

Shared Computing Cluster Usage Cheat Sheet

Useful Links

RCS website	rccs.bu.edu
SCC OnDemand	scc-ondemand.bu.edu
email	help@scc.bu.edu
Software list	rccs.bu.edu/software
Examples	rccs.bu.edu/examples

module (software environment)

<code>module avail</code>	List available packages
<code>module avail <i>matlab</i></code>	List available versions of <i>matlab</i>
<code>module spider <i>matlab</i></code>	List available versions of <i>matlab</i>
<code>module load <i>matlab</i></code>	Load module <i>matlab</i>
<code>module show <i>matlab</i></code>	Show the content (env. variables) of a module
<code>module unload <i>matlab</i></code>	Unload the module
<code>module list</code>	List all loaded modules
<code>module purge</code>	Unload all loaded modules
<code>module avail -t 2>\& 1 less</code>	Pipe module list to <i>less</i> command

qcrsh/qcsh (submit an interactive job)

<code>qcrsh</code>	Submit an interactive rsh session
<code>qcsh</code>	Submit an interactive X-windows session

`qcrsh -pe omp 4 -P project-name` Interactive job example

qsub (submit a batch job)

<code>-P <i>project-name</i></code>	Project name
<code>-N <i>job-name</i></code>	Job name
<code>-l h_rt=<i>hh:mm:ss</i></code>	Hard time limit
<code>-m e</code>	Send an email when the job ends
<code>-m ea</code>	Send an email when the job ends or is aborted
<code>-M <i>my.email@gmail.com</i></code>	Use non-BU email address
<code>-j y</code>	Merge error and output files into a single file
<code>-pe omp <i>N</i></code>	Request multiple cores (4, 8, 16, 28, 32)
<code>-pe mpi_28_tasks_per_node <i>N</i></code>	MPI job (4, 16, or 28 cores per node)
<code>-t 1-<i>N</i></code>	Submit <i>N</i> tasks
<code>-l mem_per_core=6G</code>	Request at least 6G of memory per core
<code>-l gpus=1</code>	Request a node with 1 GPU
<code>-l gpu_c=3.5</code>	Minimal GPU compute capability (3.5, 6.0, 7.0)
<code>-hold_jid <i>joblist</i></code>	Setup job dependency list
<code>-l buyin</code>	Force the job to run only on a buyin node
<code>-q <i>queue-name</i></code>	Force job to run only in a specific queue
<code>-verify</code>	Instead of submitting a job, print an info

qstat (get information about jobs)

<code>qstat</code>	List of all current jobs
<code>qstat -u <i>user-id</i></code>	All current jobs submitted by the user <i>user-id</i>
<code>qstat -s r</code>	List of running jobs
<code>qstat -s p</code>	List of pending jobs (hw, hqw, Eqw...)
<code>qstat -u <i>user-id</i> -r</code>	Display the resources requested by the job
<code>qstat -u <i>user-id</i> -ext</code>	Extended info about the user's jobs
<code>qstat -u <i>user-id</i> -s r -t</code>	Display info about sub-tasks of parallel jobs
<code>qstat -j <i>job-id</i></code>	Display job status
<code>qstat -g c</code>	Display the list of queues and load information
<code>qstat -q <i>queue</i></code>	Display jobs running on a particular queue

qdel (delete job from the queue)

<code>qdel <i>job-id</i></code>	Delete job <i>job-id</i>
<code>qdel <i>job-id</i> -t 5-7</code>	Delete tasks 5 through 7 for job <i>job-id</i>
<code>qdel -u <i>user-id</i></code>	Delete all the jobs submitted by the user

qselect (list queues corresponding to selection criteria)

<code>qselect -pe omp 16</code>	list all nodes that can execute the job with selected resources
<code>qselect -l mem_total=252G</code>	list all large memory nodes
<code>qselect -pe mpi16</code>	list all the nodes that can run 16-slot mpi jobs
<code>qselect -l gpus=1</code>	list all the nodes with GPUs

qacct (past job information)

<code>qacct -j <i>job-id</i></code>	Detailed report about job <i>job-id</i>
<code>qacct -d 3 -o <i>user-id</i> -j</code>	Detailed report about all jobs in the past 3 days
<code>qacct -d 3 -o <i>user-id</i> -q <i>queue</i> -j</code>	Detailed report about all user jobs on a <i>queue</i>
<code>qacct -P <i>project-id</i></code>	Summary report for the project (current year usage)

User Guidelines

15 min. CPU time on login nodes
12 hours - default wall clock time for a job
720 hours - wall clock time limit for a single-node job
48 hours - wall clock time limit for a GPU job
120 hours - wall clock time limit for mpi job running on multiple nodes

acctool (account information)

<code>acctool -b y</code>	Balance summary of all the projects I belong to
<code>acctool -u <i>user-id</i> -b y</code>	Balance summary of all the projects <i>user-id</i> belongs to
<code>acctool 06/18/20</code>	Number of jobs and wallclock report for the day
<code>acctool -d 2 06/18/20</code>	Number of jobs and wallclock detailed report for the day
<code>acctool -d 2 -t 5 06/18/20</code>	Display detailed report for the top 5 jobs for the day
<code>acctool -d 4 06/18/20</code>	Most detailed report for all the jobs for a particular day
<code>acctool -j <i>job-id</i> 06/18/20</code>	Report for job with given job ID.

Connecting to the Shared Computing Cluster

<code>scc1, scc2, scc3(geo), scc4</code>	SCC login nodes
<code>ssh username@scc2.bu.edu</code>	Windows (in mobaXterm)
<code>ssh -Y username@scc2.bu.edu</code>	Mac
<code>ssh -X username@scc2.bu.edu</code>	Linux

quota (home directory space usage)

<code>quota</code>	Display my Home directory usage
<code>quota -s</code>	Display my Home directory usage in human-readable format
<code>quota user-id</code>	Display Home directory usage for the user

`du -hs .[^.]* *` show size of each sub-directory

Working with the Project Disc Space

<code>groups</code>	List all projects which I belong to
<code>cd /project/myproject</code>	Change directory to the /project directory
<code>cd /projectnb/myproject</code>	Change directory to the /projectnb directory
<code>cd /restricted/project/myproject</code>	Change directory to the /restricted/project directory (from scc4 only)
<code>cd /restricted/projectnb/myproject</code>	Change directory to the /restricted/projectnb directory (from scc4 only)

pquota (project space usage)

<code>pquota</code>	Quotas for the disk space for all the projects I belong to
<code>pquota -u project</code>	Disk space quota and usage for the project

Available editors and viewers

<code>emacs</code>	Text editor ("the extensible, customizable, self-documenting, real-time display editor")
<code>vi, vim, gvim</code>	Another popular text editor
<code>gedit</code>	GNOME notepad-like text editor
<code>nano</code>	GNU text editor with command-line interface
<code>evince</code>	pdf viewer
<code>display</code>	image viewer

Commands to transfer files and Popular FTP clients

Note: The following scp commands should be executed on the local machine.

<code>scp filename username@scc1.bu.edu:~</code>	Upload file from your local machine to your home directory on the SCC
<code>scp filename username@scc4.bu.edu:/project/myproject</code>	Upload file from your local machine to your specified project directory on the SCC
<code>scp username@scc4.bu.edu:/project/myproject/filename .</code>	Download file from your project directory on the SCC to the current directory on your local machine
<code>rsync filename username@scc1.bu.edu:~</code>	sync a file from your local machine with the file in your home directory on the SCC

<code>wget http://www.site.org/file</code>	Download a file from a website
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Cyberduck	Windows and MAC FTP client
FileZilla	Windows and MAC FTP client
WinSCP	Windows FTP client

<code>dos2unix filename</code>	Convert file with DOS/MAC characters to UNIX/Linux format (execute on SCC)
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Snapshots

<code>.snapshots/yymmdd</code>	Snapshots directory structure
<code>ls .snapshots/200805</code>	View the snapshot of the directory created on August 5th, 2020