity.lbl.gov/user-guide</a>
- DockerHub: https://hub.docker.com/explore/
- SingularityHub: <a href="https://www.singularity-hub.org/">https://www.singularity-hub.org/</a>

### **Access to Singularity:**

- **Connect to cluster:** Grex, cedar, graham or beluga:
- ★ Load a module: module load singularity
- ★ Build the image: convert the image from Docker to Singularity
- Note: You may need to use your own Linux machine or VM to build the image

https://docs.computecanada.ca/wiki/Singularity
https://monitor.hpc.umanitoba.ca/doc/docs/grex/software/containers/



# Singularity

- ★ Alternative for running software: difficult to build from source
- ★ Possibilite to convert Docker images to singularity.
- ★ Singularity installed on all clusters {no Docker for security reasons}
- ★ Build the image:

module load singularity singularity build qiime2-2019.10.sif docker://qiime2/core:2019.10

- ★ Run the code via singularity:
  - singularity exec -B \$PWD:/home -B /global/scratch/someuser:/outputs \
  - -B /global/scratch/someuser/path/to/inputs:/inputs qiime2-2019.10.sif \ qiime feature-classifier fit-classifier-naive-bayes \
  - --i-reference-reads /outputs/some\_output\_feature.qza \
  - --i-reference-taxonomy /outputs/some\_output\_ref-taxonomy.qza \
  - --o-classifier /outputs/some\_output\_classifier.qza



### Links

- ★ FAQ: <a href="https://docs.computecanada.ca/wiki/Frequently\_Asked\_Questions">https://docs.computecanada.ca/wiki/Frequently\_Asked\_Questions</a>
- ★ Jobs:
  - o https://docs.computecanada.ca/wiki/Running jobs
  - o <a href="https://docs.computecanada.ca/wiki/Job-scheduling-policies#Percentage-of-the-nodes-you-have-access-to-">https://docs.computecanada.ca/wiki/Job-scheduling-policies#Percentage-of-the-nodes-you-have-access-to-</a>
  - https://docs.computecanada.ca/wiki/Advanced MPI scheduling#Whole nodes
  - https://docs.computecanada.ca/wiki/Using GPUs with Slurm

#### **★** Storage:

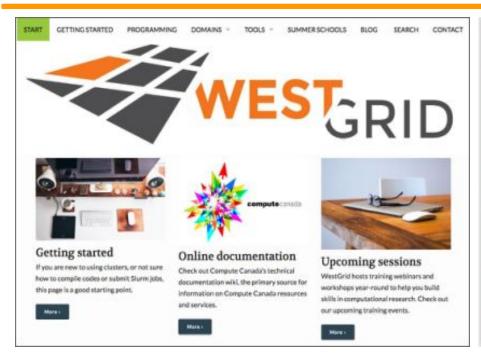
- <a href="https://docs.computecanada.ca/wiki/Storage">https://docs.computecanada.ca/wiki/Storage</a> and file management#Filesystem Quotas and Policies
- https://docs.computecanada.ca/wiki/Project\_layout?
- https://docs.computecanada.ca/wiki/Transferring\_data

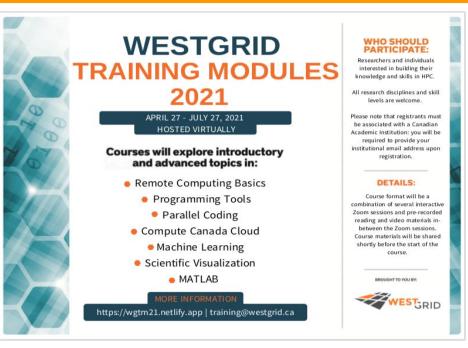
#### ★ Software:

- https://docs.computecanada.ca/wiki/Available\_software
- o https://docs.computecanada.ca/wiki/Utiliser des modules/en
- https://docs.computecanada.ca/wiki/Installing software in your home directory
- https://docs.computecanada.ca/wiki/Python
- https://docs.computecanada.ca/wiki/R



# **Upcoming WestGrid trainings**





https://westgrid.github.io/trainingMaterials https://www.westgrid.ca/events