Finding, downloading, and applying software on CURC resources

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Slides & Exercises

https://github.com/ResearchComp uting/hpc_fundamentals_micro_cre dential

In "installing_software" directory





Learning Objectives

 Learn about different methods to install and use software on Alpine



Session Overview

- The Module System (Lmod)
- Building Software on CURC Systems
- Simplifying Source Installations with Spack
- Virtual Environments with Anaconda
- Containerization with Apptainer
- Requesting Software Installations





In most cases, a supercomputer has far more software installed than the average user will ever use.

- Users may need different versions of the same software, which in general cannot be installed nor used in parallel on the same system.
- The requirements for one package may adversely affect another package or even be mutually exclusive.





HPC centers manage this complexity with environment module systems.

CURC uses the Lmod system.





Setting up for today's session.

- 1. Log in to CURC HPC system
- 2. Get on an Alpine compute node

```
$ module avail
$ acompile --help
$ acompile --time=2:00:00
$ module avail
```

Note: Login nodes do not have the full software stack!





Live Demo: Loading and unloading modules will dynamically change the software environment on the cluster.

```
$ module purge
$ module load intel/2022.1.2
$ module avail
$ module load impi
$ module avail
```

```
$ module purge
$ module load gcc
$ module avail
$ module load openmpi
$ module avail
```





Live Demo: Loading and unloading modules will set (and reset) important environment variables for you.

```
$ module purge
$ module load intel
$ module load hdf5
$ module display hdf5
$ env | grep HDF5
```

```
$ module purge
$ module load gcc
$ module load hdf5
$ module display hdf5
$ env | grep HDF5
```





Live Demo: Useful Lmod commands

```
module spider
                                     # list all available modules
module avail
                                     # list modules available to you
module load <package/version>
                                     # load a module into your env
module purge
                                     # unload all modules
module list
                                     # list currently loaded modules
module display <package>
                                     # display module info/help
                                     # view info for all version
module spider <package>
module spider <package/version>
                                     # view info for specific version
```





Points to note about CURC-managed modules:

- CURC does not update system modules; we do fresh installs of new versions and change the default when that is appropriate
- Sometimes when a module is outdated or problematic we will remove it from the software stack

Take home: pay attention to what modules you are loading, as this may be important for reproducibility!



- Definitions
 - Building- a generic term describing the overall installation process that includes compiling
 - Compiling- the process of converting source code to an executable
 - Linking- the process of combining pieces of code and data into a single file that can be loaded into memory and executed
 - Installing- any process that results in executables



- There are numerous ways to install software on CURC systems
 - Grab pre-compiled binaries
 - Within virtual environments (using Conda, Miniconda, or Mamba)
 - Using containers (Apptainer)
 - From source code
 - Using a package manager for HPC systems (Spack)



Why compile a research application manually from source code?

- It is not distributed as a pre-compiled binary, by any package manager, and is not easily containerized.
- 2. Compiling from source on the cluster will greatly improve performance.



Compilers are programs that convert code written in high level programming languages (like C/C++ or Fortran) to executable binary files.





Build systems automate the process of compiling and linking.

- 1. GNU Build System
 - your application includes instructions to run

 /bootstrap, ./autogen.sh, ./configure or make (the latter without a preceding cmake)
 - make is available in /usr/bin; Autotools available as a module

```
$ ./configure --prefix=/projects/$USER/software/bin
$ make
$ make install
```



Build systems automate the process of compiling and linking.

2. Cmake

- your application includes a cmake step
- module avail cmake



Conventions and best practices

- You will need to adapt installations for local or user installations (look for these terms in the software's docs)
- Don't install software in /home/\$USER (too small) or scratch (purged every 90 days); /projects/\$USER/software is the way to go!
- Keep your software installations organized by using a consistent file structure and naming convention
- Load the compiler first, MPI implementation second, and thirdparty libraries last



Conventions and best practices

- Don't install executables to the source directory
 - cmake -DCMAKE_INSTALL_PREFIX, ./configure --prefix
- The newest version of a compiler might not be compatible with your application. Read the package documentation and don't be afraid to try different compilers and compiler versions
- Read our 'Compiling and Linking' documentation https://curc.readthedocs.io/en/latest/compute/compiling.html



Conventions and best practices

 Make life easier for yourself by adding executables to PATH and any directories with libraries that your application links to LD_LIBRARY_PATH

```
$ export PATH=/projects/$USER/software/phyloflash/bin:$PATH
$ echo $PATH
$ export
    LD LIBRARY PATH=/curc/sw/hdf5/1.10.1/impi/17.3/intel/17.4/lib
:$LD_LIBRARY_PATH
$ echo $LD_LIBRARY_PATH
```



Want to go the extra mile?

Try our Hands-on exercise #1 provided in EXERCISES.pdf. This is not required for the micro-credential.

Objectives:

- 1) Explore CURC compilers and compiler environment variables.
- 2) Perform a simple source installation.

Estimated time to complete: 15 minutes





How can we simplify source installations?

- Package Managers Tools that automate installing, maintaining, and configuring software and any dependencies
- Environments A collection of resources that are available in a self-contained 'bubble'



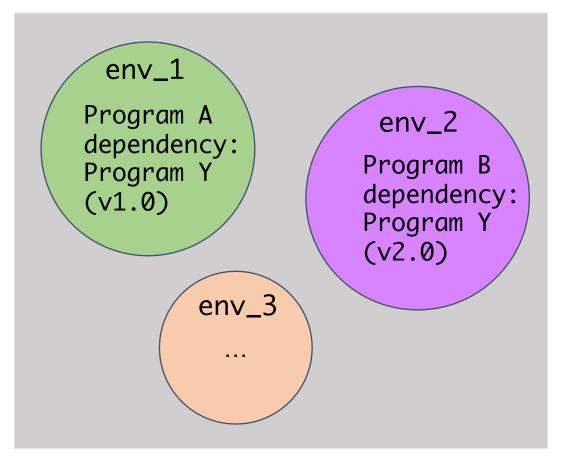


Think of virtual environments as selfcontained bubbles.

env_1 contains all the dependencies of 'Program A'.

env_2 contains all the dependencies of 'Program B'.

The environments do not interact.



Your workflow requires two programs, 'Program A' and 'Program B'.

- 'Program A' depends on 'Program Y' v1.0
- 'Program B' depends on 'Program Y' v2.0

What do you do?!





Environments are created and programs are installed in a few simple steps

```
$ module purge
$ module load spack/0.20.1
 spack env create my first env
$ spacktivate my first env
$ spack install --add samtools
```

Warning: Don't install packages outside of an environment!





 Packages are installed within activated environments

using spack install

```
$ spack install --add samtools
                                    #install default samtools
$ spack install --add samtools@1.9
                                    #install specific version
```





- Spack installations can be slow but will progress more quickly with more cores.
 - Spack builds all packages in parallel. The default parallelism is equal to the number of cores available to the process, up to 16.





Useful spack commands

```
spack env list
                                     # list all your environments
                                     # remove an environment
spack remove <env>
spack uninstall <packagename>
                                     # remove package
spack env status
                                     # check which env you're in
                                     # prints detailed package info
spack info <packagename>
spack find
                                     # show installed packages
                                     # deactivate environment
despacktivate
spack spec <packagename>
                                     # list packages plan
```





Useful spack file paths

```
# root of the spack install tree
/projects/$USER/software/spack
# location of package executables - these are symbolically linked to
      the installation tree subdirectory
/projects/$USER/spack/environments/<env>/.spack-env/view/bin
# location of spack config file
/home/$USER/.spack/config.yaml
```



Want to go the extra mile?

Try our Hands-on exercise #2 provided in EXERCISES.pdf. This is not required for the micro-credential.

Objectives:

- 1) Create a Spack environment
- 2) Install fastqc in your Spack environment

Estimated time to complete: 20 minutes **Documentation**:

https://curc.readthedocs.io/en/latest/software/spack.html





Conda is a package (software) management system

- Installs, runs, and updates packages and their dependencies
- Creates, saves, loads, and switches between virtual environments
- Created for Python programs, but can package and distribute software for any language





For our system, please be sure to add a .condarc file in your home directory.

Prevents package installs from going to your home directory

```
$ cd ~
$ nano .condarc
Add the following content:
pkgs_dirs:
  - /projects/$USER/.conda_pkgs
envs dirs:
  /projects/$USER/software/anaconda/envs
```





Environments are created and programs are installed in a few simple steps

```
$ module load anaconda
 conda create -n my first env python==3.10
$ conda activate my first env
$ python
```

Warning: Don't install packages in your base environment!





Packages are installed within activated environments

using conda install (preferred method, when available)

```
$ conda install pandas
                               #install latest pandas
$ conda install pandas==0.20.3 #install specific version of pandas
```





Packages are installed within activated environments

using pip install (if you must)

```
$ pip install --no-cache-dir pandas
                                          #install latest pandas
```

Warning: --no-cache-dir is crucial on CURC systems!





Useful conda commands

```
conda env list
                                    # list all environments
conda list
                                    # list packages in active env
conda env remove -n <envname>
                                    # remove an environment
conda config --show channels
                                    # view configured channels
conda deactivate
                                    # deactivate environment
conda create --name <clonedenv> /
                                   # clone an environment
      --clone <envtoclone>
```





Useful **conda** file paths

```
# location of python libraries
/projects/$USER/software/<env>/lib/python3.10/site-packages
 location of package executables
/projects/$USER/software/<env>/bin
 location of .condarc file
/home/$USER/.condarc
```



Want to go the extra mile?

Try our Hands-on exercise #3 provided in EXERCISES.pdf. This is not required for the micro-credential.

Objectives:

- 1) Configure your .condarc file
- 2) Create a conda environment and install samtools
- 3) Activate the environment and run samtools.

Estimated time to complete: 15 minutes **Documentation**:

https://curc.readthedocs.io/en/latest/software/python.html





Containers are portable virtualizations of an operating system, software, libraries, data, and/or workflows

- pros
 - portability- containers can run on any system equipped with its specified container manager
 - reproducibility- because containers are instances of prebuilt isolated software, the software will always execute the same every time
- cons
 - steeper learning curve than conda
 - can be difficult to troubleshoot issues
 - building containers can be tricky for multi-node MPI applications





CURC offers Apptainer (formerly Singularity) as container management software

• Apptainer comes pre-installed on all Alpine nodes, so no need to load any specific software

Many common research applications have already been containerized and can be pulled from container repositories (such as Docker Hub).



Containerization With



Useful Apptainer commands

```
apptainer exec #Execute a command to your container
apptainer run #Run your image as an executable
apptainer build #Build a container
apptainer pull #pull an image from hub
apptainer inspect #See labels/environment vars, run scripts
apptainer shell #Shell into your image
```





A container has its own file system and so needs help "seeing" files outside the container (on the host system). If not done in the .def file, this can be accomplished at runtime with bind mounting.

```
# bind mount a directory
apptainer run -B /source/directory:/target/directory sample-image.sif
```

On CURC systems, a running container automatically bind mounts these paths: /home/\$USER, \$PWD. Note that other locations will need to be manually mounted.



Want to go the extra mile?

• Try our Hands-on exercise #4 provided in EXERCISES.pdf. This is not required for the micro-credential.

Objectives:

- 1) Become familiar with basic Apptainer commands.
- 2) Pull an image from a pre-built container, then run the program from the container.

Estimated time to complete: 20 minutes

Documentation:

https://curc.readthedocs.io/en/latest/software/Containerization on.html



Requesting Software Installations

- Is the software already installed on the cluster? https://curc.readthedocs.io/en/latest/clusters/alpine/soft ware.html
- Have you considered its utility and complexity?
 - Are you the only user of this software?
 - How complex or difficult is this software to install?
- Have you tried installing the package on your own?
- Software request form: <u>https://www.colorado.edu/rc/userservices/software-request</u>



Thank you!

