INF 503 Homework 1

Problem #1 and Problem #2

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Problem #1 (of 2): Monsoon account creation and workshop

Exercise 1:

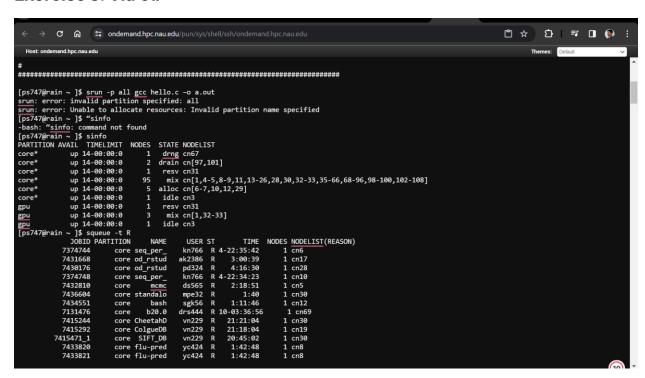


Exercise 2:



```
ሷ∣ ₹ 🗖 👰 :
      → C 🙃 ondemand.hpc.nau.edu/pun/sys/shell/ssh/ondemand.hpc.nau.edu
   Host: ondemand.hpc.nau.edu
The authenticity of host 'rain.hpc.nau.edu (10.15.108.28)' can't be established. ECDSA key fingerprint is SHA256:cdk5Vk5e4OOk3tsKwFgHvLVc112uYiuVYrBiRU+Ue/M. Are you sure you want to continue connecting (yes/no/[fingerprint])? yes Warning: Permanently added 'rain.hpc.nau.edu,10.15.108.28' (ECDSA) to the list of known hosts. ps747@rain.hpc.nau.edu's password:
# Welcome to Monsoon - login node: [rain]
# Red Hat Enterprise Linux release 8.9 (Ootpa) - Kernel: 4.18.0-513.11.1.el8_9.x86_64 # slurm 23.11.1
# You are logged in as ps747
# # Information:
# - Monsoon now running Enterprise Linux 8
# - /scratch : files auto DELETED after 30 days
#
# Issues or questions: ask-arc@nau.edu
# Maintenance:
# - None at this time
# Random tip:
# Read additional monsoon documentation here: http://nau.edu/hpc/docs
[ps747@rain ~ ]$
```

Exercise 3: Via cli



1. How many nodes make up a monsoon? - Hint: use "sinfo"

Ans: The Monsoon cluster has 105 nodes in total. The GPU division consists of five nodes.

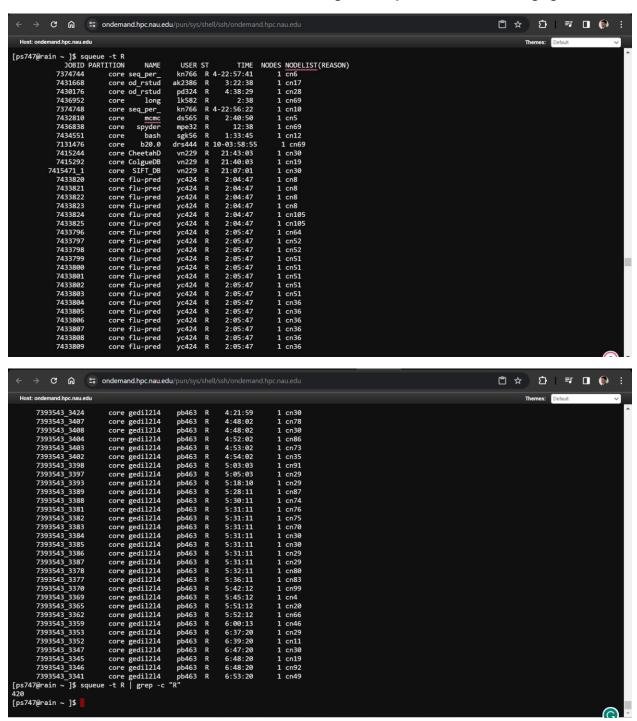
2. How many nodes are in the GPU partition?

Ans: The GPU division consists of five nodes.

3. How many jobs are currently running? – Hint: use "squeue -t R"

Ans: On the cluster, 420 jobs are executing right now.

The below screenshot shows that the current running has 420 jobs that are executing right now.



[ps747@rain ~]\$ squeue -t R | grep -c "R"

420

[ps747@rain ~]\$

4. How many jobs are currently pending? Why? - Hint: use "squeue -t PD"

```
→ C 🙃 25 ondemand.hpc.nau.edu/pun/sys/shell/ssh/ondemand.hpc.nau.edu
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[ps747@rain ~ ]$ squeue -t PD
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→ C ondemand.hpc.nau.edu/pun/sys/shell/ssh/ondemand.hpc.nau.edu
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[ps747@rain ~ ]$ squeue -t PD | grep -c
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"PD"
[ps747@rain ~ ]$
```

[ps747@rain ~]\$ squeue -t PD | grep -c "PD"

28

> 28 jobs are currently pending.

Exercise 4 via CLI

JobID	JobName	ReqMem	MaxRSS	ReqCPUS	UserCPU	Timelimit	Elapsed	State	JobEff
7430215	exercise1	1.95G	4.04M	1	00:00.252	00:20:00	00:00:33	COMPLETED	1.48
7434228	long		4.05M		00:00.149		00:05:01	COMPLETED	1.6
7437678	hostname	500M	0.1M	1	00:00.001	02:00:00	00:00:30	FAILED	
7438759	lazy		453M		00:02.002		00:01:06	COMPLETED	5.25
7438925	lazy		560M		00:01.626		00:01:06	COMPLETED	6.1
7438931	lazy		454M		00:01.581		00:01:05	COMPLETED	5.19
7438968	lazy		454M		00:01.650		00:01:06	COMPLETED	5.22
	1.16% 1.65% 1.11% 1.12 1.12 1.12 1.14								

[ps747@rain ~]\$ jobstats -rJobID JobName ReqMem MaxRSS ReqCPUS UserCPU Timelimit Elapsed State JobEff

Memory: 10.16%

CPU: 00.65%

Time Limit: 03.11%

Efficiency Score: 4.64

Below are the MaxRSS values separately:

[ps747@rain ~]\$ jobstats -r | awk 'NR > 2 {print \$4}'

4.04M

4.05M

0.1M

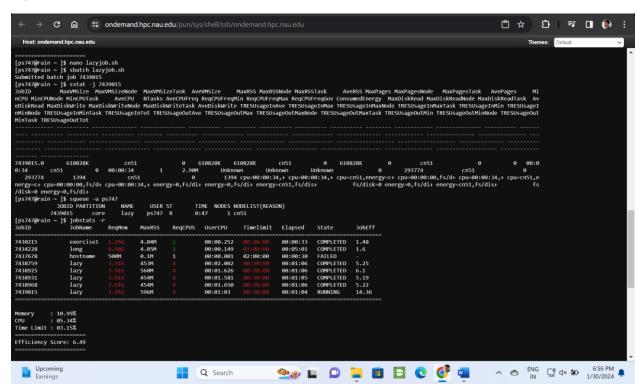
453M

560M

454M

454M

e.g. --mem=600



Memory: 10.99%

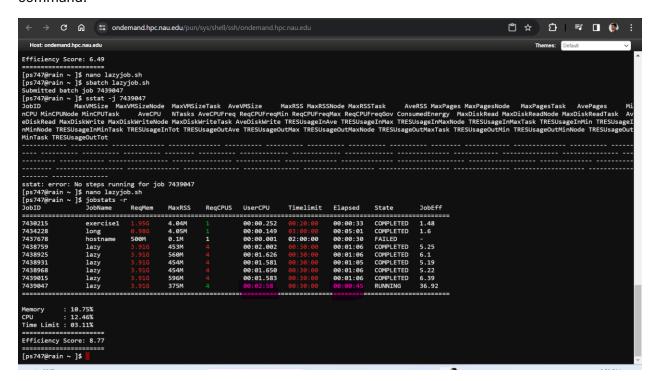
CPU: 05.34%

Time Limit: 03.15%

Efficiency Score: 6.49

===============

Edit the lazy job script, comment out the first srun command, and uncomment the second srun command.



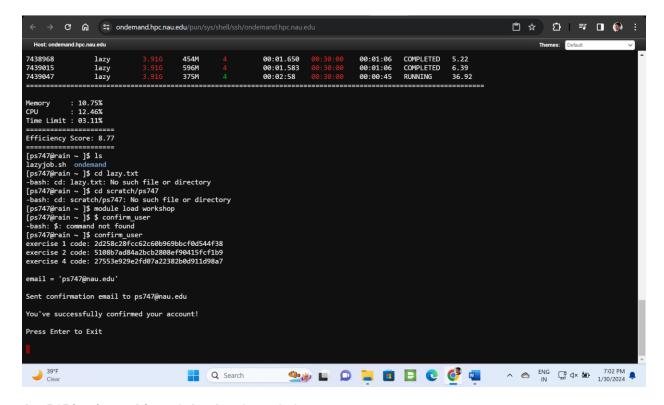


secret code from lazy.txt:

stress: info: [3865701] dispatching hogs: 3 cpu, 0 io, 1 vm, 0 hdd

stress: info: [3865701] successful run completed in 65s

The secret code for exercise4 is: 27553e929e2fd07a22382b0d911d98a7



[ps747@rain ~]\$ module load workshop

[ps747@rain ~]\$ \$ confirm_user
-bash: \$: command not found

[ps747@rain ~]\$ confirm_user

exercise 1 code: 2d258c28fcc62c60b969bbcf0d544f38

exercise 2 code: 5108b7ad84a2bcb2808ef90415fcf1b9

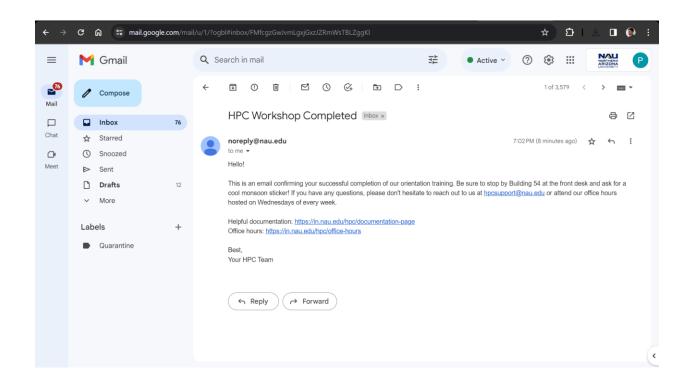
exercise 4 code: 27553e929e2fd07a22382b0d911d98a7

email = 'ps747@nau.edu'

Sent confirmation email to ps747@nau.edu

You've successfully confirmed your account!

Press Enter to Exit



Problem #2 (of 2): basic text processing

Below is the execution process and steps with commands for problem 2 of homework assignment 1:

Commands:

Use the command 'g++ gnome1.cpp -o gnome' to compile the genome.cpp file.

Next, run the code by executing the below command to run the executable file:

./gnome 1 /common/contrib/classroom/inf503/genomes/human.txt

[ps747@rain ~/Homeworks]\$ g++ gnome1.cpp -o gnome [ps747@rain ~/Homeworks]\$./gnome 1 /common/contrib/classroom/inf503/genomes/human.txt

Output screenshot for problem 1:

```
Host ondemandhpc.nau.edu

Themes: Total

Themes: The
```

To execute part 2 of homework 1, please use the below command.

./gnome 2 /common/contrib/classroom/inf503/genomes/human.txt

Output screenshot for problem 2:

Execution using makefile:

Enter the make command to run the Makefile as below in the command line:

```
← → C ♠ condemand.hpc.nau.edu/pun/sys/shell/ssh/ondemand.hpc.nau.edu

Host: ondemand.hpc.nau.edu

[ps747@rain ~/Homeworks ]$ make g++ gnome1.cpp -o gnome1 [ps747@rain ~/Homeworks ]$
```

Next, run the code by executing the below command to run the executable file:

./gnome1 1 /common/contrib/classroom/inf503/genomes/human.txt

```
Nost ordemand/spc.nau.edu

Red Hat Enterprise Linux release 8.9 (Ootpa) - Kernel: 4.18.0-513.11.1.el8_9.x86_64

# slurm 23.11.1

* You are logged in as ps747

# Information:

* Nonsoon now running Enterprise Linux 8

* - None at this time

# Random tip:

# "sinfo -1 -N" -- Get more info on node details and features

# "sinfo -1 -N" -- Get more info on node details and features

# "sinfo -1 -N" -- Get more info on node details and features

# "sinfo -1 -N" -- Get more info on node details and features

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# "sinfo -1 -N" -- Get more info on node details and features

# "sinfo -1 -N" -- Get more info on node details and features

# "sinfo -1 -N"
```

To execute part 2 of homework 1, please use the below command.

./gnome1 2 /common/contrib/classroom/inf503/genomes/human.txt

- 1. How many scaffolds were there?
- A. Total scaffolds: 607
- 2. What was the longest and 2nd longest scaffold? Provide names of scaffolds and lengths.

```
A.

B. First longest scaffold: 568815346-9606, size: 147687515

Second longest scaffold: 568815332-9606, size: 131283175
```

- 3. What was the average scaffold length?
- A. Average scaffold length: 5036552
- 4. How long does it take (in seconds) to execute this function? Hint: You will need to use system time within your code to get accurate time estimates.
- A. Time taken: 43.5763s
- 5. What was the GC content of the human genome (percent of C's and G's in the genome)?
- A. The percentage(%) of GC content is: 40.9955%