## **Reference Sheet for Slurm**

| Command       | What it does  |
|---------------|---|
| sinfo         | reports the state of partitions and nodes   |
| squeue        | reports the state of jobs in the batch queue  |
| sbatch script | submits a job script  |
| scancel jobid | cancels a pending or running job jobid  |
| sacct         | reports accounting information about active and completed jobs                          |
| srun          | submits job for execution or initiate job steps in real time (used inside batch script) |

| Directive         | Description   |
|-------------------|---|
| Partition         | #SBATCHpartition=partitionname OR #SBATCH -p partitionname                    |
| Account           | #SBATCHaccount=accountname OR #SBATCH -A accountname                          |
| Wall time         | #SBATCHtime=01:00:00 OR #SBATCH -t 1:00:00                                    |
| Node count        | #SBATCHnodes=2 OR #SBATCH -N 2  |
| Constraint        | #SBATCHconstraint="c12" OR #SBATCH -C "c12"                                   |
| Total tasks count | #SBATCHntasks=24 OR #SBATCH -n 24   |
| Memory            | #SBATCHmem=24576 (NOTE: memory given in MB)                                   |
| Mail options      | #SBATCHmail-type=FAIL,BEGIN,END   |
| Mail user         | #SBATCHmail-user=user@mail.com  |
| Job name          | #SBATCHjob-name=jobname OR #SBATCH -J jobname                                 |
| Stderr            | #SBATCH -e slurm-%j.err-%N (where %j is job number and %N is first node name) |
| Stdout            | #SBATCH -o slurm-%j.out-%N  |

## **Useful Slurm Aliases** that provide information in a useful manner for our clusters

Bash:

alias si="sinfo -o \"%20P %5D %14F %8z %10m %10d %11l %16f %N\"" alias sq="squeue -o \"%8i %12j %4t %10u %20q %20a %10g %20P %10Q %5D %11l %11L %R\""

alias si 'sinfo -o "%20P %5D %14F %8z %10m %11l %16f %N"'

alias sq 'squeue -o "%8i %12j %4t %10u %20q %20a %10g %20P %10Q %5D %11l %11L %R"'

To slurm commands such as squeue, sinfo (and si, sq) can add –M all to see all clusters

Can also submit to any cluster from nodes on a different cluster by using full path, e.g., from kingspeak you can submit a job to notchpeak by using /uufs/notchpeak.peaks/sys/pkg/slurm/std/bin/sbatch