Babraham Compute Cluster Cheat Sheet

qsub Queues a task on a compute node wi	ithout any interactive session links. Shell can be safely closed. Prints STOUT and STERR to files.
echo "fastqc test.fq" qsub -cwd -V -l	vf=2G -pe orte 8 -o \$f.out -j y -N \${f}_fastqc -hold_jid sra_dump
-cwd	Use current directory. Without this will run job from home directory.
-V	Transfer loaded modules to compute node
-1 <i>vf=2G</i>	Assign 2gb memory to job
-pe <i>orte 8</i>	Assign 8 cores to job
-p -1000	Set job priority, -1023 to 0. Avoid hogging the cluster.
-o [output.out] -j y	Write STOUT and STERR to output.out
-N [jobname]	Call job "jobname" in queue
-hold_jid [jobname]	Hold in queue until job "jobname" completes
-m eas -M [email address]	Send e-mail at <u>e</u> nd of job, or when <u>a</u> borted or <u>s</u> uspended
qrsh -cwd -V -1 vf=2G -pe orte 4 perl myscript.pl > output.txt & qlogin Starts a shell session on a compute node. Good for running interactive tasks, such as SeqMonk. Don't use for jobs. Exit the shell before closing. qlogin -l vf=1G -pe orte 1	
	ode. Good for running interactive tasks, such as SeqMonk. Don't use for jobs. Exit the shell before closing.
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