

FALL 2018



APPLIED ENGINEERING DATA ANALYSIS, OPTIMIZATION AND VISUALIZATION

Getting started with TACC

JOSHUA RHODES, PHD

Research Fellow/Adjunct Professor, The University of Texas at Austin

TACC is the Texas Advanced Computing Center

- Started in 1960s with a CRAY CDC 6600
 - Your phone is more powerful than this thing now
- TACC became TACC in 2001
- Mostly housed at the PRC campus
 - Visualization lab on Main Campus

TACC is available for you as part of UT

- Computing is free
- Your advisor needs to get a project set up
- TACC can be brought on for research projects and as computing support for grants
- Deeper dives into using TACC, by TACC
 - <https://learn.tacc.utexas.edu/>

Logging into TACC

- From Terminal
 - ssh
 <username>@maverick.tacc.utexas.ed
 u
 - Password
 - 2-factor authentication

The first thing you see is your account balance

```
-----, - -----, names marked in green -----,
----- Project balances for user joshdr -----
| Name                Avail SUs    Expires | Name                Avail SUs    Expires |
| ME397M-Applied-Engin  75000  2018-06-30 | WebberEnergy         4497  2018-03-31 |
----- Disk quotas for user joshdr -----
| Disk                Usage (GB)    Limit    %Used    File Usage    Limit    %Used |
| /home                1.0          10.0     10.28     4722        750100    0.63 |
| /work                696.7       1024.0   68.03     123818      3000000   4.13 |
-----
```

\$Home vs. \$Work directory

- \$Home: where you run programs from
 - 10GB storage
- \$Work: where you store your data
 - 1TB storage
- You can access your data in work from home
 - i.e. function running in \$home can read file in \$work

You land on your home folder

- `home/XXXXXX/<tacc_username>`
- The same commands we learned in Lecture 2:Terminal apply here
 - `cd`, `mkdir`, `ls`, `rm`, `rmdir`, etc.

When you first log on, you are on a login node

- YOU CANNOT RUN PROGRAMS ON THE LOGIN NODE – YOU WILL GET KICKED OFF
- You can make folders, move files, edit file, install programs, queue files to run

There is a way to get a compute node all to yourself

- You can create an interactive development session
 - `idev -m 90`
 - `-m 90` = get computer for 90 min, default = 30 min
 - Will kick you off and not save anything at end of time!!

So now that I am in, what do I do?

- You have to load programs (modules) before you can use them
 - Like double-clicking on an icon
- To check what you can run
 - module avail

What modules are available?: module avail

```
login1.maverick(1008)$ module avail
```

----- /opt/apps/intel14/mvapich2_2_0/modulefiles -----					
Rstats/3.0.3	petsc/3.5-complexdebug	petsc/3.5-cxxcomplexdebug	petsc/3.5-uni	python/2.7.6	
paraview/4.3.1	petsc/3.5-cxx	petsc/3.5-cxxdebug	petsc/3.5-unidebug	visit/2.7	
petsc/3.5-complex	petsc/3.5-cxxcomplex	petsc/3.5-debug	petsc/3.5	visit/2.9	(D)
----- /opt/apps/intel14/modulefiles -----					
boost/1.51.0	gs1/1.16	hdf5/1.8.12 (m)	mvapich2/2.0b (L)	nco/4.5.4	netcdf/4.2.1.1
					udunits/2.2.19
----- /opt/apps/xsede/modulefiles -----					
CTSSV4	TERAGRID-paths	cue-comm	globus-5.0	jdk32/1.7.0	tginfo/1.1.3
GLOBUS-5.0	apache-ant/1.6.5	cue-login-env	globus/5.2.5	pacman/3.29	tgproxy/0.9.1
TERAGRID-BASIC	ctssv4	cue-math	gsissh/4.3	teragrid-basic	tgresid/2.3.4
TERAGRID-DEV	cue-build	cue-tg	gx-map/0.5.3.3	teragrid-dev	uberftp/2.6
----- /opt/apps/modulefiles -----					
Rstudio/0.98.501	cuda/7.5	gcc/5.4.0	lmod/7.7.1	pylauncher/2.1	vapor/2.5.0
Rstudio/1.0.153 (D)	cuda/8.0 (g,D)	git/2.7.0	luatools/1.1	qt/4.8.4	vtk/6.1.0
amira/6.0.1	cudnn/4.0	idev/1.0	matlab/2013a	remora/1.7	xalt/1.8 (L)
autotools/1.0	ddt/5.0.1	idl/7.0.6	matlab/2015a	sanitytool/1.3	
cmake/2.8.12.2	ffmpeg/2.1.4	idl/8.4 (D)	matlab/2017a (D)	settarg/7.7.1	
cmake/3.7.1 (D)	gcc/4.7.1	intel/14.0.1.106 (L,D)	mcr/v81	swr/17.0	
cuda/6.5	gcc/4.9.1 (D)	intel/15.0.3	nc1_ncarg/6.3.0	tacc-singularity/2.3.1	
cuda/7.0	gcc/4.9.3	launcher/2.0	ospray/1.3.1	tacc_tips/0.5	
----- /opt/modulefiles -----					

Is that it?

- Check for ALL installed programs
 - module spider
- Check for installed programs that start with R
 - module spider R
 - >

We have multiple versions of R installed and available, v3.4.0 has dependencies (so it d/n show up earlier)

Rstats:

Versions:

Rstats/3.0.3

Rstats/3.4.0

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

\$ module spider Rstats/3.4.0

RstatsPackages: RstatsPackages/3.4.0

You will need to load all module(s) on any one of the lines below before the "RstatsPackages/3.4.0" module is available to load.

intel/15.0.3 mvapich2/2.1

Help:

This is the R statistics (RstatsPackages) package built on October 04, 2017.

It includes the following accessory packages including but not limited to:

Rmpi, snow, snowfall

pdbMPI, pbdSLAP, pbdBASE, pbdDMAT, pbdDEMO, pbdNCDF4, pmclust

multicore

doMC, doSNOW, doMPI, doParallel

BH, bigmemory, biganalytics, bigtabulate, synchronicity

Rdsm, SparseM, slam, cluster, randomForest, bit, ff, mchof

BioConductor (base installation plus some common packages)

ggplot2, rjags/r2jags, rgdal, rstan

The RstatsPackages modulefile extends the PATH and LD_LIBRARY_PATH paths as appropriate.

Version 3.4.0

For the packages we used to build our R homework, we need R v3.4.0

- We must load the dependencies first

```
login1.maverick(1003)$ module load intel/15.0.3 mvapich2/2.1
```

The following have been reloaded with a version change:

1) intel/14.0.1.106 => intel/15.0.3 2) mvapich2/2.0b => mvapich2/2.1

```
login1.maverick(1004)$ module avail
```

```
----- /opt/apps/intel15/mvapich2_2_1/modulefiles -----
Rstats/3.4.0      amber/14.0      fftw3/3.3.4      paraview/5.4.1    python/2.7.9      vtk/7.0.0 (D)
RstatsPackages/3.4.0  fftw2/2.1.5    parallel-netcdf/4.3.3.1  phdf5/1.8.16      visit/2.12

----- /opt/apps/intel15/modulefiles -----
cxx11/4.9.1      gsl/1.16      hdf5/1.8.16      impi/5.0.3      mvapich2/2.1 (L)  netcdf/4.3.3.1  python/2.7.9      python/2.7.13 (D)

----- /opt/apps/xsede/modulefiles -----
CTSSV4          TERAGRID-paths  cue-comm          globus-5.0        jdk32/1.7.0        tginfo/1.1.3      xdusage/1.0
GLOBUS-5.0      apache-ant/1.6.5  cue-login-env     globus/5.2.5      pacman/3.29        tgproxy/0.9.1
TERAGRID-BASIC  ctssv4           cue-math          gsissh/4.3        teragrid-basic     tgresid/2.3.4
TERAGRID-DEV    cue-build        cue-tg            gx-map/0.5.3.3    teragrid-dev       uberftp/2.6
```

Installing your own software

- Here we will install GAMS
 - https://www.gams.com/24.8/docs/userguides/userguide/_u_g_u_n_i_x_i_n

Platform

MS Windows 32 bit

Windows Vista or newer on AMD- or Intel-based (x86_32) architectures.

DOWNLOAD**MS Windows 64 bit**

Windows Vista or newer on AMD- or Intel-based (x86_64) architectures.

DOWNLOAD**Linux 64 bit**

AMD- or Intel-based 64-bit (x86_64) Linux systems with glibc 2.12 or higher.

DOWNLOAD

You can create shortcuts for yourself

- `alias gams=~ /opt/gams/gams25.0_linux_x64_64_sfx/gams`
- `alias`
`gamslib=~ /opt/gams/gams25.0_linux_x64_64_sfx/gamslib`

To use TACC for more than 90 minutes, you will need to submit a job to the queue

- There is a sequence to get in the queue
 - Shell script
 - R script

Aside: you can use VIM to make some quick edits



```
jdr2823 — joshdr@login1.maverick:~/me397 — ssh joshdr@maverick.tacc.utexas.edu
joshdr@login1.maverick:~/me397 — ssh joshdr@maveri...  joshdr@login2.maverick:~/FCe — ssh joshdr@maverick....

##

source('RHODES_WU_HWK1.R')

wdata <- RHODES_WU_HWK1(month = 1, day = 1, year = 2017, station_id = 'KILFRANK2')

head(wdata)

write.csv(wdata, 'wu_output.csv', row.names = F)

~
~
~
~
```

You can use a utility like Cyberduck to make life easier

