HPC Speed Cluster

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<u>High-Performance Computing (HPC) Facility: Speed - Concordia University</u>

<u>GitHub - NAG-DevOps/speed-hpc: Speed: Gina Cody School HPC Facility: Scripts, Tools, and Refs</u> **Support**: <u>rt-ex-hpc@encs.concordia.ca</u>

HPC Team

The team can be reached at rt-ex-hpc@encs.concordia.ca

- Serguei Mokhov, PhD
 [Manager, Networks, Security, and HPC, AITS; Affiliate at CSSE and CIISE]
- **Gillian Roper**[Senior Systems Administrator, HPC and Research Support, AITS]
- Carlos Alarcón Meza
 [Systems Administrator, HPC and Networking, AITS]
- Farah Salhany
 [IT Instructional Specialist, AITS]

Speed Resources

Speed official page:

https://www.concordia.ca/ginacody/aits/speed.html

- Speed manual:
 - PDF: https://github.com/NAG-DevOps/speed-hpc/blob/master/doc/speed-manual.pdf
 - HTML: https://naq-devops.github.io/speed-hpc/
- GitHub page includes examples (PRs are welcome):

https://github.com/NAG-DevOps/speed-hpc

Slurm project account: <>

ENCS Account and GCS Webmail

- To set up ENCS account for the first time <u>OR</u> if you cannot remember your ENCS credentials, go to the ENCS service desk (you could also send an email to <u>help@concordia.ca</u> if off campus)
- To change your existing ENCS password, visit the <u>password reset page</u>
- To access ENCS webmail, visit https://webmail.encs.concordia.ca/
 - access from off campus needs VPN
 - use same login as in the lab PCs or Speed

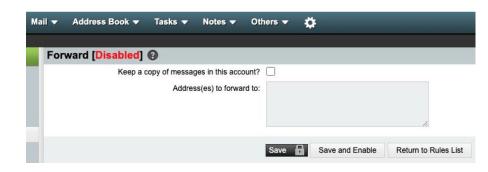
IMPORTANT

If you don't read it regularly, set up forwarding to a desired email address:

- Mail => Filters => Forward
 - => Save and Enable

WHY?

- HPC announcements are sent to your ENCS email.
- → Job notifications are also sent there by default



Policies and Procedures

- Initial quota is 32 CPUs and 4 GPUs.
 - Can be expanded, for a fixed duration, upon request.
- A mailing list has been established for announcements and discussions. All users of the Speed cluster are automatically subscribed to it: hpc-ml@encs.concordia.ca
- Support requests: <u>rt-ex-hpc@encs.concordia.ca</u>

IMPORTANT

- speed-submit is a tiny virtual used for dispatching to execution nodes and is SSH accessible from Concordia subnets or from the VPN.
 - Excessive process or IDE executed on speed-submit will be terminated.
 - Only compile things on execution nodes (same for using IDEs)

Getting Started

Connect to Speed

- Requires VPN (if off-campus)
- SSH to the cluster's submit node [speed-submit]
 - PuTTY or MobaXterm on Windows
 - Terminal on Mac or Linux

```
> ssh <ENCSusername>@speed.encs.concordia.ca
```

Go to a working directory

• First time (create a directory):

\$USER is a parameter (either manually replace it by your ENCS username or the system will automatically do the work.)

```
[speed-submit] [/home/../..] > mkdir /speed-scratch/$USER
[speed-submit] [/home/../..] > cd /speed-scratch/$USER
[speed-submit] [/speed-scratch/..] >
```

Not the first time (navigate to your directory):

```
[speed-submit] [/home/../..] > cd /speed-scratch/$USER
[speed-submit] [/speed-scratch/..] >
```

Storage

Students have 10GB quotas (50GB for faculty), however, NFS speed scratch has 10TB

To check your quota [in terminal]

> quota							
	Diskspace			Files			
WHERE	used	warn	limit	used	warn	limit	volume/qtree
Basic Home and Web	9.2G	9.8G	10.0G	56K			vol: users
INBOX	7.4M	80.0M	85.0M	1	-	-	vol: mailspool
/nettemp	0.0K	750.0M	M0.008	0	-	-	vol: nettemp
/speed-scratch	5.6G	9.8T	10.0T	48K	_	_	<pre>vol: speed_scratch</pre>

"Out of quota" errors can be diagnosed with the /encs/bin/bigfiles command

Refer to: https://nag-devops.github.io/speed-hpc/#how-to-resolve-disk-guota-exceeded-errors

Storage

If you need to copy between **G**: or **U**: drive and /speed-scratch/\$USER

Unix/Linux home U:

```
Speed to U:
```

```
> cp -r /speed-scratch/$USER/project ~/
```

U: to Speed

```
> cp -r ~/project /speed-scratch/$USER/
```

Windows home G:

• Speed to G:

```
> cp -r /speed-scratch/$USER/project /winhome/<1st letter of $USER>/$USER/
```

o G: to Speed

```
> cp -r /winhome/<1st letter of $USER>/$USER/project /speed-scratch/$USER/
```

SLURM tiny Intro

Slurm is workload manager, used for job scheduling

- Batch job (to run unattended tasks).
 - Job script acts as a wrapper for your actual job my-script.sh
 - Slurm options are usually embedded in a job script prefixed by #SBATCH directives.

for example:

```
#!/encs/bin/tcsh
#SBATCH --job-name=my-job --account=<>
#SBATCH --mail-type=ALL
#SBATCH --mem=10G -p pt -t 600
or #SBATCH --mem=10G -p ps -t 600
#SBATCH --gpus=1 -p pg
```

- Use **srun** in your script for every complex compute step.
- Submit job to the cluster

```
sbatch [options] my-script.sh
```

Required/common Job Parameters:

```
→ --job-name (-J)
```

- → --account (-A)
- \rightarrow --partition (-p)
- → --nodes (-N)
- **→** --mem
- → --time (-t)
- → --ntasks (-n)
- → --cpus-per-task (-c)
- → --gpus (-G)
- → --chdir (-D)
- → --output (-o)
- → --mail-type

SLURM tiny Intro

• Interactive jobs (to set up environments, debugging, and optimizing)

Similar to submitting a batch job but allows you to run shell commands interactively within the allocated resources.

 To request an interactive session, allocate resources with salloc [options] for example:

```
salloc -J my-interactive-job -p pt --mem=10G
salloc: Granted job allocation 339377
salloc: Waiting for resource configuration
salloc: Nodes magic-node-04 are ready for job
```

For quick, short job to compile on a GPU node, you can use an interactive srun directly, however salloc is preferred.

```
for tcsh:
```

```
for bash:
    srun --pty --gpus=1 --mem=1G -t 60 /encs/bin/tcsh
    srun --pty --gpus=1 --mem=1G -t 60 /encs/bin/bash
```

Running Batch Jobs

Check job output (pay attention to requested vs. used resources and adjust accordingly)

```
[speed-submit] [/speed-scratch/../speed-hpc/src] > seff 102406

Dob ID: 102406
Cluster: speed
User/Group: carlos/carlos
State: COMPLETED (exit code 0)
Cores: 1
CPU Utilized: 00:00:00
CPU Efficiency: 0.00% of 00:00:30 core-walltime
Job Wall-clock time: 00:00:30
Memory Utilized: 4.84 MB
Memory Efficiency: 0.09% of 5.00 GB
```

Other commands to check:

- sinfo -la
- squeue
- squeue -A comp6831f24
- showjob JOBID
- jobsummary JOBID
- sstat JOBID #for each step

To cancel a job:

- Batch:
 - > scancel JOBID
- Interactive:
 - > exit

REFERENCES

- HPC GitHub: https://github.com/NAG-DevOps/speed-hpc
- Speed Manual: https://naq-devops.github.io/speed-hpc/
- High-Performance Computing (HPC) Facility: Speed: https://www.concordia.ca/ginacody/aits/speed.html
- Connect to VPN: https://www.concordia.ca/it/support/connect-from-home.html
- Connect to a GCS server: https://www.concordia.ca/ginacody/aits/support/fag/ssh-to-gcs.html
- MobaXterm: https://mobaxterm.mobatek.net/download-home-edition.html
- Linux and scripting lessons: https://software-carpentry.org/lessons/
- Udemy: https://www.concordia.ca/it/services/udemy.html
- Calcul Québec's example: https://github.com/calculquebec/formation_cpu-gpu
- Launch X (Graphical) applications: https://www.concordia.ca/ginacody/aits/support/faq/xserver.html
- The Unix Shell: https://swcarpentry.github.io/shell-novice/index.html

For help, email: <u>rt-ex-hpc@encs.concordia.ca</u>

THE END

Thank You