

O2 – Tips and Tricks

RC INFORMATIONAL SESSION

Presenters: Research Computing Consultants (RCCs)



HARVARD
MEDICAL SCHOOL

RESEARCH COMPUTING 1

How to know if you're requesting too much RAM or CPUs on your jobs?

O2_jobs_reports:

- Customizable queries allow you to select specific time intervals, partitions, jobs states and more.
- Three available output formats
- Creates a comprehensive report for your jobs or everyone in your Lab

```
-bash-4.2$ O2_jobs_report --lastdays=14 --report
```

JOBS STATES COUNT FROM 2023-04-18 TO 2023-05-03							
USERNAME	COMPLETED	FAILED					
rc	49	2					

JOBS PARTITIONS COUNT FROM 2023-04-18 TO 2023-05-03		
USERNAME	transfer	priority
rc	43	8

JOBS STATISTICS FROM 2023-04-18 TO 2023-05-03							
User	Total Jobs	Median Pending Time(hr)	Average Allocated RAM(GB)	Average Used RAM(GB)	Max Used RAM(GB)	Jobs Using > 1/2 Alloc RAM	RAM Efficiency(%)
rc	51	0.01	1.9	0	0	0	0.04

User	Average Allocated CPU	CPU Efficiency(%)	Average Runtime(hr)	WallTime Efficiency(%)	Jobs Using > 1/2 WallTime
rc	1	26.4	0.01	0.1	0

```
-bash-4.2$ O2_jobs_report
```

JOBID	USER	ACCOUNT	PARTITION	STATE	STARTTIME	WALLTIME(hr)	nCPU,RAM(GB),nGPU	PENDINGTIME(hr)	CPU_EFF(%)	RAM_EFF(%)	WALLTIME_EFF(%)
7636669	rc	rccg	transfer	COMPLETED	2023-05-01	24.0	1,2,0,0	0.02	27.9	0.0	0.1
7647004	rc	rccg	priority	COMPLETED	2023-05-01	0.1	1,0,1,0	0.03	22.2	0.0	3.0
7647127	rc	rccg	priority	COMPLETED	2023-05-01	0.1	1,0,1,0	0.03	8.3	0.0	4.0
7653780	rc	rccg	transfer	COMPLETED	2023-05-01	24.0	1,2,0,0	0.02	0.0	0.0	0.0
7687145	rc	rccg	transfer	COMPLETED	2023-05-01	24.0	1,2,0,0	0.02	0.0	0.0	0.0
7727291	rc	rccg	transfer	COMPLETED	2023-05-02	24.0	1,2,0,0	0.01	20.9	0.0	0.2

Create your own modulefiles

Here's an example modulefile (hmmer/3.2.2.lua)

```
help([[
For detailed instructions, go to:
    http://hmmer.org

PDF documentation current available at:
    http://eddylib.org/software/hmmer/Userguide.pdf
]])

whatis("Version: 3.3.2")
whatis("Keywords: sequence, HMM, hidden, Markov, model, homolog, alignment")
whatis("URL: http://hmmer.org ")
whatis("Description: biosequence analysis using profile hidden Markov models")

prepend_path( "PATH", "/n/app/hmmer/3.3.2-gcc-9.2.0/bin")

prepend_path( "MANPATH", "/n/app/hmmer/3.3.2-gcc-9.2.0/share/man")

family("hmmer")
```

``help()`` can let you document the purpose of the module, as well as any helpful usage tips/whatever you'd like to include for informational purposes. This text shows up when using ``module help <modulename>``.

``whatis()`` allows you to specify metadata for your module.

``prepend_path()`` allows you to modify environment variables that manage paths, such as PATH, MANPATH, etc. ``append_path()`` does something similar, but appends to the existing environment variable instead of prepends.

``family()`` allows you to create mutually exclusive versions for a given software type (e.g., only one version of "hmmer" can be loaded at a time).

Strictly speaking, every element of this file is optional. For function, likely only one or more instances of ``prepend_path()`` are needed to set up a working module. For more advanced utilities to use in modulefile creation, refer to the [Lmod documentation](#) or contact rchelp@hms.harvard.edu for assistance.

Creating your own modulefiles, cont.

- O2 modulefile general hierarchy:

```
Core/<program>/<version>.lua
Linux/<program>/<version>.lua
Compiler/<compiler>/<version>/<program>/<version>.lua
MPI/<compiler>/<version>/<mpi>/<version>/<program>/<version>.lua
```

- We suggest for custom modulefile organization to at least have a central `software` location, with software titles underneath, e.g.,
 - `software`
 - `software1`
 - `version1.lua`
 - `version2.lua`
 - `software2`
 - `version1.lua`
 - `version2.lua`
 - Etc.
- Do whatever makes the most sense for yourself/the lab
- Once your modulefile hierarchy location/path is determined, add it to MODULEPATH via the following:
 - `module use /path/to/software`
 - Where `/path/to/software` is the location of your software hierarchy
 - It may be useful to add it to your `.bashrc` file to prevent having to run the above command every time you log in.
- Note that if you install a version of software that matches a version on O2, yours will take precedent.



Are you running out of space? Or seeing errors with your jobs?

- Perhaps you have hit a [storage limit](#)!
- Common error message: "No space left on device"
- Being over a storage limit will prevent you from:
 - Adding/modifying files
 - Writing hidden configuration/"housekeeping" files
- Use quota-v2 to check your storage usage and limits!
- If you're over a storage limit, remove files or move them to another storage location.



Are you running out space? Or seeing errors with your jobs? Cont.

- Use quota-v2 to check your storage usage and limits!

[rc@login06 ~ 2023-05-02 14:48:29]\$ quota-v2

Usage for rc (As of 2023-05-02 14:00:00)

path	type	username	usage	storage limit	last update
/home	user	orchestra_rc	3.30 GiB	100.00 GiB	2023-05-02 14:00:00
/n/scratch3/users	user	orchestra_rc		10.00 TiB	2023-05-02 14:00:00
/n/standby/hms/it/rccg	group		10.53 TiB	14.00 TiB	2023-05-02 14:00:00

Exceeded quotas

path	type	username	usage	storage limit	last update	grace remaining
/n/standby/hms/it/rccg_q1	group		502.93 MiB	500.00 MiB	2023-05-02 14:00:00	None
/n/standby/hms/it/rccg_q2	group		502.93 MiB	500.00 MiB	2023-05-02 14:00:00	None
/n/standby/hms/it/rccg_q6	group		1.18 GiB	500.00 MiB	2023-05-02 14:00:00	None

You have lost write access to one or more of the above paths. Please clear them up to regain write access.

[rc@login06 ~ 2023-05-02 14:49:13]\$

Annotations:

- Your usage:** Points to the first two rows of the first table (user paths).
- Group usage:** Points to the third row of the first table (group path).
- Exceeded limits:** Points to the three rows of the second table (paths where quotas are exceeded).

*Available on the [O2 command line](#) (see above) or on the [O2Portal](#).

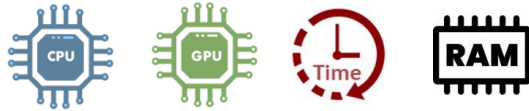


HARVARD
MEDICAL SCHOOL

RESEARCH COMPUTING

O2 Portal – An alternative or complement to the command line

- Access to O2 resources



- Access to popular applications



- Transfer data to/from O2
- Move data between O2 filesystems
- View/edit files



Tmux - why is important?

- Tmux is a terminal multiplexer in Linux
 - Enable pseudo-terminal instances
 - Detach/Attach from a pseudo-terminal
- Use cases
 - Data Transfer on the O2 transfer cluster
 - Continue to run a command *after* disconnecting from a session



Loops - save time and streamline a computational workflow

```
for fq in *fq; do
```

```
    sbatch -p short -t 10:0 --wrap "fastq_to_fasta -i $fq -o ${fq%.fq}.fa"
```

```
Done
```

```
for file in *fq; do
```

```
    batch="$batch $file"
```

```
    [ ${#batch} -gt 100 ] && sbatch -p short -t 1:0:0 --wrap "for fq in $batch; do fastq_to_fasta -i $fq -o ${fq%.fq}.fa; done"
```

```
    Batch=""
```

```
done
```

```
[ -z "$batch" ] || sbatch -p short -t 1:0:0 --wrap "for fq in $batch; do fastq_to_fasta -i $fq -o ${fq%.fq}.fa; done"
```



Don'ts

- Do not run your analysis on the login nodes
- Do not submit short [seconds] jobs to O2
- Do not request too much memory
- Do not be afraid to ask for help! Contact at rchelp@hms.harvard.edu

