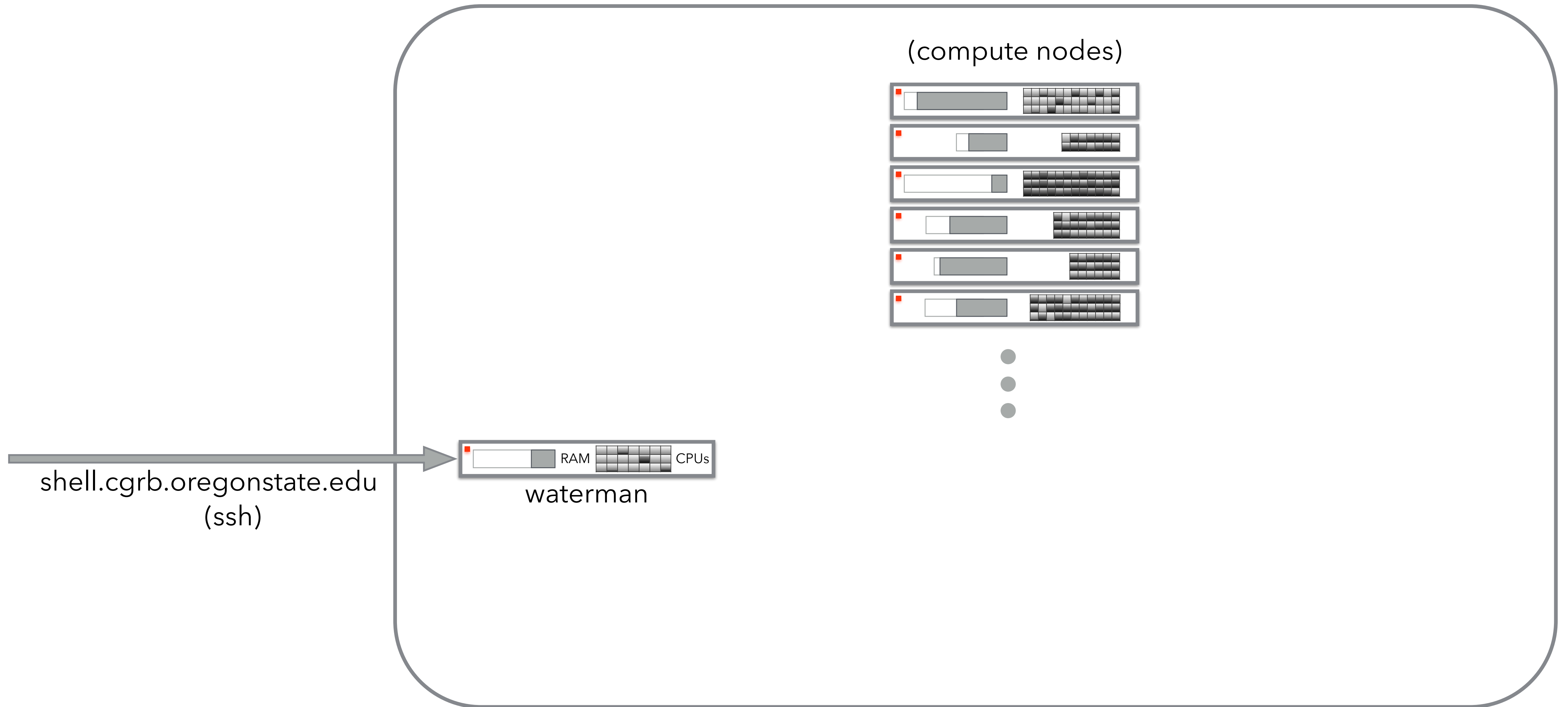


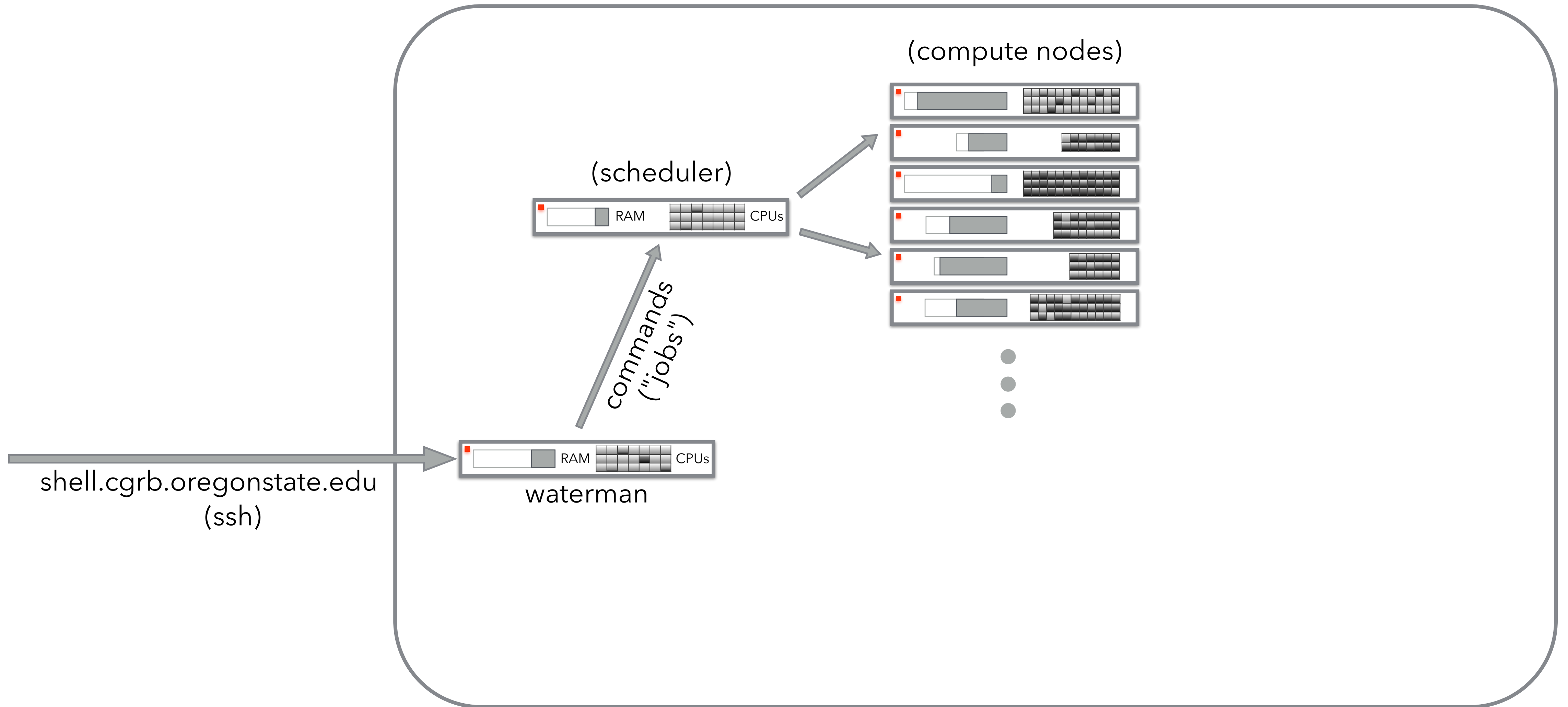
# CGRB Computational Infrastructure



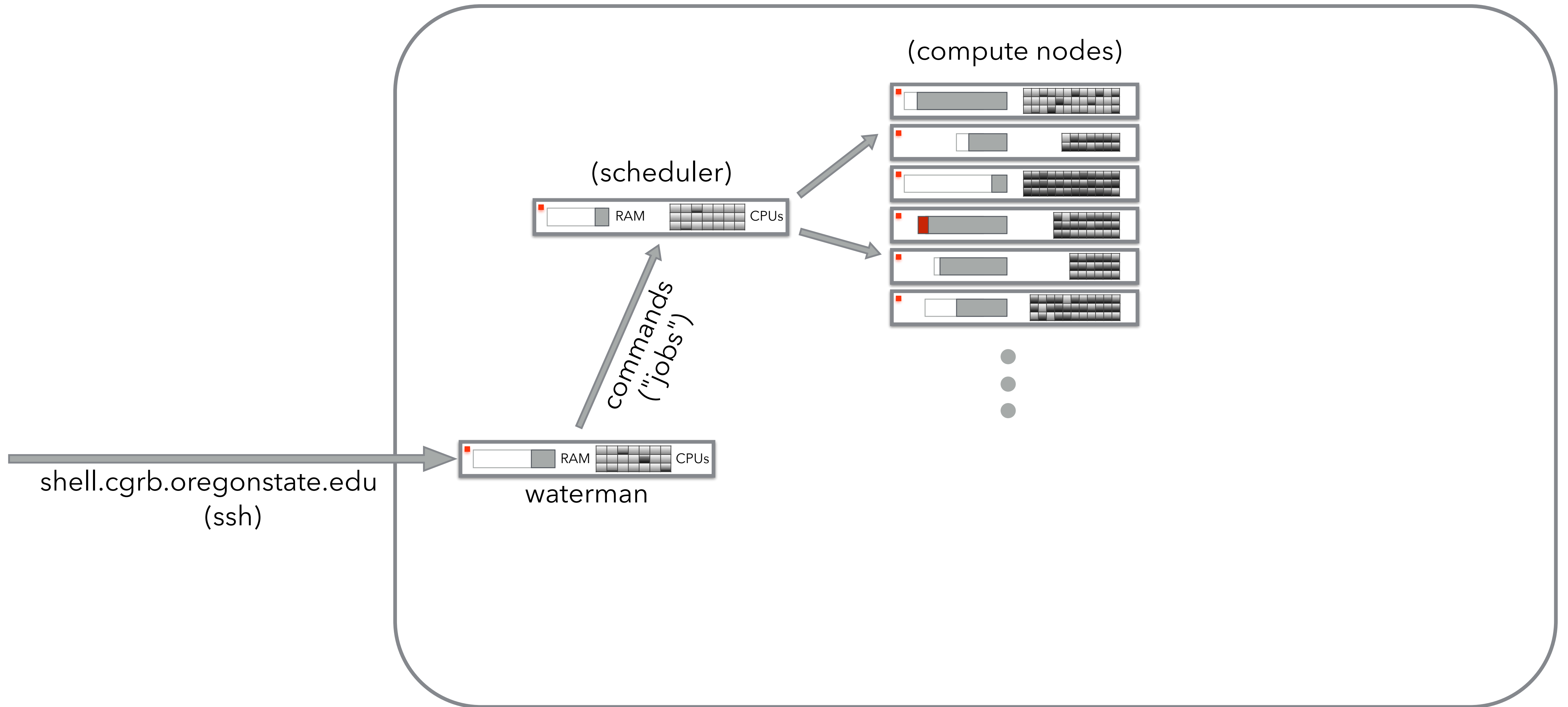
# CGRB Computational Infrastructure



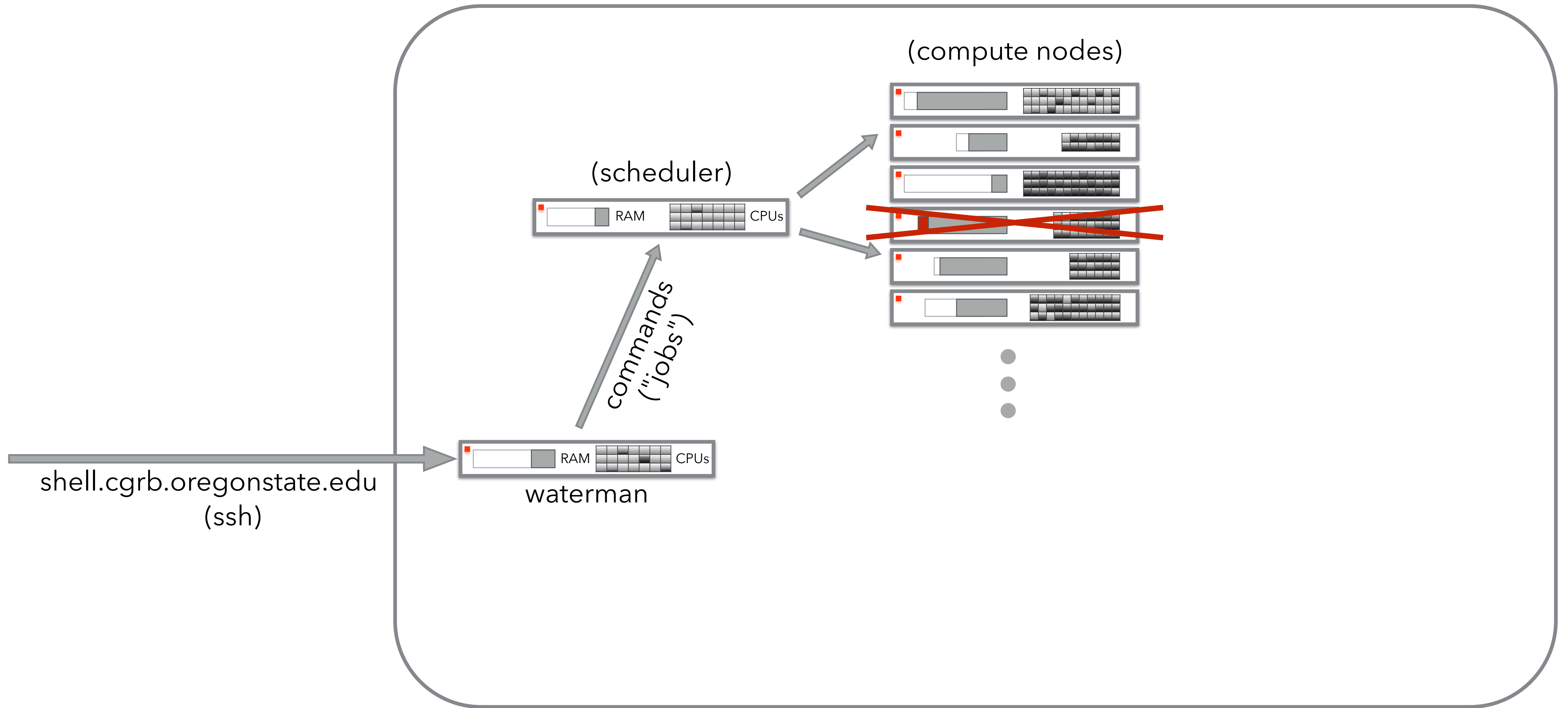
# CGRB Computational Infrastructure



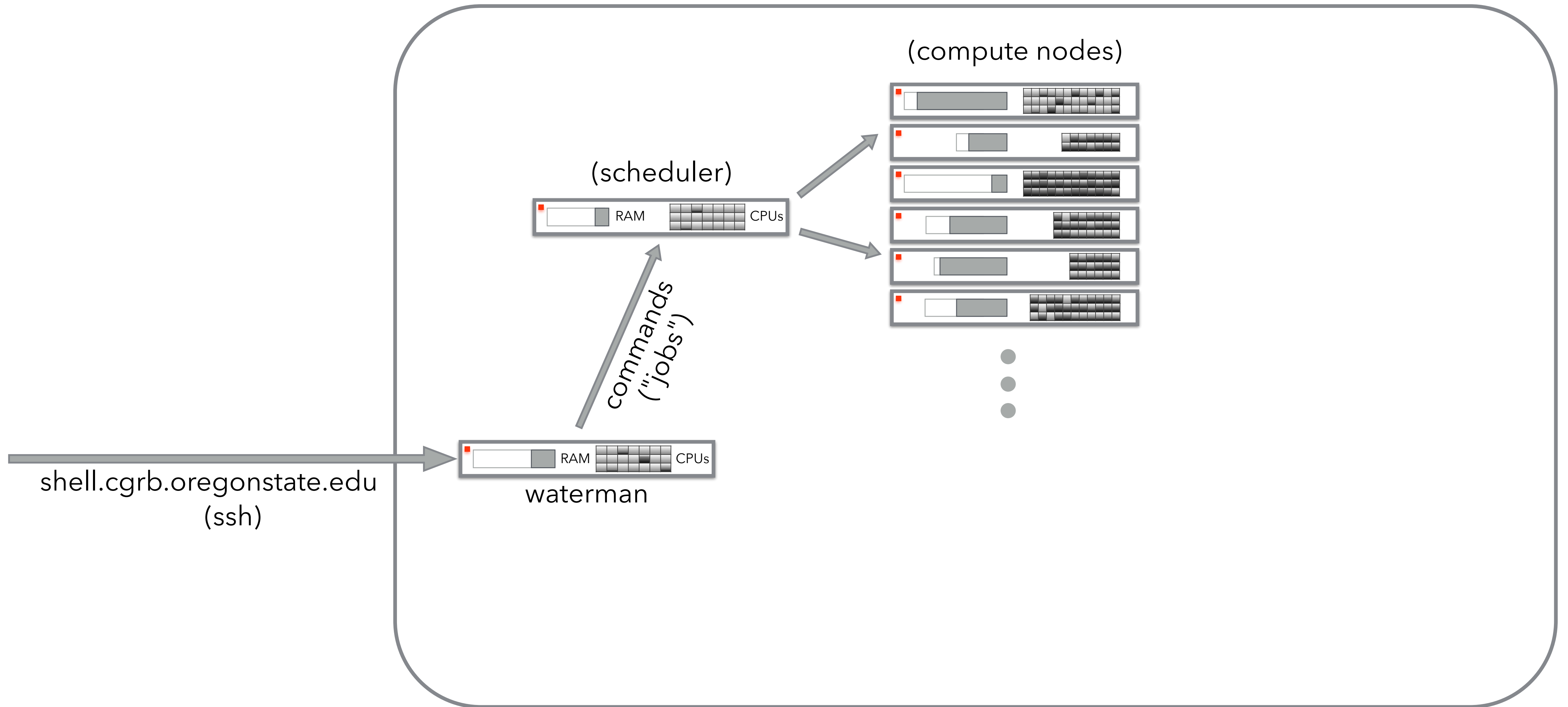
# CGRB Computational Infrastructure



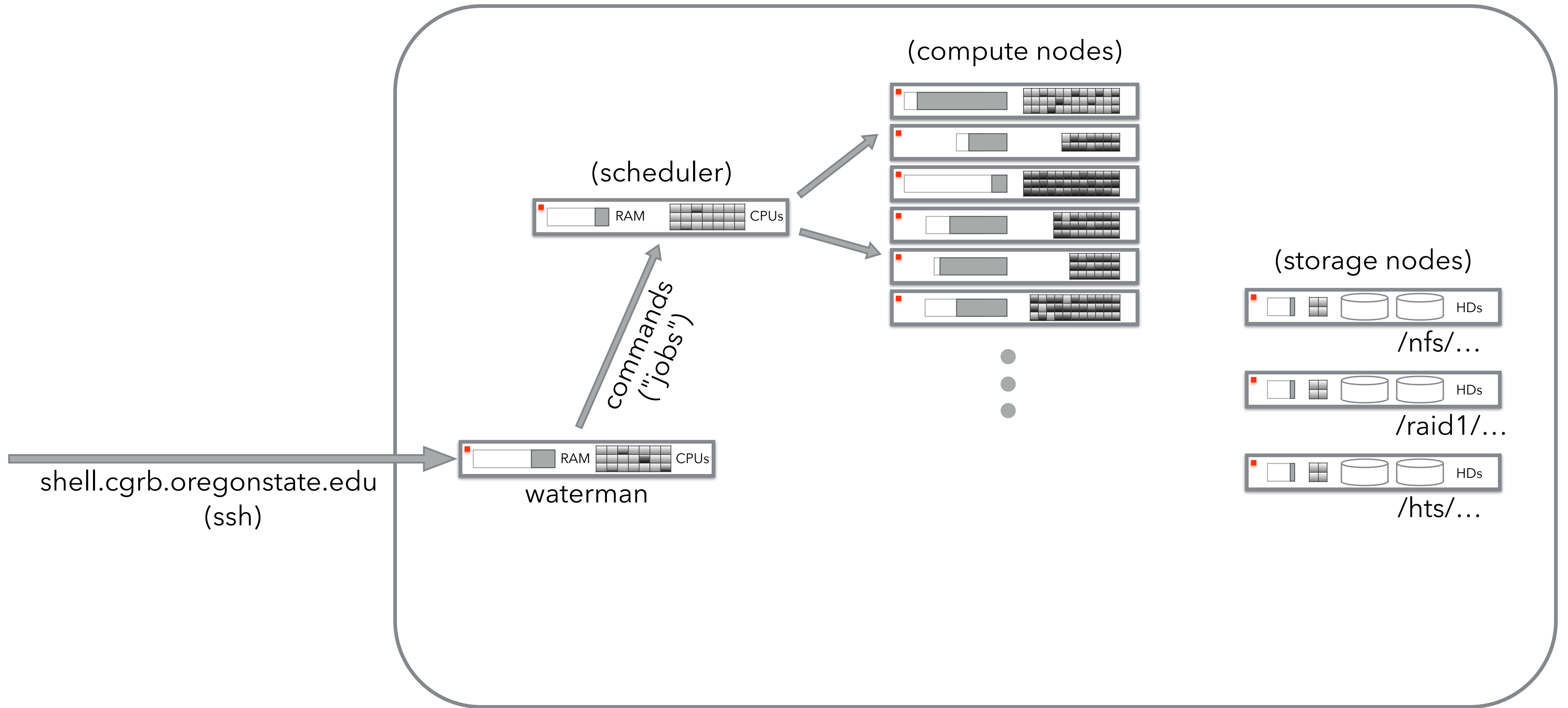
# CGRB Computational Infrastructure



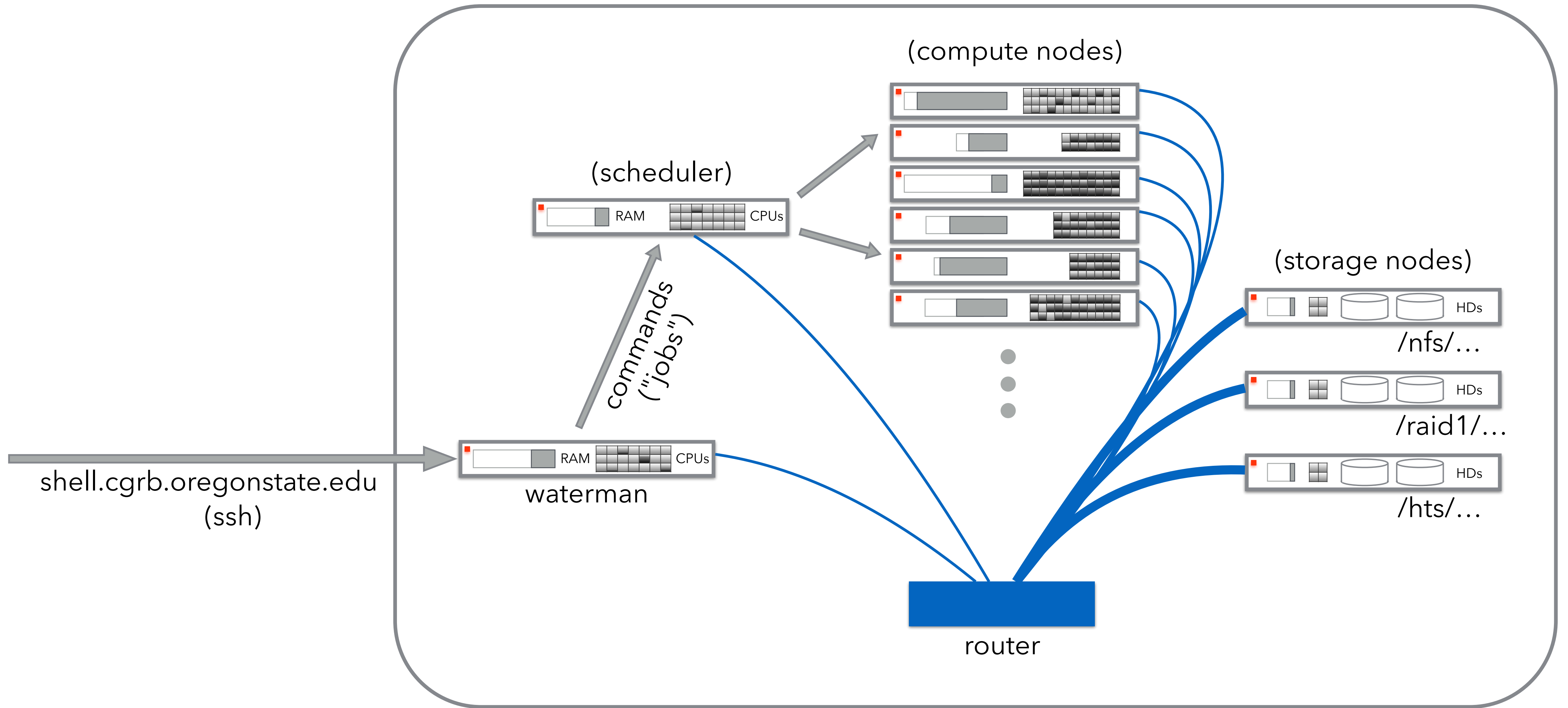
# CGRB Computational Infrastructure



# CGRB Computational Infrastructure

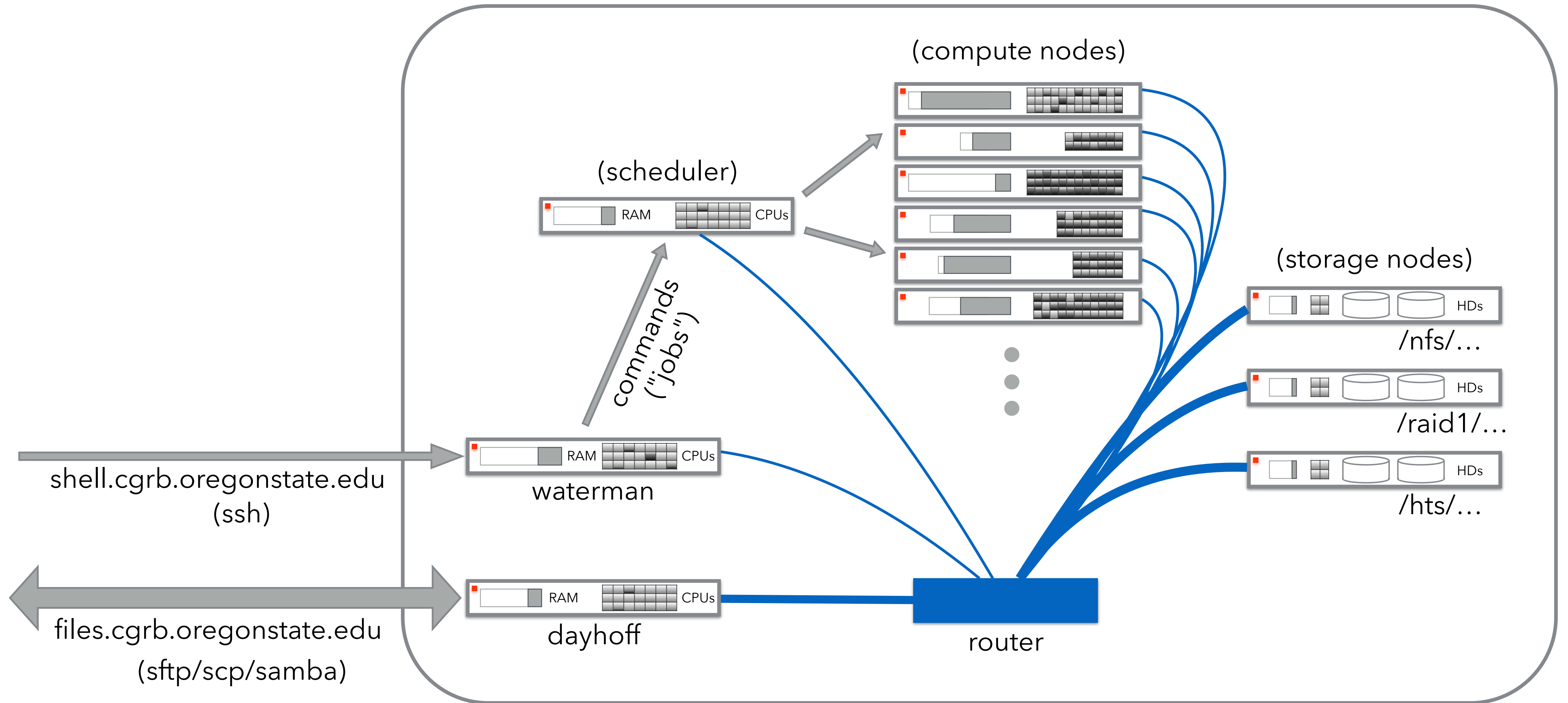


# CGRB Computational Infrastructure





# CGRB Computational Infrastructure



Kinda secret

```
[soneil@mbp ~]$ ssh -p XXX oneils@shell.cgrb.oregonstate.edu
[Linux@waterman ~]$ cd bug
/raid1/home/cgrb/oneils/bug
[Linux@waterman ~/bug]$ ls
[Linux@waterman ~/bug]$ date ; sleep 10 ; date > result.txt
Wed Jan 17 10:07:24 PST 2018
[Linux@waterman ~/bug]$ ls
result.txt
```

Some bioinformatics command  
that creates an output file

Create this directory for logs  
and printed output

My command (in quotes)

```
[Linux@waterman ~/bug]$ SGE_Batch -c 'date ; sleep 10 ; date > result.txt' -r log_dir -P 4 -m 10G
```

```
* Beginning the Data run  
  RunID = log_dir  
  Dir = log_dir
```

```
* Your job 2863594 ("log_dir") has been submitted
```

I'll need 4 CPUs and 10G RAM  
(only start job if this much is free when it starts;  
kill it if it tries to use more than this)

```
[Linux@waterman ~/bug]$ qstat
```

job-ID	prior	name	user	state	submit/start at	queue	slots	ja-task-ID
2863594	0.00000	log_dir	oneils	qw	01/17/2018 10:25:45		4	

```
[Linux@waterman ~/bug]$ qstat
```

job-ID	prior	name	user	state	submit/start at	queue	slots	ja-task-ID
2863594	0.50976	log_dir	oneils	r	01/17/2018 10:25:56	roots@roots1.cgrb.oregonstate.	4	

```
[Linux@waterman ~/bug]$ qstat
```

```
[Linux@waterman ~/bug]$ ls
```

```
log_dir/  result.txt
```

# How much time + RAM did that job take?

```
[Linux@waterman ~/bug]$ SGE_Plotdir log_dir
#RAM(Gigs) Time(Seconds)
0.000166893 0.00
[Linux@waterman ~/bug]$ cd log_dir/
/raid1/home/cgrb/oneils/bug/log_dir
[Linux@waterman log_dir]$ ls
log_dir_1_sge.sh  log_dir.e2863594  log_dir.o2863594  log_dir.pe2863594  log_dir.po2863594
[Linux@waterman log_dir]$ cat log_dir.e2863594
Full Command:          date
Memory (kb):           700
# SWAP (freq):         0
# Waits (freq):        3
CPU (percent):         50%
Time (seconds):        0.00
Time (hh:mm:ss.ms):    0:00.00
System CPU Time (seconds): 0.00
User CPU Time (seconds): 0.00
[Linux@waterman log_dir]$ cat log_dir.o2863594
Started on:             roots1
Started at:             Wed Jan 17 10:25:56 PST 2018
Wed Jan 17 10:25:56 PST 2018
Finished at:           Wed Jan 17 10:26:06 PST 2018
```

Inspect the “std error” log of the command (none to show here)

Inspect the “std output” log of the command (result of “date” printed)

# What do I have access to, right now?

```
[Linux@waterman ~/bug]$ SGE_Avail
```

#HOST	TOTRAM	FREERAM	TOTSLOTS	Q	QSLOTS	QFREESLOTS	QSTATUS	QTYPE
megraw3	503.7	503.7	64	megraw	64	64	au	BIP
megraw2	503.7	503.7	64	bpp	64	64	au	BIP
roots1	503.7	484.9	64	roots	64	56	normal	BIP
plant0	86.4	86.4	8	plant	8	8	au	BIP
otter	503.7	493.8	64	otter	64	7	normal	BIP

Machine roots1 has 503.7G of RAM, currently 484.9G available.

It has 64 total CPUs.

64 of those are allocated to the "roots" queue

56 of those are available

Machine + queue are in "normal" state (au or E: not good)

B: SGE\_Batch-able  
I: qrsh (login)-able  
P: multi-CPU (-P option)-able

## Resources available to burnsg at this moment

```
[Linux@waterman ~/bug]$ SGE_Avail -u burnsg
```

#HOST	TOTRAM	FREERAM	TOTSLOTS	Q	QSLOTS	QFREESLOTS	QSTATUS	QTYPE
beagle	913.2	900.9	64	beagle	64	56	normal	BIP
reef	125.7	62.8	48	reef	12	12	normal	BIP
chrom22	3.7	3.7	4	all.q	4	4	u	BP
chrom20	3.7	3.7	4	all.q	4	4	u	BP
chrom18	3.7	3.49	4	all.q	4	3	normal	BP
epi1	31.3	23.2	8	all.q	6	2	normal	BP

Practice...

epi1 has 6 CPUs in the all.q, 2 are available;  
notice no 'l' - *not* qrsh (login)-able

all.q: for everyone!

other queues: for specific lab, department  
members

Log me in interactively, somewhere

```
[Linux@waterman log_dir]$ qrsh
```

```
[Linux@roots1 ~]$ exit
```

```
logout
```

```
[Linux@waterman log_dir]$ qrsh -pe thread 8
```

```
[Linux@roots1 ~]$ SGE_Avail
```

#HOST	TOTRAM	FREERAM	TOTSLOTS	Q	QSLOTS	QFREESLOTS	QSTATUS	QTYPE
megraw3	503.7	503.7	64	megraw	64	64	au	BIP
megraw2	503.7	503.7	64	bpp	64	64	au	BIP
roots1	503.7	486.9	64	roots	64	48	normal	BIP
plant0	86.4	86.4	8	plant	8	8	au	BIP
otter	503.7	492.7	64	otter	64	7	normal	BIP

```
[Linux@roots1 ~]$ qstat
```

job-ID	prior	name	user	state	submit/start at	queue	slots	ja-task-ID
2885254	0.51611	QRLOGIN	oneils	r	01/17/2018 10:54:43	roots@roots1.cgrb.oregonstate.	8	

```
[Linux@roots1 ~]$ exit
```

```
logout
```

```
[Linux@waterman log_dir]$ qrsh -pe thread 8 -q roots@roots1
```

```
[Linux@roots1 ~]$
```

Same, but allocate me 8 CPUs

Same, but I specifically want the roots queue, on the roots1 machine

(-q roots would get me the roots queue on any machine that has it)

If using qrsh, do not forget to log back out!

You're telling the scheduler you are using those resources, so others can't.



I want to run a *lot* of jobs....

```
[Linux@waterman log_dir]$ head -n 5 lotsa_commands.txt
```

```
sleep 2; echo job1 > result1.txt  
sleep 2; echo job2 > result2.txt  
sleep 5; echo job3 > result3.txt  
sleep 1; echo job4 > result4.txt  
sleep 3; echo job5 > result5.txt
```

Per-job requirements

Similar to SGE\_Batch

Only run 10 at a time

```
[Linux@waterman ~/bug]$ SGE_Array -c lotsa_commands.txt -r log_dir -P 4 -m 10G -b 10
```

```
[Linux@waterman ~/bug]$ qstat
```

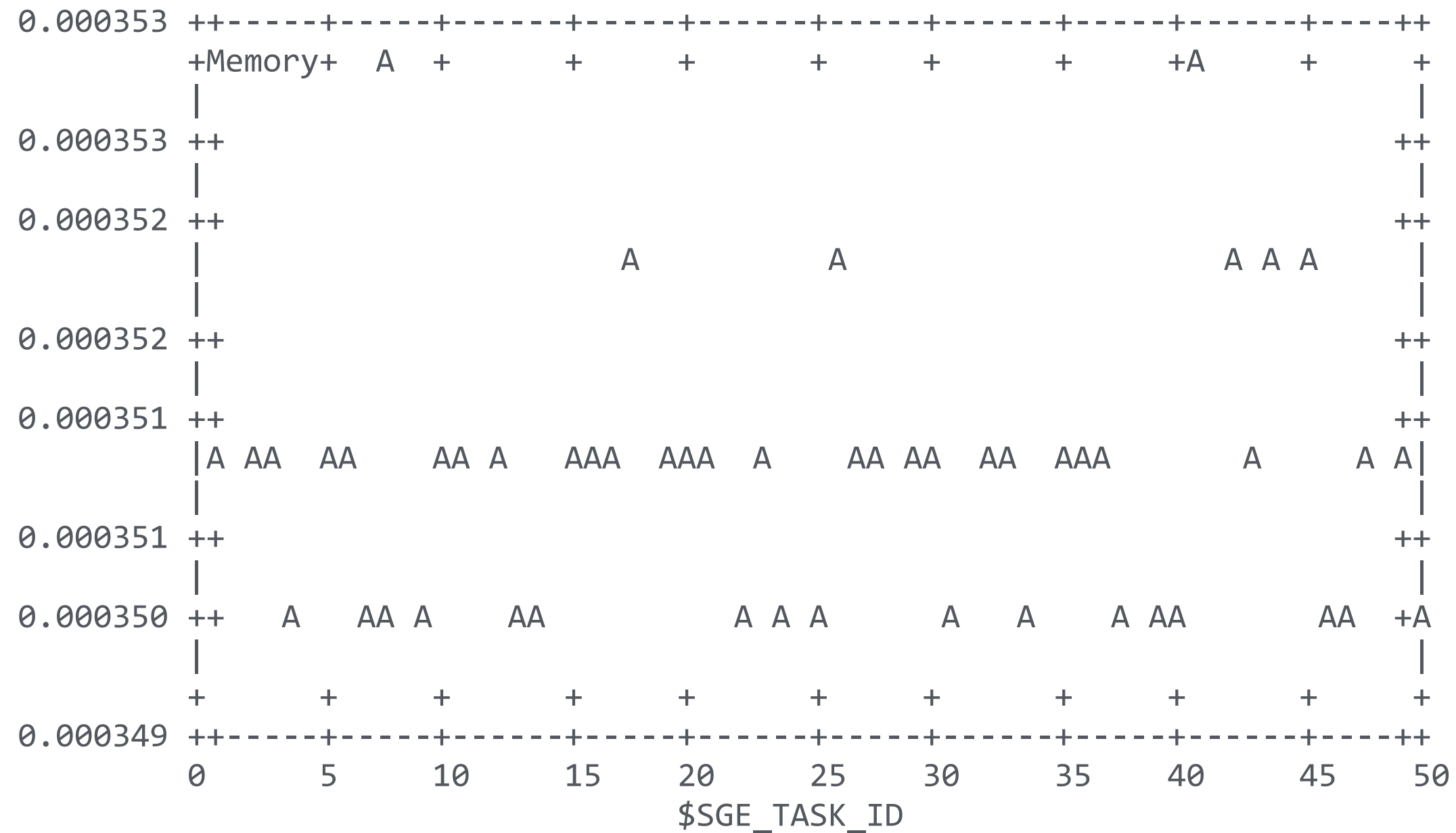
job-ID	prior	name	user	state	submit/start	at	queue	slots	ja-task-ID
2893513	0.50976	log_dir	oneils	r	01/17/2018	11:08:12	roots@roots1.cgrb.oregonstate.	4	1
2893513	0.50976	log_dir	oneils	r	01/17/2018	11:08:12	roots@roots1.cgrb.oregonstate.	4	2
2893513	0.50976	log_dir	oneils	r	01/17/2018	11:08:12	roots@roots1.cgrb.oregonstate.	4	3
2893513	0.50976	log_dir	oneils	r	01/17/2018	11:08:12	roots@roots1.cgrb.oregonstate.	4	4
2893513	0.50976	log_dir	oneils	r	01/17/2018	11:08:12	roots@roots1.cgrb.oregonstate.	4	5
2893513	0.50976	log_dir	oneils	r	01/17/2018	11:08:12	roots@roots1.cgrb.oregonstate.	4	6
2893513	0.50976	log_dir	oneils	r	01/17/2018	11:08:12	roots@roots1.cgrb.oregonstate.	4	7
2893513	0.50976	log_dir	oneils	r	01/17/2018	11:08:12	roots@roots1.cgrb.oregonstate.	4	8
2893513	0.50976	log_dir	oneils	t	01/17/2018	11:08:12	roots@roots1.cgrb.oregonstate.	4	9
2893513	0.50976	log_dir	oneils	t	01/17/2018	11:08:12	roots@roots1.cgrb.oregonstate.	4	10
2893513	0.00000	log_dir	oneils	qw	01/17/2018	11:08:07		4	11-50:1

10 Are running (or starting); the other 40 are waiting!

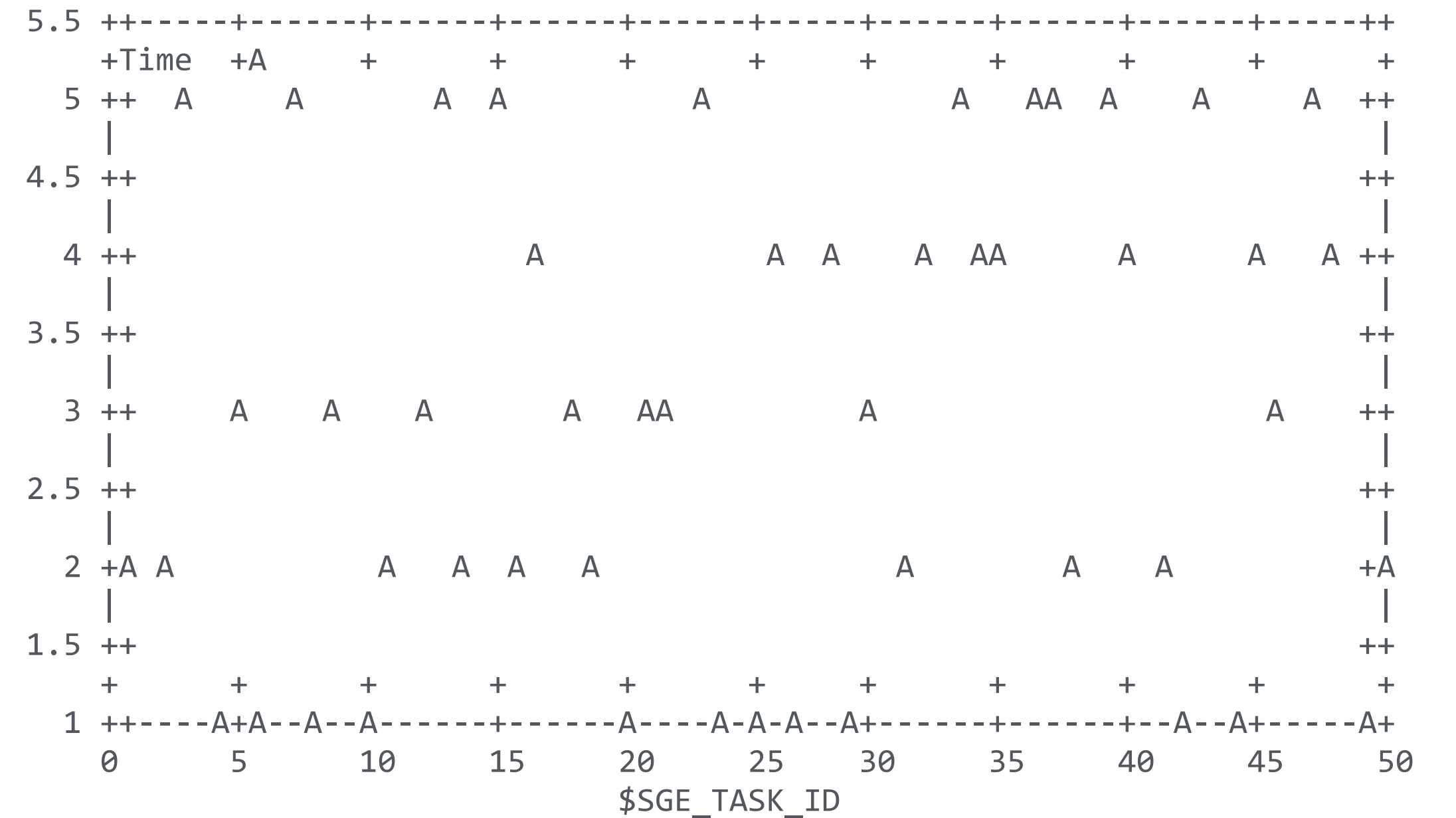
```
[Linux@waterman ~/bug]$ SGE_Plotdir log_dir
```

#ArrayJob#	GigsRAM	Seconds
1	0.000350952	2.00
2	0.000350952	2.00
3	0.000350952	5.00
4	0.000349998	1.00
5	0.000350952	3.00
...		

How much time/RAM did those jobs use?



RAM plot per job (gigs)



time plot per job (seconds)

Delete job # 2899950 (if it belongs to me)

```
[Linux@waterman ~/bug]$ qdel 2898850
```

Delete all my jobs (oh %\*!\$)

```
[Linux@waterman ~/bug]$ qdel -u oneils
```

See all jobs by all users

```
[Linux@waterman ~/bug]$ qstat -u '*'
```