Installing Software and Writing Modules

Introduction

In the intro portion of the workshop you will learn:

- About downloading code
- About compiling code
- How to build a package from source code
 - configure
 - build
 - install
- What are yum, rpm, get-apt, & sudo
- How to write modules

Downloading Code

Source vs Executable

- In most cases you are better off downloading the source and building the code (aka the executable) yourself.
- Downloading an executable is easier but likely will not to work.

Downloading Executables

Some developers provide pre-built executables of their software. There are instances when available executables will run flawlessly on Hydra, but make sure that:

- 1 you can trust the origin,
- 2 you get a version compatible with Hydra,
 - i.e., CentOS 7.x for Intel/AMD CPUs (x86_64)

Remember

- Hydra configuration is specific:
 - pre-built code may need *stuff* (dependencies) not on Hydra.



Notes on Downloading Executables

Risks

- Since users on Hydra do not have elevated privileges (root access) you are very unlikely to damage the cluster, but malicious software can still damage your files.
- In rare cases it may install a Trojan horse that could exploit a known vulnerability.
 - Be vigilant and responsible.
 - In case of doubt, never hesitate to contact us.

Compiling code

Steps

Creating executable from source code is typically done as follows:

- 1 compile the source file(s) to produce object file(s),
- 2 link the object file(s) and libraries into an executable.

In Practice

- Often aided by a makefile,
- Configuring is creating such makefile or equivalent.

This will be illustrated in the hands on section.

Building from Source

1. Configure

- Most packages come with a configuration script, a list of prerequisites (dependencies/libraries) and instructions,
- Some packages allow you to build the code without some features in case you cannot satisfy some of the prerequisites,
- You most likely need to load the right module(s) to use the appropriate tools (compilers).
- The configuration step will test if the code can be built:
 - check dependencies, versions, etc.
 - if this fails, the code cannot be built as is.

1.b Makefile only

- Other (simpler) packages come with a makefile that needs to be edited,
 - check the instructions.



Building from Source (cont'd)

2. Build

- make sure you have loaded the right modules,
- run make to compile and link (aka build) the code.

2.b Test

some packages come with the optional step of testing the built before installing it, using something like make test.

3. Install

- copy the executable(s) to the right place(s),
 - usually defined by the configuration,
- best practice is to separate build from install locations.



Setting up Your Environment to Run Your Code

Likely Needed

You likely will need to adjust your environment to run some code:

- 1 the location of the code: path or PATH,
- 2 the location of the libraries: LD_LIBRARY_PATH,
- 3 you may also need to set some environment variables, etc.

Easier Way: modules

This is where using a module makes things easy:

- compact, and
- works with any shell.

The yum, rpm, get-apt and sudo Soup

Definitions

- yum: is a package-management utility for CentOS
- rpm: pre-built software package
 - both are for sys-admin,
 - help handle dependencies,
 - yet ...
- get-apt: Debian's version of yum, does not work on CentOS.

Also

sudo: allows to run a command as 'root': you can't!

BTW

- Instructions that mention yum, rpm, apt-get or sudo
 - will not work on Hydra,
 - **yet** in most cases there is another way.



How about Hydra

Using yum

- While you cannot install packages with yum,
- you can check if we've installed a prerequisite package

In practice

• if the instructions say
sudo yum -y install <package>

you can run

yum info <package>

Using yum info

Example

```
yum info libXt-devel
... stuff and may be slow the first time ...
Installed Packages
Name
    : libXt-devel
Arch : x86_64
Version: 1.1.5
. . .
Description: X.Org X11 libXt development package
. . .
    You want the Arch: x86 64 to be listed as "Installed"
    not just "Available".
```

How to avoid sudo

sudo make install

- if the instructions says
- sudo make install
 - instead, set the installation directory to be under your control,
 - in most cases at the configuration step
- ./configure -prefix=/home/username/big-package/3.5
 - and use

make install

Final Notes

Remember

- there is a way to use yum as a non privileged user
 - not recommended, unless you're an expert!
- you can always ask about a missing prerequisite,
- most of those can be build from source since Linux is an open source OS.

Module and Module Files

The Command module

- convenient mechanism to configure your environment,
- reads a file, the module file, that holds instructions,
- a shell independent way to configure your environment:
 - same module file whether sh/bash or csh/tcsh.

Examples

- We provide module files, users can write their own.
 - look at all the module files we wrote,
 - they can be found in /share/apps/modulefiles/

Module File Syntax and Concepts

Special Instructions

Instructions to configure your environment:

prepend-path PATH /location/of/the/code

setenv BASE /scratch/demo

set-alias crunch "crunch --with-that-option *"

Syntax

- Module files can be complex, using tcl language
 - you **do not** need to know tcl to write module files.

Simple or Complex

- A simple module file can just list the modules that must be loaded to run some analysis.
- Can write complex module files and leverage tcl.



Example of module Commands

Basic

	Info		Config	Details
module module			load unload	list help <name></name>
module	whatis	<name></name>	swap	show <name></name>

More help

man module

A Simple Module File

Example

```
#%Module1.0
#
# load two modules and set the HEASOFT env variable
module load gcc/10.1.10
module load python/3.8
setenv HEASOFT /home/username/heasoft/6.3.1
```

Example of More Elaborate and Complex Module Files

Will be illustrated in the hands on section.

Module Files Organization

Recommended Approach

- use a central location under you home directory
 ~/modulefiles,
- use a tree structure
- use version numbers if/when applicable,
- let module know where to find the module files.

Customization/Examples

Tree structure

- ~/modulefiles/crunch/
- ~/modulefiles/crunch/1.0
- ~/modulefiles/crunch/1.2
- ~/modulefiles/crunch/2.1
- ~/modulefiles/crunch/.version
- ~/modulefiles/viewit

Define a Default Version

An optional file .version can be used to set the default version:

```
#%Module1.0
```

set ModulesVersion "1.2"

Hence

module load crunch module swap crunch/2.1



Customization (cont'd)

Let module Know Where to Find the Module Files

```
module use --append ~/modulefiles
```

Either

- in your initialization file ~/.bashrc or ~/.cshrc
- 2 or better yet in a ~/.modulerc file

```
#%Module1.0
# adding my own module files
module use --append /home/username/modulefiles
```



Hands-on Section

Hands-On

In the hands-on portion of the workshop you will

- Build and install software using best-practices,
 - trivial case,
 - simple/didactic example,
 - somewhat complex examples.
- Write simple and more elaborate module files.
- Run the software you installed in jobs.

But first, log in to Hydra

- If you need a reminder about how to log into Hydra and how to change your password, check the *Intro to Hydra* tutorial.
 - If the link does not work:

https://github.com/SmithsonianWorkshops

- > Hydra-introduction
 - > hydra_intro.md

Switch to github for the Hands-on

Go to

https://github.com/SmithsonianWorkshops/advanced-hydra-workshops/

Convention

- I use % as prompt
 - your prompt might be different, like \$
 - you type what is after the prompt
 - no prompt: result from previous command.

But First

Create a location where to run things

- For SAO (CfA) ppl
- % cd /pool/sao/\$USER
- % mkdir -p advanced-workshop/sw+m
- % cd advanced-workshop/sw+m
 - For others (biologists+)
- % cd /pool/genomics/\$USER
- % mkdir -p advanced-workshop/sw+m/hands-on
- % cd advanced-workshop/sw+m/hands-on
 - \$USER will be replaced by your user name,
 - feel free to put this elsewhere.

Exercise 1

Install a simple prebuilt executable: rclone

Create a directory

% mkdir ex01

```
% cd ex01
 2 Get rclone
 ■ Google "download rclone linux" -> https://rclone.org/install/
% wget https://downloads.rclone.org/rclone-current-linux-am
--2023-03-14 14:20:17-- https://downloads.rclone.org/rclone.
Resolving downloads.rclone.org (downloads.rclone.org)... 9
Connecting to downloads.rclone.org (downloads.rclone.org) |
HTTP request sent, awaiting response... 200 OK
Length: 17790831 (17M) [application/zip]
Saving to: 'rclone-current-linux-amd64.zip'
100%[=========]] 17
```

2023-03-14 14:20:21 (5.54 MB/s) - 'rclone-current-linux-amo

Exercise 2

Compiling a trivial program

```
Let's build from source a very very very simple code:
 1 create a directory and copy the source file
% cd ..
% mkdir ex02
% cd ex02
% cp -pi /pool/sao/hpc/aw/ex02/hello.c ./
 2 look at the code
% cat hello.c
#include <stdio.h>
#include <stdlib.h>
/* simple hello world demo code in C */
int main () {
  printf ("hello world!\nEasy peasy ;-P\n");
  exit(0):
}
```

Excercise 3

- Similar simple code
 - but let's use a makefile and
 - a different compiler by loading a module
- create a directory and copy the source and makefile files

```
% cd ..
% mkdir ex03
% cd ex03
% cp -pi /pool/sao/hpc/aw/ex03/hello.c ./
% cp -pi /pool/sao/hpc/aw/ex03/makefile ./
2 look at the files
```

% more hello.c makefile
....

3 load the Intel compiler, build and run it

Build a Bio Package

Build an Astro Package

Write a More Elaborate Module File

The rclone/1.62.0 module file for your private version

```
create a file with the following content:
#%Module1.0
#
# set some internal variables
set ver 1.62.0
set base /pool/<genomics|sao>/username>/advanced-worksho
#
# what to show for 'module whatis'
module-whatis "System paths to run rclone $ver"
#
# configure the PATH and the MANPATH
prepend-path MANPATH $base/rclone/$ver/man
 where?
 1 under the ex01/ directory
```

How to use it:

% module unload tools/rclone

```
% module load /pool/<genomics|sao>/$USER/advanced-workshop/
% which rclone
/pool/<genomics|sao>/$USER/advanced-workshop/sw+m/ex01/bin/
% module unload /pool/<genomics|sao>/$USER/advanced-workshop/
% module load tools/rclone
% which rclone
/share/apps/bioinformatics/rclone/1.53.1/rclone
```

Using Instead a Central Location

```
% cd ~
% mkdir modulefiles
% mkdir modulefiles/rclone
% cp -pi /pool/sao/hpc/aw/ex01/modulefiles/rclone/1.62.0 mc
% cd -
% module use --append ~/modulefiles
% module load rclone/1.62.0
% which rclone
% module unload rclone
% module load tools/rclone
% which rclone
% module unload tools/rclone
```

Make it permanent

```
% cat <<EOF ~/.modulerc
#%Module1.0
# adding my own module files
module use --append /home/username/modulefiles
EOF</pre>
```

Check this out

- % module whatis rclone
- % module whatis tools/rclone

Examples of Complex Module Files

Plenty of Examples

```
cd /share/apps/modulefiles
```

more intel/2022.2 more idl/8.8 more bio/blast2go/1.5.1 more bio/trinity/2.9.1

Questions?