

Research Computing Noor GPU Cluster Introduction

GPU Cluster Capacity

Total Number Of Compute Nodes (Intel Xeon E5-2670)	28
Total Number Of Cores	448
Total Memory	1 TB
Total Compute Capacity (Theoretical Max)	9.31 TFLOPS
Local Scratch /tmp	400 GB
Shared NFS Scratch	20 TB

Total Number Of GPU	88
Fermi M2070-Q / Kepler K20m	24 / 64
Total Number Cuda Cores	170496
Total Memory	464 GB
Total Peak single precision floating point performance	250 TFLOPS
Total Peak double precision floating point performance	87.24 TFLOPS

Cluster Specification & Configuration

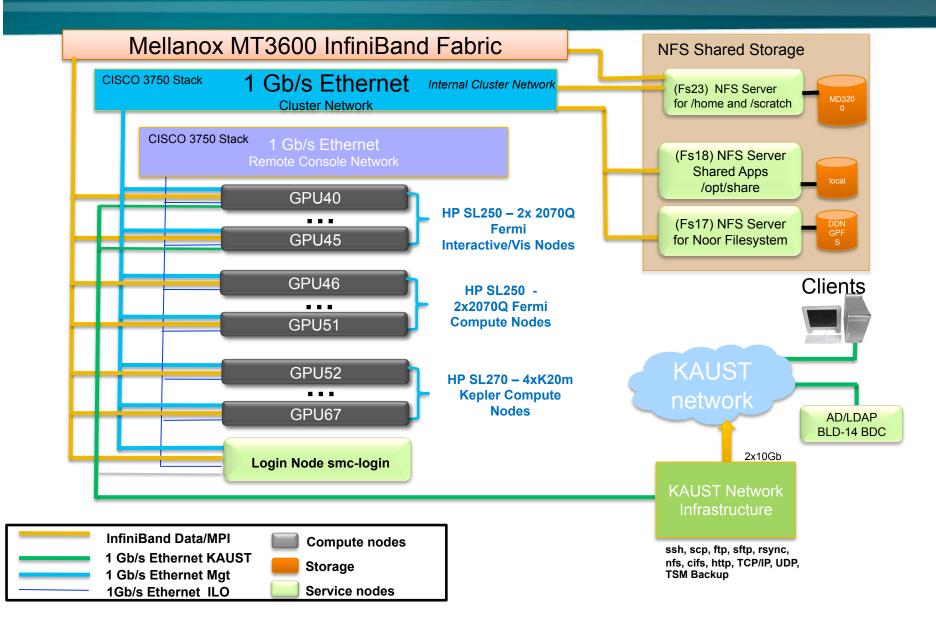
Hardware

- HP ProLiant SL250s Gen8 (Intel(R) Xeon(R) CPU E5-2670 0 @ 2.60GHz)
 - 2 Processors, 16 cores, Memory 64GB per node
 - GPU Fermi Tesla M2070-Q
 - Processor Speed 1.56GHz, Memory 6GB, Memory Bandwidth 150 GB/Sec
 - Peak double precision/single precision floating point performance 515/1030 Gigaflops
 - Cuda Cores / Node 448
- HP ProLiant SL270s Gen8 (Intel(R) Xeon(R) CPU E5-2670 0 @ 2.60GHz)
 - 2 Processors, 16 cores, Memory 64GB per node
 - GPU Kepler K20m
 - Processor Speed 1.56GHz, Memory 5GB, Memory Bandwidth 208 GB/Sec
 - Peak double precision/single precision floating point performance 1.17/3.52 TFLOPS
 - Cuda Cores / GPU 2496

Software

- OS
 - RHEL6.3, Kernel 2.6.32-279.el6.x86_64
- Drivers
 - NVRM version: NVIDIA UNIX x86_64 Kernel Module 319.17 (Fermi Nodes)
 - NVRM version: NVIDIA UNIX x86_64 Kernel Module 319.32 (K20 Nodes)
- Compilers
 - Cuda5 Toolkit and SDK, PGI and CAPS OpenACC compilers available on all the nodes NFS shared
 - Use module command to check and load the compilers
- Job Scheduler
 - LSF HPC 7 update 6

Cluster Design Document



User Guide

- Requires Noor Account to login to Noor GPU Cluster
- How submit the Batch job
 - \$ ssh smc-login.kaust.edu.sa (Use KAUST Portal User ID and Password)
 - Transfer requires files under /scratch/<uid>
 - \$ cd /scratch/<uid>
 - · Load require compilers
 - · \$ module load compilers-extra
 - \$ module load cuda
 - \$ module load pgi
 - \$ bsub -q gpu -n1 -x -R select[k20] -o out%J -e error.%J ./myjob
 - Use –x option if your job going to use all the GPU so other job cannot start on the same node
- Simple Job submission script

```
$vi myjob_script
#/bin/bash —I

#BSUB —q gpu

#BSUB —n 16 # Number of compute cores

#BSUB —R select[k20] # to select k20 nodes —R select[fermi] for Fermi nodes

#BSUB —e erro.%J

#BSUB —o output.%J

#BSUB —x #For Exclusive

cd /scratch/<uid>
module load compilers-extra

module load cuda

module load pgi
./myapps
:wq
$ bsub < myjob_script
```

- http://rcweb.kaust.edu.sa/KAUST/ResearchComputing/wiki/NoorGuide#ExampleofusingbatchjobscriptsOpenMPI
- CAUTION! Make sure to submit the job using /scratch/<uid> dir for heavy I/O please use /tmp
- How to submit Interactive Job
 - \$ ssh –X smc-login.kaust.edu.sa (Make sure Xserver" is running on your desktop (Xming, Cygwin etc.))
 - \$ bsub -XF -I -q interactive xterm
 - http://rcweb.kaust.edu.sa/KAUST/ResearchComputing/wiki/NoorGuide#InteractiveLSFJob

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Important LSF Command

- bsub # submit the job
 bjobs # check the job status
 btop # Move the job up
 bbot # Move the job down
 bstop # Suspend the job
- bresume # Resume suspended job
- bkill # kill the jobbqueues # queues list

How to request Noor Account

- KAUST USERS
 - Go to http://researchcomputing.kaust.edu.sa
 - IT Request Forms Section
 - Noor Account Request Form
- External USERS (KAUST Collaborators) require
 - In addition to Noor Account Request form submit Remote access request form
 - Remote Access Request Form

** Note KAUST internal Researcher or Faculty must make the Noor and ssh Gateway account

^{**} Note all the command applies to only your jobs.