# Minerva GPU (graphics processing unit)

## Minerva Scientific Computing Environment

https://labs.icahn.mssm.edu/minerva/ab

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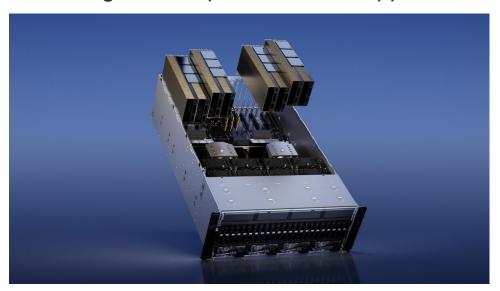


### **Outline**

- What is GPU?
- GPU resources on Minerva
- User GPU Software Environment on Minerva
- Run GPU jobs in LSF
- Reminder and future training sessions

### What is GPU?

- A graphics processing unit (GPU) is a specialized electronic circuit initially designed to accelerate computer graphics and image processing.
- GPUs can be used across a wide range of compute-intensive applications:
  - Al/Machine Learning
  - Simulations
  - Professional visualization
  - Gaming



## **CPU vs GPU**

	CPU vs GPU	
	CPU	GPU
Function	Generalized component that handles main processing functions of a server	Specialized component that excels at parallel computing
Processing	Designed for serial instruction processing	Designed for parallel instruction processing
Design	Fewer, more powerful cores	More cores than CPUs, but less powerful than CPU cores

<u>reference</u>

## **GPU** resources on Minerva

- GPU resources on Minerva
  - Current:
    - Interactive queue (1 GPU node)
    - gpu queue for batch (23 GPU nodes)
  - By Jan. 2025:
    - 51 additional nodes with 220 GPUs in total.

		С	Jan. 2025			
	V100	A100	A100-80GB	H100	H100-NVlink	L40S
# of nodes	12	8	2	2	47	4
GPU card/node	4 V100	4 A100	4 A100	4 H100	4 H100	8 L40S
CPU cores	32	48	64	64	96	96
host memory	384 GB	384 GB	2 TB	512 GB	1.5 TB	1.5 TB
GPU memory	16 GB	40 GB	80 GB	80 GB	80 GB	48 GB

## **User GPU Software Environment - Major packages**

#### OS: Centos 7.9 with glibc-2.17(GNU C library) available

- Packages with GPU support:
  - Schrödinger Suite, Amber tools, NAMD, Gromacs, Alpha Fold2, etc.
- Al tools with python/3.7.3
  - CuPy, cuDF, cuML, Numba, scikit-learn, Scanpy, Squidpy, etc.
  - Minerva Python instruction
- Al tools with conda
  - MONAI, Rapids, NVFlare, tensorflow, pytorch, etc.
  - Minerva conda instruction
- Al tools with singularity
  - Holoscan, BioNeMo, Parabricks, DeepVariant, etc.
  - Minerva singularity instruction
  - Minerva Singularity training
- Cuda toolkit versions up to 12.1.1
- Nsight Systems

### **User Software Environment: Lmod**

> 1000 modules, and different versions are supported on Minerva

#### **Lmod Software Environment Module system implemented:**

Search for module: \$module avail or \$module spider
 Check all available R versions \$ml spider R

......R/3.3.1, R/3.4.0-beta, R/3.4.0, R/3.4.1, R/3.4.3\_p, R/3.4.3, R/3.5.0, R/3.5.1\_p, R/3.5.1, R/3.5.2, R/3.5.3

- To check the detailed PATH setting in module files: \$\frac{\$ml show R}{\text{}}\$
- Load module: \$ml python or \$module load python or \$ml python/2.7.16 (for a specific version)
- Unload module \$mI -gcc or \$module unload gcc gail@1@1603c03: ~ \$ ml python.
- List loaded modules: \$ml or \$module list
- Purge ALL loaded modules \$ ml purge
- Autocompletion with tab
- More at:
  - Minerva Lmod quide
  - Lmod user guide

```
gail01@li03c03: ~ $ ml python
gail01@li03c03: ~ $ ml

Currently Loaded Modules:
   1) gcc/8.3.0   2) python/3.7.3

gail01@li03c03: ~ $ ml python/2.7.16

The following have been reloaded with a version change:
   1) python/3.7.3 => python/2.7.16

gail01@li03c03: ~ $ ml -gcc
```

### **Interactive access to GPU resources**

- Set up an interactive environment on compute nodes
- Useful for testing and debugging jobs

bsub -P acc\_hpcstaff -q interactive -n 4 -W 2:00 -R rusage[mem=4000] -R span[hosts=1] --gpu num=1 -R v100 -XF -Is /bin/bash

- Interactive, gpu, gpuexpress can be specified for -q.
- -ls: Interactive terminal/shell
- -XF: X11 forwarding
- /bin/bash: the shell to use
- GPU option specification:
  - V100: -gpu num=Ngpus -R v100
  - A100: -gpu num=Ngpus -R a100
  - A100-80G: -gpu num=Ngpus -R a10080g
  - H100: -gpu num=Ngpus -R h10080g

## Minerva LSF queues with GPUs

Queue structure in Minerva									
Queue	Wall time limit	available resources							
interactive (Dedicated to interactive jobs)	12 hours	4 nodes+1 V100 GPU nodes							
gpu	6 days	40 V100*, 16 A100, 4 A100-80GB, 4 H100-80GB							
gpuexpress	15 hours	40 V100*, 16 A100, 4 A100-80GB, 4 H100-80GB							

## **bhosts**

- bhosts queue\_name
  - o bhosts gpu
  - bhosts gpuexpress

[[yuj25@li03c03	~]\$	bhosts	gpu							
HOST_NAME		STATUS		JL/U	MAX	NJOBS	RUN	SSUSP	USUSP	RSV
dg-p100		ok		:	20	19	8	0	0	11
lg03a03		ok		_	32	4	4	0	0	0
lg03a04		ok		-	32	5	5	0	0	0
lg03a07		ok		_	32	4	4	0	0	0
lg03a08		ok		-	32	8	8	0	0	0
lg03a10		ok		9 <u></u> 6	32	8	8	0	0	0
lg03a11		ok		-	32	8	8	0	0	0
lg03a12		ok		_	32	0	0	0	0	0
lg06g28		ok		: <del></del> /	64	13	13	0	0	0
lg07c05		ok		_	48	37	37	0	0	0
lg07c06		ok			48	29	29	0	0	0
lg07c07		ok		9 <u>—</u> 0	48	17	17	0	0	0
[lg07c08		ok		-	48	11	11	0	0	0
lg07c09		ok			64	24	24	0	0	0

## **Batch job submission example**

#### \$ cat myfirst.lsf

```
#!/bin/bash
#BSUB -J myfirstjob
                                            # Job name
#BSUB -P acc hpcstaff
                                             # REQUIRED; To get allocation account, type "mybalance"
#BSUB -q premium
                                             # queue; default queue is premium
#BSUB -n 1
                                             # number of compute cores (job slots) needed, 1 by default
#BSUB -W 6:00
                                             # REQUIRED; walltime in HH:MM
                                             # 4000 MB of memory request per "-n"; 3000 MB by default
#BSUB -R rusage[mem=4000]
#BSUB -oo %J.stdout
                                             # output log (%J : JobID)
#BSUB -eo %J.stderr
                                            # error log
                                             # Initialize the execution environment
#BSUB -L /bin/bash
ml gcc
                                            # Commands that you need to run
cd /sc/arion/work/MyID/my/job/dir/
../mybin/serial executable < testdata.inp > results.log
```

#### \$ bsub < myfirst.lsf

Job <87426883> is submitted to queue premium>.

## **GPGPU** - batch jobs

```
#BSUB -q gpu
                                                       # submit to gpu queue
                                                       # Ncpu is 1~48 on A100
#BSUB -n Ncpu
                                                       # request 4 GPUs per node on A100 node
#BSUB -gpu num=4
#BSUB -R a100
#BSUB -R span[hosts=1]
                                                       # request all gpu cards on the same node
module purge
module load anaconda3 (or 2)
                                                       # to access tensorflow
module load cuda
                                                       # to access the drivers and supporting
source activate tfGPU
                                                       subroutines
python -c "import tensorflow as tf"
```

## **GPGPU** - batch jobs (continue)

- LSF will set CUDA\_VISIBLE\_DEVICES to the list of GPU cards assigned to the job.
   E.g: 2,1,3 Most standard packages honor these assignments
  - DO NOT MANUALLY CHANGE THE VALUE OF CUDA\_VISIBLE\_DEVICES.
- Multiple GPU cards can be requested across different GPU nodes

```
#BSUB -q gpu # submit to gpu queue

#BSUB -n 8 # 8 compute cores requested

#BSUB -R span[ptile=2] # 2 cores per node, so 4 nodes in total requested

#BSUB -R v100 # request specified gpu node v100, change to a100,

#BSUB -gpu num=2 # 2 GPUs requested per node
```

Note that 2 GPU cards will be reserved on each of 4 nodes for your job. If your job cannot /does not run in distributed mode, you will still lock these resources on the nodes that you are not using and prevent others from being dispatched to those node.

CUDA\_VISIBLE\_DEVICES may be defined differently on each of the nodes allocated to your job.

### **GPGPU - Local SSD**

- Make your own directory under /ssd and direct your temporary files there.
- Clean up your temporary files after completion.

	A100	1.8 TB SATA SSD			
Current	A100-80GB	7.0 TB NVMe PCIe SSD			
	H100	3.84 TB NVMe PCle SSD			
Jan. 2025	L40S	3.84 TB NVMe PCle SSD			
	H100	3.84 TB NVMe PCle SSD			

```
#BSUB -q gpu

#BSUB -gpu num=2

#BSUB -R v100

#BSUB -R span[hosts=1]

#BSUB -E "mkdir /ssd/YourID_$LSB_JOBID"

#BSUB -Ep "rm -rf /ssd/YourID_$LSB_JOBID"
```

### nvidia-smi

- A monitoring and management command line utility, nvidia-smi
- Only available on nodes with GPUs

NVID	IA-SMI	525.1	05.17 [	Driver	Version:	525.105.1	7 CU	DA Versi	on: 12.0
GPU   Fan	Name Temp	Perf	Persiste Pwr:Usaç		The second second of the second secon	Dis Memory-Usa			
	NVIDIA 74C	A100 P0	-PCI 242W /	7700 T		0:06:00.0 ( iB / 40960		80%	E. Proce Disabl
1   N/A 	NVIDIA 49C	A100 P0	-PCI 126W /	(T) (1) (1)		0:2F:00.0 ( iB / 40960		45%	E. Proce Disabl
2 N/A	NVIDIA 28C		-PCI 32W /	3.38 N		0:86:00.0 ( iB / 40960		0%	E. Proce Disabl
3 N/A	NVIDIA 28C		-PCI 33W /	(C)		0:D8:00.0 ( iB / 40960		0%	E. Proce Disabl
+ +									
Proce   GPU 	esses: GI ID	CI	PII	) Тур	oe Proc	ess name			GPU Memo Usage
======   0   1   2   3	N/A N/A N/A N/A N/A	N/A N/A	13439 113104 13439 13439	4 9	Ci	======== acs/bin.AV 11/progs/b acs/bin.AV on	in/gmx	_24_tmpi	744M

### **Minerva Ticket Submission**

- Send an email to: <a href="mailto:hpchelp@hpc.mssm.edu">hpchelp@hpc.mssm.edu</a> to start a ticket.
- Information to include:
  - The error message or the location of the log file on Minerva.
    - use command "pwd" to check the current working directory.
  - All the commands you used to get the error message.
  - The location of the scripts used and how did you run the script.
  - Which node.
    - yuj25@<mark>li03c04</mark>
  - The job ID and location of the job script.
    - Job <123456789> is submitted to queue premium>.
  - Please do not include any "<" in the email.</li>

## **Friendly Reminder**

- Never run jobs on login nodes
  - For file management, coding, compilation, etc., purposes only
- Never run jobs outside LSF
  - Fair sharing
  - Scratch disk not backed up, efficient use of limited resources
  - Job temporary dir configured to /local/JOBS instead of /tmp.
- Logging onto compute nodes is no longer allowed
- Follow us by visiting <a href="https://labs.icahn.mssm.edu/minervalab">https://labs.icahn.mssm.edu/minervalab</a>
- Acknowledge Scientific Computing at Mount Sinai and NIH funding in your publications with the template
  - https://labs.icahn.mssm.edu/minervalab/policies/acknowledge-scientific-computing-at-mount-sinai/

## **Minerva Seven Training Sessions Spring 2024**

Seven training sessions in person/Zoom this Spring with more info <a href="here">here</a>

Session	Topic	Date	Time
1	Minerva Intro	27-Mar	1PM
2	LSF Job Scheduler	3-Apr	1PM
3	Intro to GPU resources on Minerva	10-April	1PM
4	5 Ways to Get Started with GPUs*	12-Apr	1PM
5	Accelerated general data science in medicine with RAPIDS, CuPy and Numba*	17-Apr	1PM
6	Data Ark	24-Apr	1PM
7	How to Accelerate Genome Analysis Toolkit (GATK) by using Parabricks*	1-May	1PM

### Last but not Least

Got a problem? Need a program installed? Send an email to:

hpchelp@hpc.mssm.edu

## Acknowledgements

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# Thank you!