



CPU

top ps

sar

**RAM** 

top

vmstat

free

Disk

df

du

Jobs

qstat

qconf

qacct

qdel

jobs

# Monitoring jobs: CPU



top: continuous look at processes

	15.51.02	6	400	22	50	4				warran A (	03, 0.13, 0.16
	615 tota										) zombie
	: 0.0%u					99.5				a, 0.0%hi	
	65957020										52k buffers
	51199996										204k cached
										,	
PID	USER	PR	NI	VIRT	RES	SHR	S	%CPU	<b>%MEM</b>	TIME+	COMMAND
3423	root	20	0	420m	231m	9.9m	S	0.7	0.4	27:26.44	splunkd
1345	dikowr	20	0	15436	1636	896	R	0.7	0.0	0:00.50	top
104	root	20	0	0	0	0	s	0.3	0.0	0:16.01	events/5
2374	root	20	0	18372	812	468	S	0.3	0.0	5:00.50	irqbalance
34494	root	20	0	0	0	0	S	0.3	0.0		
1	root	20	0	19348	1504	1180	s	0.0	0.0	0:01.80	init
2	root	20	0	0	0	0	s	0.0	0.0		kthreadd
3	root	RT	0	0	0	0	S	0.0	0.0		migration/0
4	root	20	0	0	0	0	S	0.0	0.0	0:02.81	ksoftirqd/0
5	root	RT	0	0	0		s	0.0	0.0		stopper/0
6	root	RT	0	0	0		s	0.0	0.0		watchdog/0
7	root	RT	0	0	0		s	0.0	0.0		migration/1
	root	RT	0	0	0		S	0.0	0.0		stopper/1
	root	20	0	0	0		s	0.0	0.0		ksoftirqd/1
	root	RT	0	0	0		S	0.0	0.0		watchdog/1
		RT	0	0	0		S	0.0	0.0		migration/2
12	root	RT	0	0	0		S	0.0	0.0		stopper/2
	root	20	0	0	0		S	0.0	0.0		ksoftirqd/2
	root	RT	0	0	0		S	0.0	0.0		watchdog/2
	root	RT	0	0	0		S	0.0	0.0		migration/3
	root	RT	0	0	0		S	0.0	0.0		stopper/3
	root	20	0	0	0		s	0.0	0.0		ksoftirgd/3
	root	RT	0	0	0		S	0.0	0.0		watchdog/3
	root	RT	0	0	0		S	0.0	0.0		migration/4
	root	RT	0 0	0	0		S	0.0	0.0		stopper/4
	root	20	0	0	0		8	0.0	0.0		ksoftirqd/4
22	root	$\mathbf{R}^{\mathbf{T}}$	U	0	U	0	s	0.0	0.0	0:00.40	watchdog/4

ps: snapshot of processes

# Monitoring jobs: CPU



sar: System Activity Report, part of Sysstat

	01075 10								
000					SAR.txt				
03.30.01		acc	22.27	0.00	0.01	0.00	0.00	77.75	
06:00:01		all	22.09	0.00	0.09	0.00	0.00	77.82	
06:10:01		all	22.23	0.00	0.01	0.00	0.00	77.76	
06:20:01	PM	all	22.09	0.00	0.09	0.00	0.00	77.82	
06:20:01	PM	CPU	%user	%nice	%system	%iowait	%steal	%idle	
06:30:01	PM	all	22.24	0.00	0.01	0.00	0.00	77.76	
06:40:01	PM	all	22.08	0.00	0.09	0.00	0.00	77.83	
06:50:01	PM	all	22.18	0.00	0.02	0.00	0.00	77.80	
07:00:01	PM	all	22.14	0.00	0.08	0.00	0.00	77.78	
07:10:01	PM	all	22.08	0.00	0.09	0.00	0.00	77.83	
07:20:01	PM	all	22.24	0.00	0.01	0.00	0.00	77.76	
07:30:01	PM	all	13.38	0.00	0.01	0.00	0.00	86.61	
07:40:01	PM	all	0.69	0.00	0.01	0.00	0.00	99.30	
07:50:01	PM	all	7.49	0.00	0.08	0.00	0.00	92.43	
08:00:02	PM	all	22.21	0.00	0.03	0.00	0.00	77.75	
08:10:01	PM	all	22.11	0.00	0.09	0.00	0.00	77.80	
08:20:01	PM	all	22.24	0.00	0.01	0.00	0.00	77.76	
08:30:01	PM	all	22.09	0.00	0.09	0.00	0.00	77.82	
08:40:01	PM	all	22.24	0.00	0.01	0.00	0.00	77.75	
08:50:01	PM	all	22.08	0.00	0.09	0.00	0.00	77.83	
09:00:01	PM	all	22.24	0.00	0.01	0.00	0.00	77.75	
09:10:01	PM	all	22.09	0.00	0.09	0.00	0.00	77.82	
09:20:01	PM	all	22.24	0.00	0.01	0.00	0.00	77.76	
09:30:01	PM	all	22.08	0.00	0.09	0.00	0.00	77.83	
09:40:01	PM	all	22.24	0.00	0.01	0.00	0.00	77.76	
09:50:01	PM	all	22.09	0.00	0.09	0.00	0.00	77.81	
10:00:01	PM	all	22.24	0.00	0.01	0.00	0.00	77.75	
10:00:01	PM	CPU	%user	%nice	%system	%iowait	%steal	%idle	
10-10-01	DM	211	22 10	0 00	0 00	0 00	a aa	77 81	

Can access for a particular day or set of days in the past month sar -b -f /var/log/sa/sa20 (for the 20th of the month) syntax varies by system

# Monitoring jobs: RAM



vmstat: get info on memory usage

```
dikowr@hydra-3% vmstat
procs -----memory----- ---swap-- ----io---- --system-- ----cpu-----
                     buff
                                            bi
                                                       in
                                                            cs us sy id wa st
              free
                                  si
                                                  bo
r b
       swpd
                          cache
                                       so
          0 60039684 890464 3309216
                                                1
                                           0
                                                           3
                                      0
```

#### free: similar to vmstat, output looks a bit different

```
[dikowr@login-3-1 ~]$ free
                                                         buffers
                                                                      cached
             total
                                               shared
                          used
                                      free
          32998248
                       2193796
                                                  224
                                                          271480
                                                                     1162592
                                 30804452
Mem:
-/+ buffers/cache:
                        759724
                                 32238524
Swap:
          31211516
                                 31211516
```

# Monitoring jobs: disk



disk free: [-h] display in KB, MB, or GB df

disk usage: [-sh] sum of directories du

## Monitoring jobs: Jobs 💯



qstat

qstat -f job\_id

Produce detailed report about job

qstat -u dikowR

To list all the jobs belonging to a particular user (me)

qstat -q

To view a summary of all queues

qstat -j

Prints info about pending jobs



Hydra website:

https://www.cfa.harvard.edu/~sylvain/hpc/status/



qconf get info on queue configuration

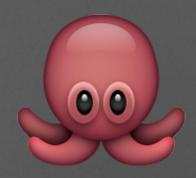
qdel delete a job (e.g. \$qdel job-id)

qacct report on SGE usage

jobs display status of jobs in foreground/ background



How to use shell scripting with Hydra to be productive in your job submission



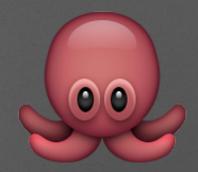
```
#!/bin/bash
for i in *.nex; do

    qsub -q mThC.q -N MrBayes_$i -S /
bin/sh -cwd -j y -o MrBayes_$i.out -l
mres=1G,h_data=1G,h_vmem=1G mb.job $i

done
```

check out:

man qsub



```
mb $1
```

# Compute intensive RAM intensive I/O intensive



#### Compute intensive



- -Limitation is processing
- -How to parallelize: decompose the overall application process into separate tasks, which can then be executed in parallel

Common examples:

Phylogenetic tree search

Alignment

#### I/O (data) intensive



- -Spends more time reading and writing than processing
- -Should scale linearly according to the size of the data
- -How to parallelize: split the data into parts that can be processed independently and then re-assemble

Common examples:

**BLAST** 

lots of little assemblies

#### RAM intensive



-Limitation is memory

-Will likely crash if not enough RAM, more time won't help

Common examples:

Genome/transcriptome assembly (~300GB for 1Gbp genome)

Phylogenetic tree building on a very\* large matrix (# of taxa vs. # of characters)

#### Resources



Google

**Biostars** 

**Stack Overflow** 

Individual program email lists/google groups

#### Resources



https://hydra-3.si.edu/wordpress/

https://hydra-3.si.edu/ganglia/

http://si-vmatlassian:8090/display/ORIS/ Migration+Notes%3A+Hydra-2+to +Hydra-3

http://si-vmatlassian:8090/display/ORIS/ Hydra+Software

