

# How to deal with common Scheduling Problems

# Troubleshooting, Make sure you gather data that may be lost

- Keep any error messages you received.
- What environment setting were set, which module have you loaded when you submitted the job?
- When did the problem occur?
- The state of a cluster changes, jobs start and finish, nodes fail and are repaired. What were the output of commands that you ran on the system that make you think there is a problem or that you think analyst should see.

Keep a record of any output with script command.

“script <filename to write>”

Do your work

“exit” to stop writing the script.

If you need to add more information to the file

Use “Script –a <filename>”

**tip:** Put a date and time in the file name

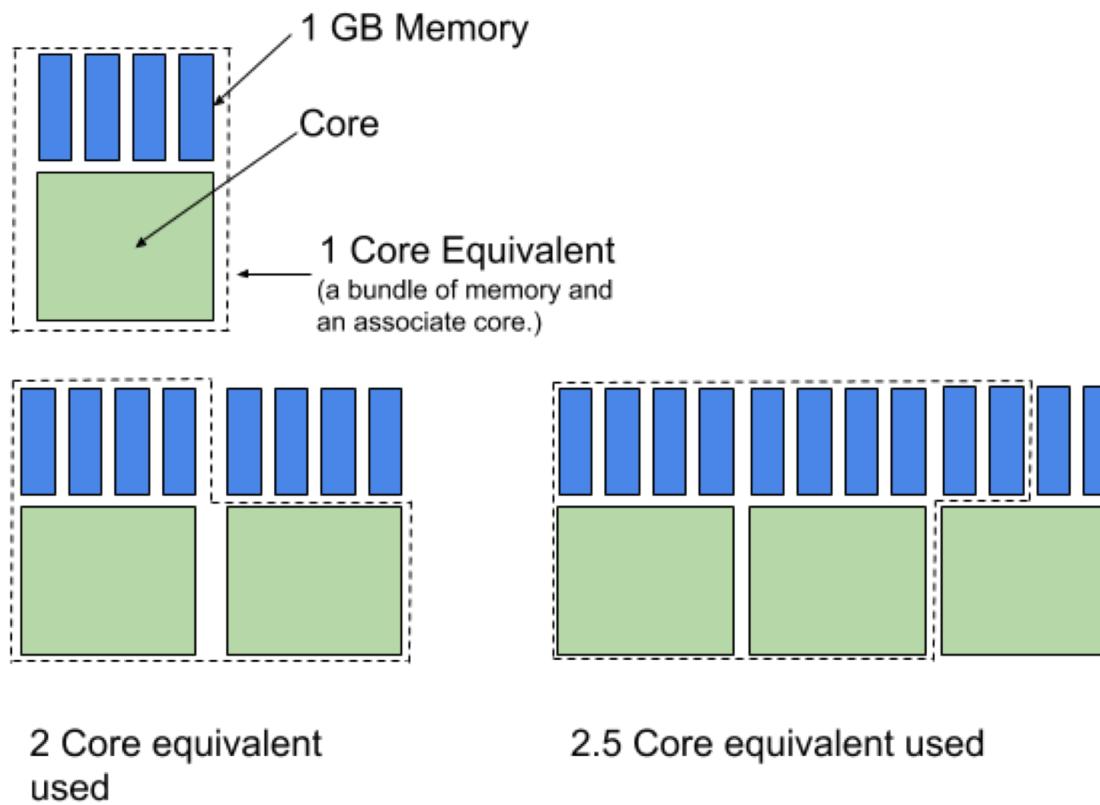
# Memory Request Problems

- The Memory your jobs requests is what the system must have available for your job to use.
  - If your requests to use 1.5 TiB it will never run on a node with 1.5TiB of RAM as some RAM will be used by the operating system, and there will not be enough resources to run your job.
  - **Recommended** that all request for RAM in 1000's of MiB
    - $1.5\text{TiB}=1,572,864 \text{ MiB}$  and if you request 1,500,00 MiB there would be 72,864 MiB to run the OS and services.
    - This recommendation is also good for smaller single core jobs as well, On a 32 core 128 GiB RAM node, the scheduler can fit 31 jobs asking for 1 core and 4GiB of RAM or 32 jobs asking for 1 core an 4000MiB.

# Running out of Memory

- Ask for more memory
- Be careful don't ask for too much
  - Asking for more memory makes your job more difficult to schedule.
  - CC cluster only have 4GB RAM per core on most nodes.
  - You can ask for more memory than the cluster has
  - Your group will be assessed as having used more resources, (for the purpose priority and allocation)

# Core Equivalent



Compute Canada core equivalent [documentation](#)

# Node types on Cedar - Compute

Number of Nodes	% of total	Memory per core (GiB)	Total Mem (GiB)	Cores	GPUS	Partition type
1408	73.64%	4	192	48		cpubase
576	20.08%	4	128	32		
128	4.46%	8	256	32		
24	0.84%	16	512	32		cpularge
24	0.84%	48	1536	32		
4	0.14%	96	3072	32		

# Node types on Graham - Compute

Number of Nodes	% of total	Memory per core (GiB)	Total Mem (GiB)	Cores	GPUS	Partition type
884	85.35%	4	128	32		cpubase
72	9.36%	4.2	192	44		
56	5.79 %	8	256	32		
24	2.48 %	16	512	32		cpularge
3	0.31 %	96	3072	32		

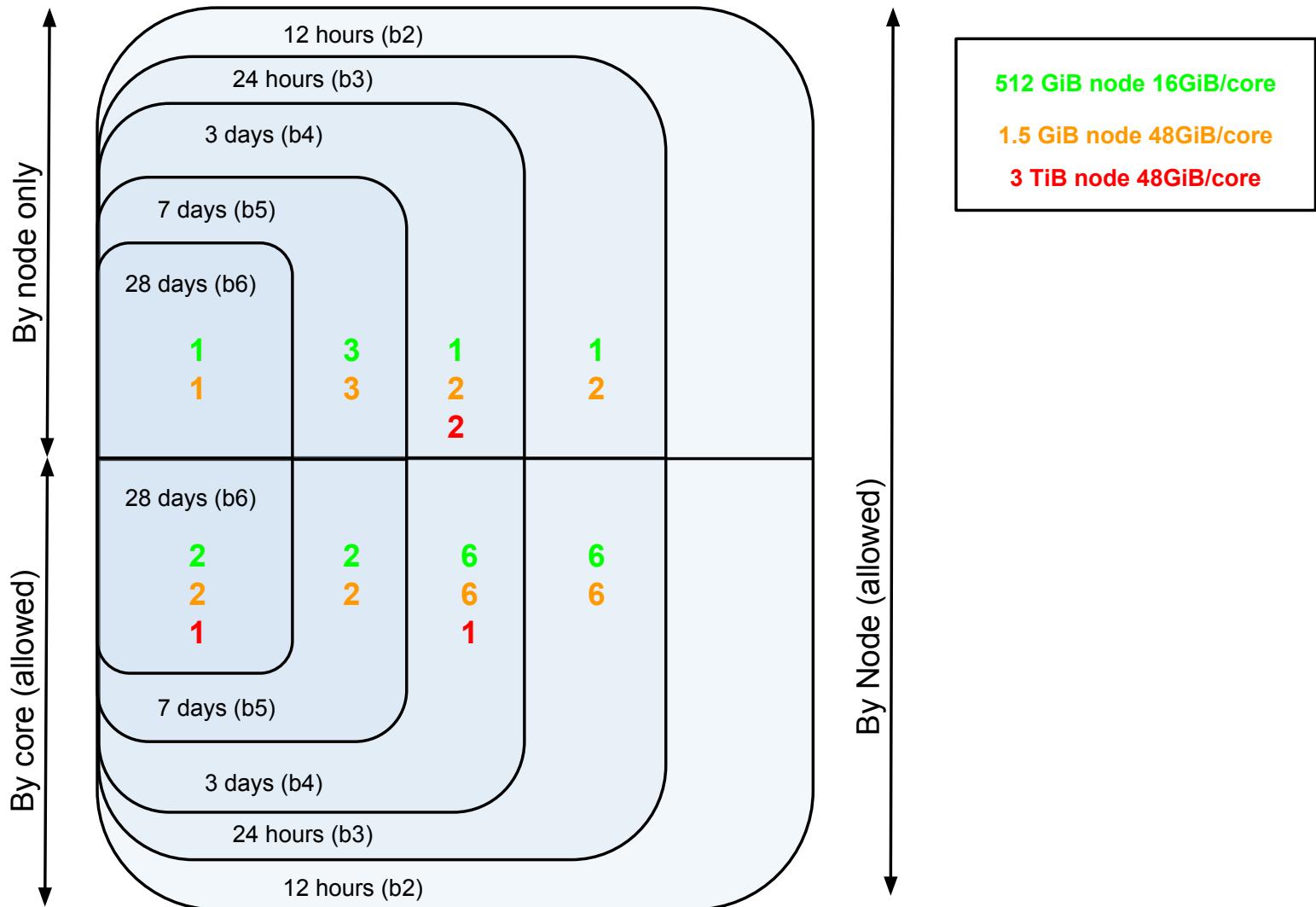
# Node types on Beluga - Compute

Number of Nodes	% of total	Memory per core (GiB)	Total Mem (GiB)	Cores	GPUS	Partition type
172	24.57%	2.4	96	40		cpubase
516	73.71%	4.8	192	40		
24	1.71%	19.2	768	40		cpularge

# Node types on Niagara - Compute

Number of Nodes	% of total	Memory per core (GiB)	Total Mem (GiB)	Cores
2016	100%	4.7	188	40

# Sept 2018 Setup large mem partitions on Cedar



# Virtual and Physical memory

- Your program can ask to use a chunk of memory. This is virtual or requested MaxVMSIZE
- MaxRSS is the amount of memory used by your code.

# Find out how much memory a job used.

Command	Flags	What its used for
sstat		Display various status information of a running job
	-j <jobid>	Displays information about the specified job
	--format= AveCPU,MaxRSS,MaxVMSize,JobID	limits the information to that about memory (MaxVMSize is requested memory) (MaxRSS is memory used)
sacct		Displays slurm accounting data
	-j <jobid>	Displays information about the specified job
	-u \$USER	Displays information about jobs belong to a specific user
	--format= JobID,AveCPU,MaxRSS,MaxVMSize	limits the information to that about memory
salloc		Submit to run Job Interactively
	Same flags as sbatch	Note not all sbatch flags work

# Interactive Jobs for debugging

Use salloc instead of sbatch to launch interactive jobs.

```
salloc --ntasks=4 --mem-per-cpu 4000 -t 0-00:20
```

List environment variables with printenv

```
printenv | grep ^S
```

To check memory usage use

```
top -u $USER
```

Look at the jobs cgroup,

```
cat  
/sys/fs/cgroup/cpuset/slurm/uid_${SLURM_JOB_UID}/job_${SLURM_JOB_ID}/  
step_${SLURM_STEPID}/tasks
```

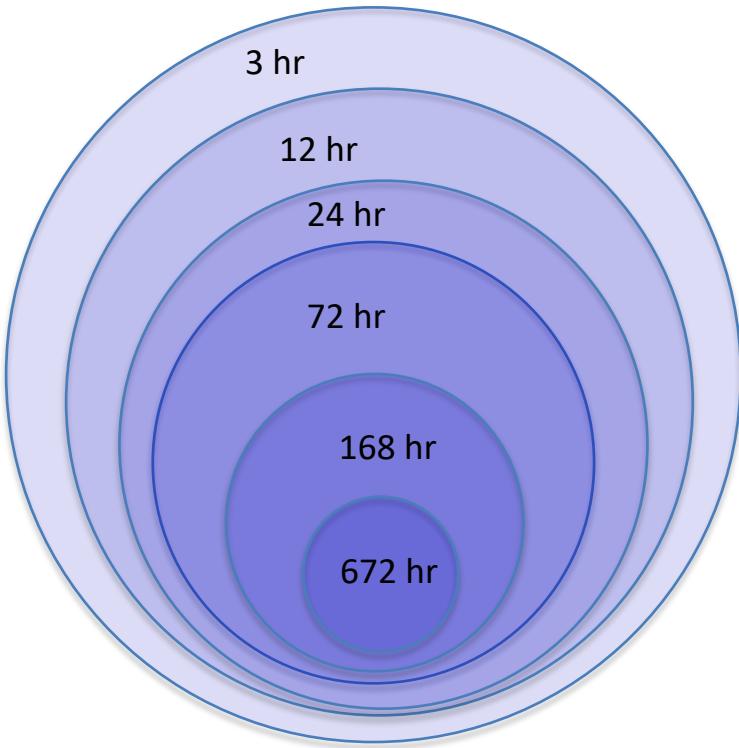
Remember that you are logged in only on one machine  
and your job may span more than one

# Running out of runtime

If your jobs are running out of time.

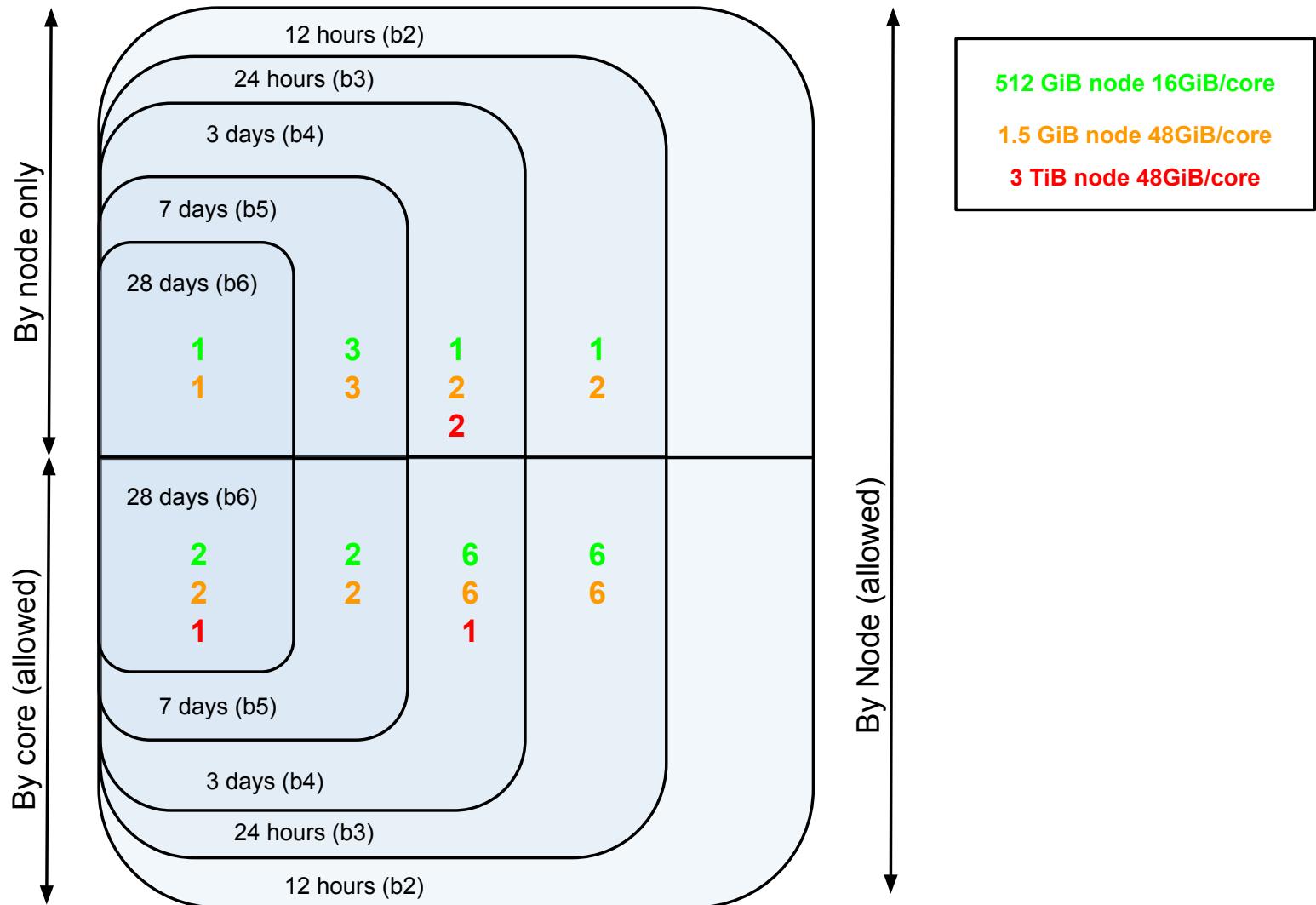
- Ask for more time.
- Don't ask for too much runtime.
- Asking for more runtime may limit you to how many resources can run your job. This may interact with how much memory you asked for.

# Partitions on Cedar and Graham



- There are partitions based upon how long the maximum walltime your job has.
- Your job ends up in the shortest walltime partition that has a longer walltime than your job
- The shorter walltime partitions include all the nodes of longer walltime partitions.

# Sept 2018 Setup large mem partitions on Cedar



# Partition Stats

## (CC script)

Node type	Max walltime					
	3 hr	12 hr	24 hr	72 hr	168 hr	672 hr
-----						
Number of Queued Jobs by partition Type (by node:by core)						
Regular	1:15	2:31	2:145	11:187	86:69	3:2
Large Mem	0:0	0:0	0:0	0:0	0:1	0:1
GPU	0:1	0:526	10:10	0:0	189:4	0:0
-----						
Number of Running Jobs by partition Type (by node:by core)						
Regular	60:6	4:2	45:836	5:90	11:1065	1:4
Large Mem	0:0	0:0	0:0	0:0	0:0	1:0
GPU	0:20	2:10	13:2	0:0	0:0	0:3
-----						
Number of Idle nodes by partition Type (by node:by core)						
Regular	0:0	0:0	0:0	0:0	0:0	0:0
Large Mem	3:1	0:0	0:0	0:0	0:0	0:0
GPU	17:1	11:1	0:0	0:0	0:0	0:0
-----						
Total Number of nodes by partition Type (by node:by core)						
Regular	851:411	821:391	756:346	636:276	180:100	90:50
Large Mem	27:12	24:11	24:11	20:3	3:2	2:1
GPU	156:78	144:72	116:58	104:52	13:12	13:12

# My job is not running right now

- `scontrol show job <jobid>`
  - pay special attention to **JobState**, and **Reason**
- Look at resource availability with CC's `partitions-stats` command, make sure you look up the resources the resources available to your jobs memory and time requirements.
- Looks at the jobs priority with the `sprio` command.

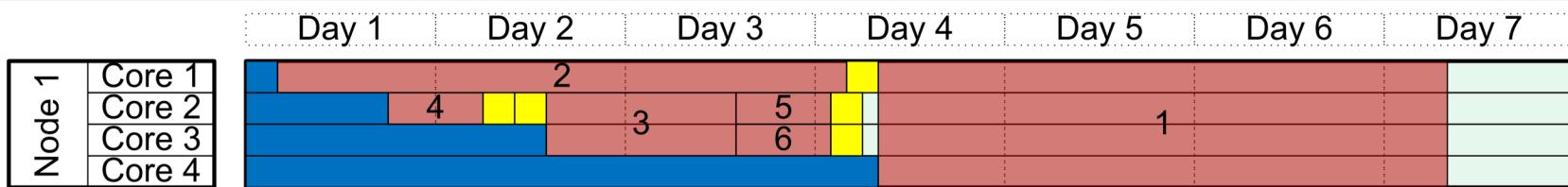
# Fixing priority

- Priority is based upon your groups recent usage compared to allocation in the recent past.
  - Wait and ask group members to use less resources.
  - Ask your PI to apply for a RAC allocation.
  - Change your job to ask for resources that are in less demand. Ex: checkpointing.

# Checkpointing

- Some programs can save its state and restart from save state.
- This allows long running jobs to be broken up into smaller runs.
- This minimizes effect of hardware failures, system downtime etc... on your ability to get your work done. This is particularly important for simulations using large amount of hardware.
- If your application does allow this make sure to understand how long a checkpoint and or restart takes. If it takes a long time to do, don't do it very often.

Big high priority jobs in the scheduler leave “holes” that can be filled with smaller shorter Jobs



# How to ask support for help

- Read status page and any support notices  
<http://status.computecanada.ca/>.
- Being able to ask for help in a helpful manner will likely result in your questions being help in a much more responsive manner.
- An email “Something is wrong”, “Nothing works” will take a long time to resolve.
  - A analyst responsible for a system will fix problems with the system they are responsible for, before they start answering asking you questions in order to determine which system you are having a problem. The problem could be with the cluster, cloud, website, network etc..
- In the subject of the email, include the system/cluster name a a few words of what may be wrong.
  - “job 123456 fails to run on the Cedar cluster” will much more likely to handled by a person who can help quickly.

# Don't make the Analysts play detective unnecessarily

- Compute Canada has multiple clusters.
- Your Compute Canada user name may not be apparent from your email.
- You may have 1000s of job queued, running, completed, failed on the system with which one did you have an issue?
- When did this happen?
- Which jobsript did you launch your job with, have you modified it since?
- What version of software are you running?

TO: support@computeccanda.ca

Subject: Job 123456 gives errors on the CC Cedar cluster

Hello, My name is Alice, user asmith . Today at 10:00 am MST I submitted a job 123456 on the Cedar cluster. The Job script is located /my/job/script/path I have not changed it since submitting my job, since it is short I included it in the email bellow.

```
#!/bin/bash
#SBATCH --account=def-asmith-ab
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=16
#SBATCH --time=00:05:00
{ time mpiexec -n 1 ./sample1 ; } 2>out.time
```

A list of the following modules were loaded at the time follow:

```
[asmith@cedar5]$ module list
Currently Loaded Modules:
 1) nixpkgs/16.09 (S)  5) intel/2016.4 (t)
 2) icc/.2016.4.258 (H)  6) imkl/11.3.4.258 (math)
 3) gcccore/.5.4.0 (H)  7) openmpi/2.1.1 (m)
 4) ifort/.2016.4.258 (H)  8) StdEnv/2016.4 (S)
```

The job ran quickly and the myjob-123456.out and myjob-123456.err files were created. There was no output in the myjob-123456.out file but there was an message in the myjob-123456.err output

```
[asmith@cedar5 scheduling]$ cat myjob-123456.err
slurmstepd: error: *** JOB 123456 ON cdr692 CANCELLED AT 2018-09-06T15:19:16 DUE TO TIME LIMIT ***
```

Can you tell me how to fix this problem?

# User debugging questions

Any questions.

If there is time and we have a volunteer who consents for others to listen and learn and that there will be a recording made of this and posted on line. We can do a live debugging session example.

The end

Material bellow will not be in the presentation unless diagram is needed to answer a question.

# **SLURM SCRIPT REFERENCE MATERIAL BELOW**

# Basic Slurm script commands

Slurm script command	Description
<code>#!/bin/bash</code>	Sets the shell that the job will be executed on the compute node
<code>#SBATCH --ntasks=1</code> <code>#SBATCH --n1</code>	Requests for 1 processors on task, usually 1 cpu as 1 cpu per task is default.
<code>#SBATCH --time=0-05:00</code> <code>#SBATCH -t 0-05:00</code>	Sets the maximum runtime of 5 hours for your job
<code>#SBATCH --mail-user= &lt;email&gt;</code>	Sets the email address for sending notifications about your job state.
<code>#SBATCH --mail-type=BEGIN</code> <code>#SBATCH --mail-type=END</code> <code>#SBATCH --mail-type=FAIL</code> <code>#SBATCH --mail-type=REQUEUE</code> <code>#SBATCH --mail-type=ALL</code>	Sets the scheduling system to send you email when the job enters the following states: BEGIN,END,FAIL,REQUEUE,ALL
<code>#SBATCH --job-name=my-named-job</code>	Sets the Jobs name

# Slurm script commands

Slurm script command	Description
#SBATCH --ntasks=X	Requests for X tasks. When cpus-per-task=1 (and this is the default) this requests X cores. When not otherwise constraint these CPUs may be running on any node
#SBATCH --nodes=X	Request that a minimum of X nodes be allocated to this job
#SBATCH --nodes=X-Y	Request that a minimum of X nodes and a maximum of Y nodes be allocated to this job
#SBATCH --cpus-per-task=X	Request that a minimum of X CPUs per task be allocated to this job
#SBATCH --tasks-per-node=X	Requests minimum of X task be allocated per node

# Slurm script commands

Slurm script commands	Description of effects
#SBATCH --ntasks=1 #SBATCH --cpus-per-task=1	Requests 1 CPU (Serial) cpus-per-task is set to 1 by default and may be omitted.
#SBATCH --cpus-per-task=X #SBATCH --ntasks=1 #SBATCH --nodes=1	Requests for X CPUs in 1 task on 1 node (OpenMP) Both ntasks and nodes are set to 1 by default and may be omitted
#SBATCH --ntasks=X #SBATCH --tasks-per-node=X #SBATCH --cpus-per-task=1	Requests for X CPUs and tasks on 1 node cpus-per-task is set to 1 by default and may be omitted.
#SBATCH --ntasks=X #SBATCH --nodes=1 #SBATCH --cpus-per-task=1	Requests for X CPUs and tasks on 1 node cpus-per-task is set to 1 by default and may be omitted.

# Slurm script commands

Slurm script commands	Description of effects
#SBATCH --ntasks=X #SBATCH --cpus-per-task=1	Requests X CPUs and tasks (MPI) cpus-per-task is set to 1 by default and may be omitted.
#SBATCH --ntasks=X #SBATCH --ntasks-per-node=Y #SBATCH --cpus-per-task=1	Requests for X CPUs and tasks with Y CPUs and tasks per node cpus-per-task is set to 1 by default and may be omitted.
#SBATCH --ntasks=X #SBATCH --nodes=1 #SBATCH --cpus-per-task=1	Requests for X CPUs and tasks on the same node, cpus-per-task is set to 1 by default and may be omitted.
#SBATCH --ntasks=X #SBATCH --nodes=1 #SBATCH --cpus-per-task=1	Requests for X CPUs and tasks on the 1 node cpus-per-task is set to 1 by default and may be omitted.

# Slurm script commands

Slurm script command	Description
#SBATCH --ntasks=1 #SBATCH --cpus-per-task=1	Requests 1 cpu in 1 task. (Serial) cpus-per-task is set to 1 by default and may be omitted.
#SBATCH --cpus-per-task=N #SBATCH --ntasks=1 #SBATCH --nodes=1	Requests for X processors on the same node (OpenMP) Both ntasks and nodes are set to 1 by default and may be omitted
#SBATCH --ntasks=X	Requests for X processors which may be running on any node (MPI).
#SBATCH --nodes=X #SBATCH --ntasks=Y	Requests minimum of X nodes for the Y tasks. ( MPI job)
#SBATCH --array=0-4	Requests Job array of 5 jobs with indexes 0-4
#SBATCH --array=1,3,5,7,9	Requests Job array of 5 jobs with indexes 1,3,5,7,9
#SBATCH --array=0-X%Y ex: #SBATCH --array=0-4%2	Requests Requests Job array of X jobs with only a maximum of Y jobs running at the same time

# Slurm script commands

PBS script command	Description
#SBATCH --mem=4000	Requests 4000 MB of memory in total
#SBATCH --mem-per-cpu=4000	Requests 4000 MB of memory per cpu
#SBATCH --licenses=sas:2	Requests 2 SAS licenses
#SBATCH --gres=gpu:1	Requests that your job get 1 GPU allocated per node
#SBATCH --exclusive	Requests that your job run only on nodes with no other running jobs
#SBATCH --dependency=after:job_id1	Requests that the the job start after job (jobid1) has <b>started</b>
#SBATCH --dependency=afterany:job_id1, job_i2	Requests that the the job start after ether job (jobid1) or job (jobud2) has <b>finished</b>
#SBATCH --dependency=afterok:job_id1	Requests that the the job start after job (jobid1) has <b>finished successfully</b>

# Slurm script commands

PBS script command	Description
#SBATCH --account=acc_name	To submit a job to a specific accounting group such as RAC/RAS allocation or different role
#SBATCH --tmp=200G	Asks for 200Gb of temporary disk space
#SBATCH --constraint=blue	To ask for a node feature or constraint set by cluster admin. Here we are looking for “blue” nodes.
#SBATCH --partition=partition_name	To ask for the job to run in a specific partition or queue by name, (unlike Moab there can be more than 1 partition per Job)
--prolog=<executable>	Run by srun only, runs the executable before the step
--epilog=<executable>	Run by srun only, runs the executable after the step finishes

# SLURM Environment Variables

Environment Variable	Description
SLURM_JOB_NAME	User specified job name
SLURM_JOB_ID	Unique slurm job id
SLURM_NNODES	Number of nodes allocated to the job
SLURM_NTASKS	Number of tasks allocated to the job
SLURM_ARRAY_TASK_ID	Array index for this job
SLURM_ARRAY_TASK_MAX	Total number of array indexes for this job
SLURM_MEM_PER_CPU	Memory allocated per CPU
SLURM_JOB_NODELIST	List of nodes on which resources are allocated to Job
SLURM_JOB_CPUS_PER_NODE	Number of CPUs allocated per Node
SLURM_JOB_PARTITION	List of Partition(s) that the job is in.
SLURM_JOB_ACCOUNT	Account under which this job is run.

# Getting information on your Job

Command	What its used for
squeue -u <username>	List all current jobs for a user
squeue -u <username> -t PENDING	List all pending jobs for a user
squeue -u <username> -t RUNNING	List all running jobs for a user
squeue -p <partitionname>	List all the jobs in a partition
scontrol show job <jobid>	List information on Job
scontrol show jobid -dd <jobid>	List detailed information on Job
sstat --format=AveCPU,MaxRSS,MaxVMSize,JobID -j <jobid>	List info resource used by your completed job : average cpu time, Max memory, Max virtual memory, JobId
sacct -u <username> --format=JobID,JobName,AveCPU,MaxRSS,MaxVMSize,JobID,Elapsed	List resources used by all jobs of a user
sprio	List job priority information

# Controlling jobs

Command	What its used for
scancel <jobid>	Cancel job
scancel -u <username>	Cancel all the jobs for a user
scancel -t PENDING -u <username>	Cancel all the pending jobs for a user:
Scancel -name JobName	Cancel one or more jobs by name
scontrol hold <jobid>	Hold a job, prevent it form starting
scontrol resume <jobid>	Release a job hold, allowing the job to try to start
scontrol requeue <jobid>	Requeue a running, suspended or finished job into pending state
scontrol requeuehold<jobid>	First requeue the job than put a hold on it.
squeue -u <username> -ho %A -t R	List running jobs by user
squeue --start	Show expected start time of jobs. (This can change)

# Getting information on you and your group

Command	What its used for
sacctmgr list Users USERS=<username>	List user and their default account (accounting group)
sacctmgr show user <username> withassoc	List user and their default account (accounting group) and shows more extensive information
sshare	Shows usage info for user.

# Getting information on your Cluster

Command	What its used for
sinfo --states=idle	Show idle node on cluster
sinfo -R	Show down, drained and draining nodes and their reason
sinfo --Node --long	Show detailed node info.
scontrol show reservation  user=root starttime=now duration=infinite flags=maint nodes=<nodeid>	Shows reservations on the cluster

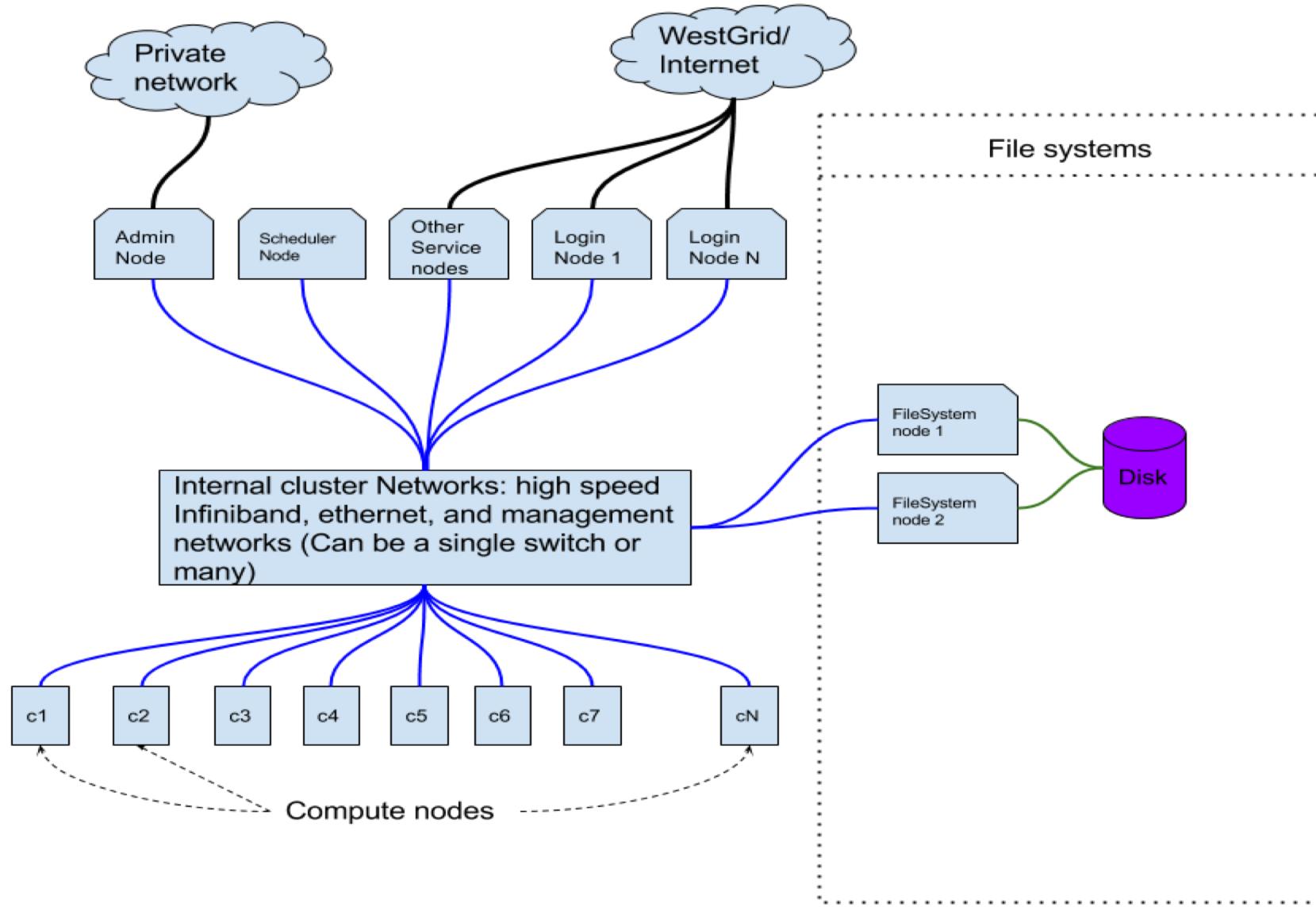
# Administrating your Cluster

Command	What its used for
scontrol create reservation user=root starttime=now duration=infinite flags=maint nodes=<nodeid>	Create a maintaince reservation on node nodeid
sacctmgr modify account where name=def-<account> set rawusage=0	Zero account usage fairshare stats
sacctmgr modify user where account=def-<account> name=<uname> set RawUsage=0	Zero user usage fairshare stats

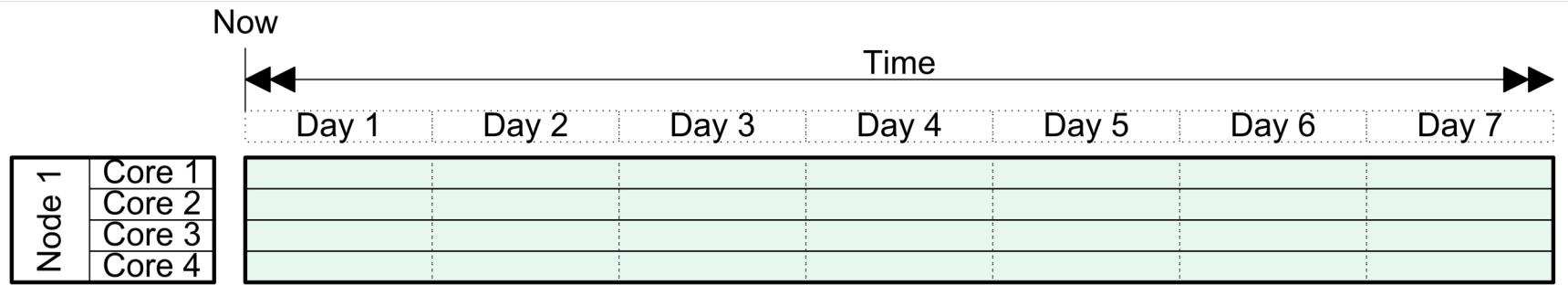
Material bellow will not be in the presentation unless diagram is needed to answer a question.

## **OTHER REFERENCE MATERIAL BELOW**

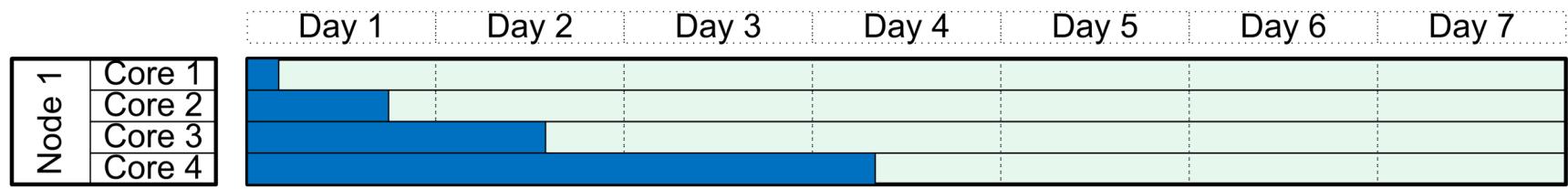
# Typical HPC Cluster



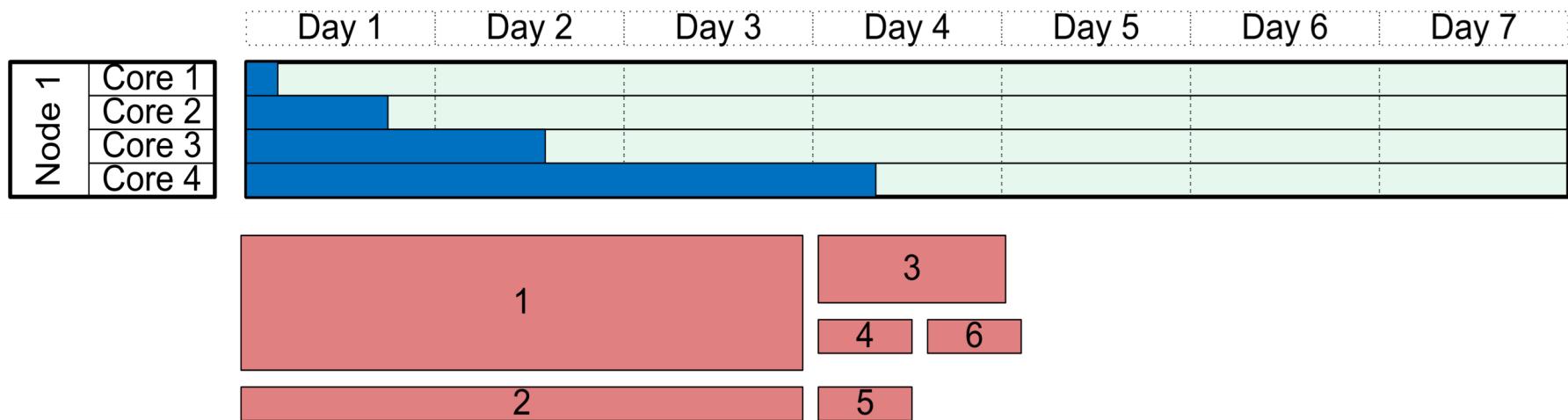
# Visualizing single node cluster



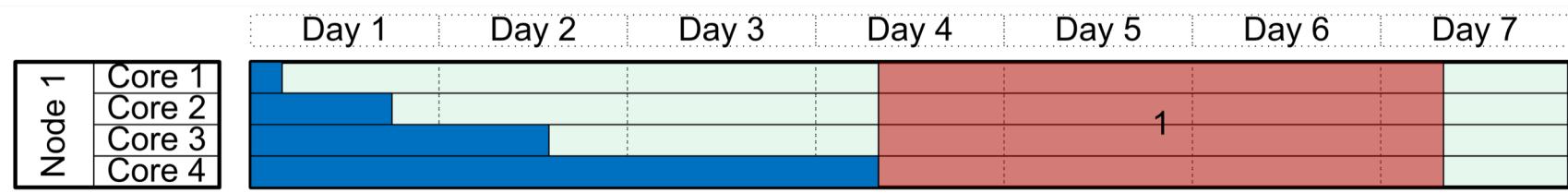
# Running jobs



# Scheduling jobs in order of priority



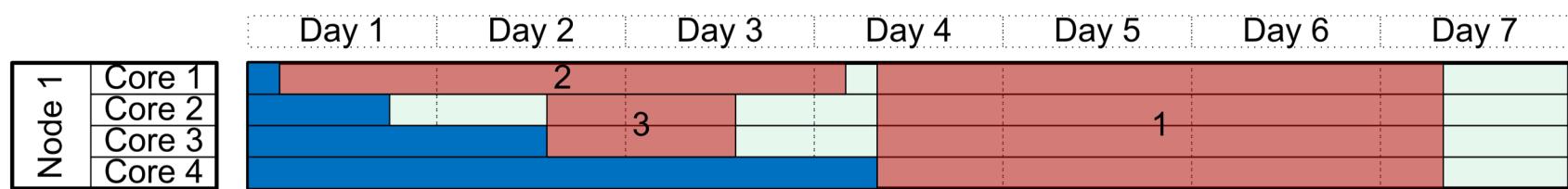
# Scheduling jobs in order of priority



# Scheduling jobs in order of priority

	Day 1	Day 2	Day 3	Day 4	Day 5	Day 6	Day 7
Node 1	Core 1	2					
	Core 2						
	Core 3						
	Core 4						

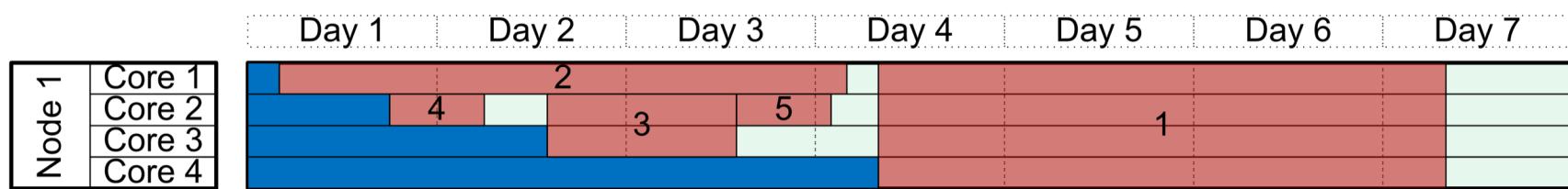
# Scheduling jobs in order of priority



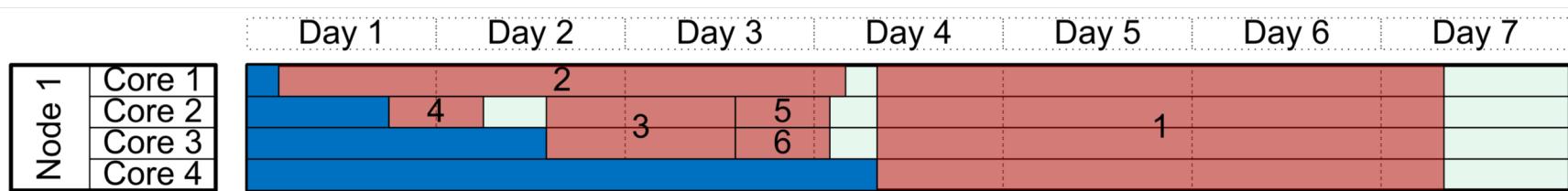
# Scheduling jobs in order of priority

	Day 1	Day 2	Day 3	Day 4	Day 5	Day 6	Day 7
Node 1	Core 1	2					
	Core 2	4	3		1		
	Core 3						
	Core 4						

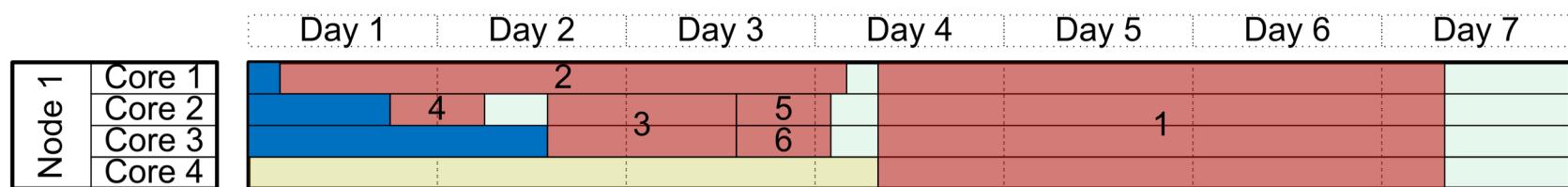
# Scheduling jobs in order of priority



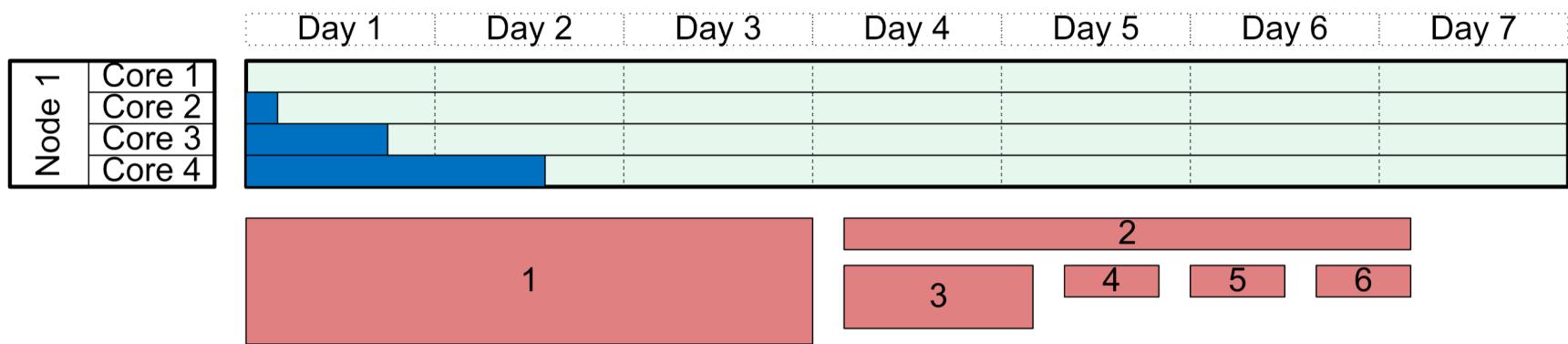
# Scheduling jobs in order of priority



# A Job finishes early



# Jobs are rescheduled



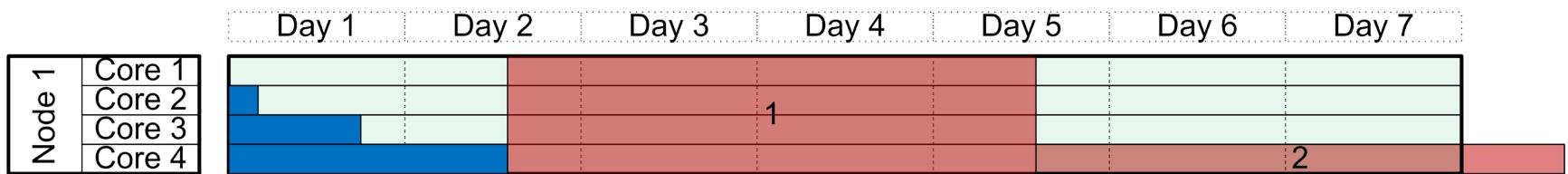
# Jobs are rescheduled

	Day 1	Day 2	Day 3	Day 4	Day 5	Day 6	Day 7
Node 1	Core 1						
	Core 2						
	Core 3						
	Core 4						

A Gantt chart illustrating job scheduling across 7 days (Day 1 to Day 7) for 4 cores (Core 1 to Core 4) on Node 1. The chart shows the start and end times of tasks assigned to each core.

- Core 1:** Task starts on Day 1 and ends on Day 1.
- Core 2:** Task starts on Day 1 and ends on Day 1. A second, longer task starts on Day 3 and ends on Day 6.
- Core 3:** Task starts on Day 1 and ends on Day 4.
- Core 4:** Task starts on Day 1 and ends on Day 5.

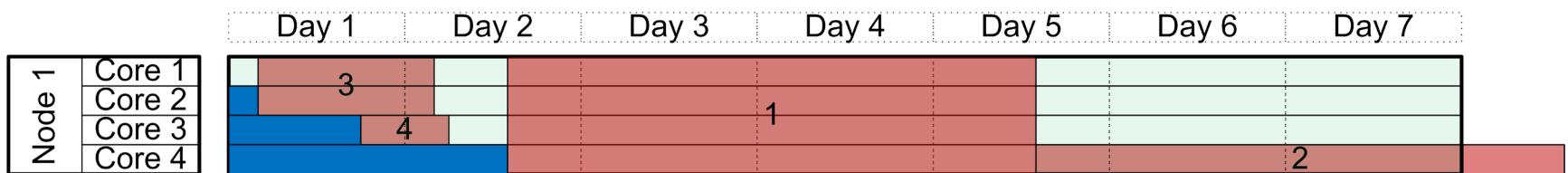
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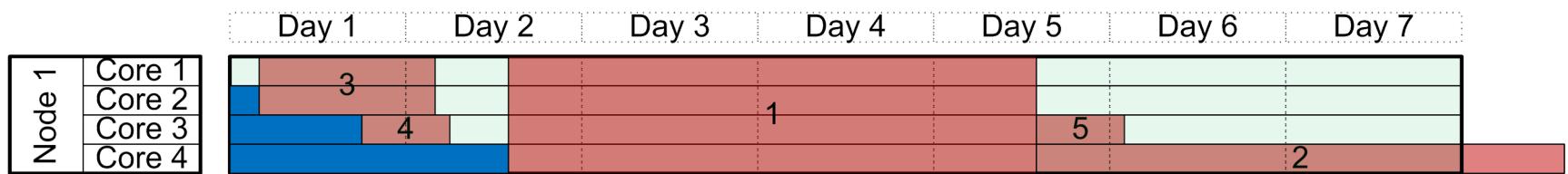
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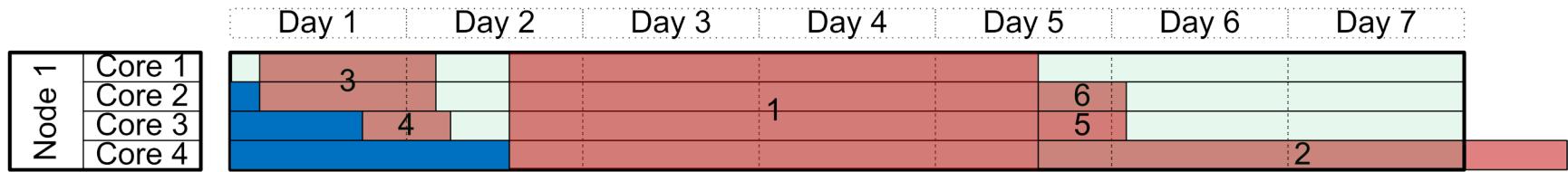
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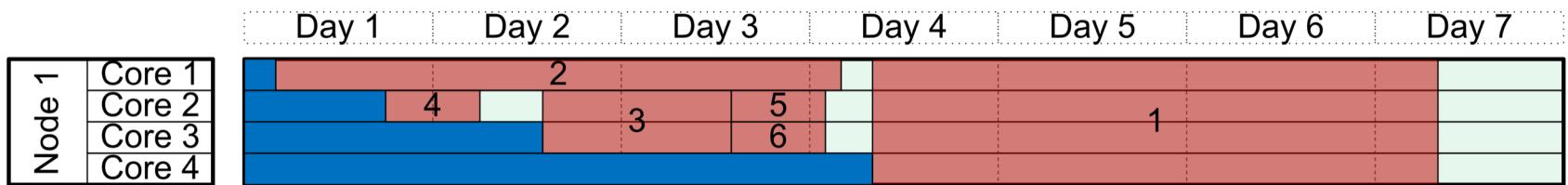
# Jobs are rescheduled



# Jobs are rescheduled



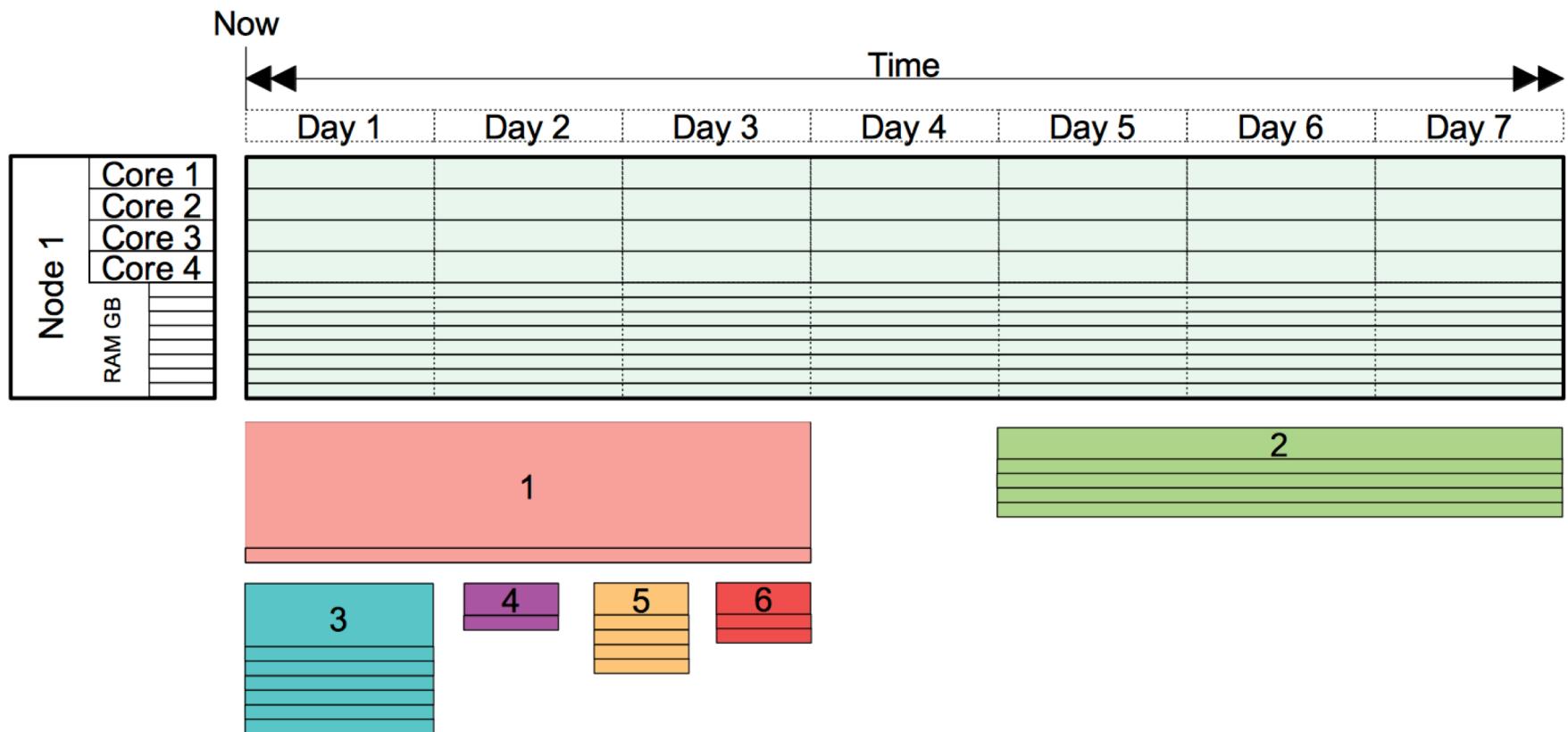
# Single node cluster



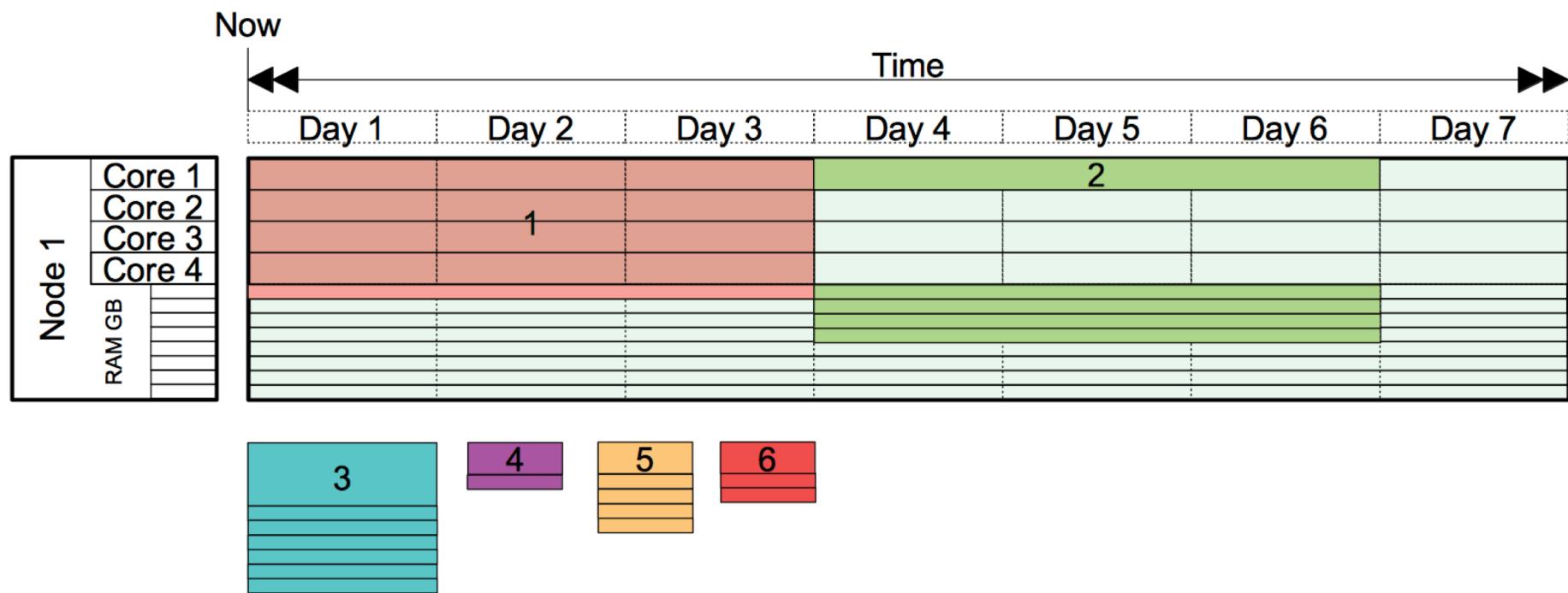
# Short serial jobs and Backfill

	Day 1	Day 2	Day 3	Day 4	Day 5	Day 6	Day 7
Node 1	Core 1 Core 2 Core 3 Core 4	2 4 3 6	1	5	1		

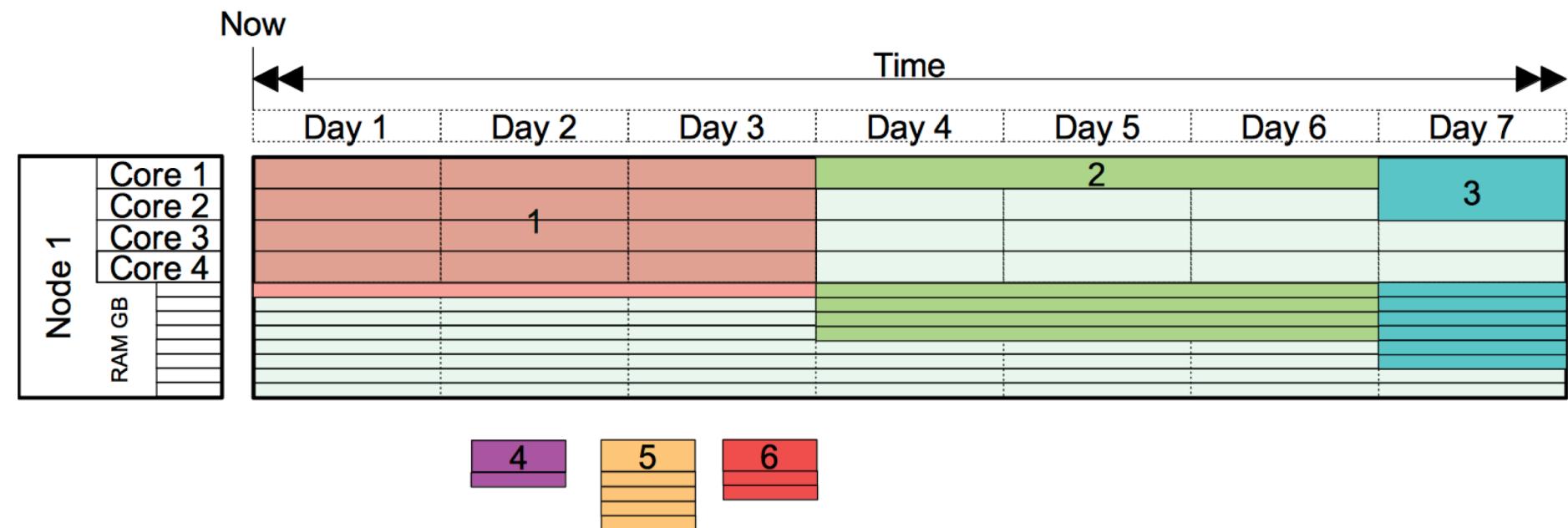
# Scheduling Cores and Memory



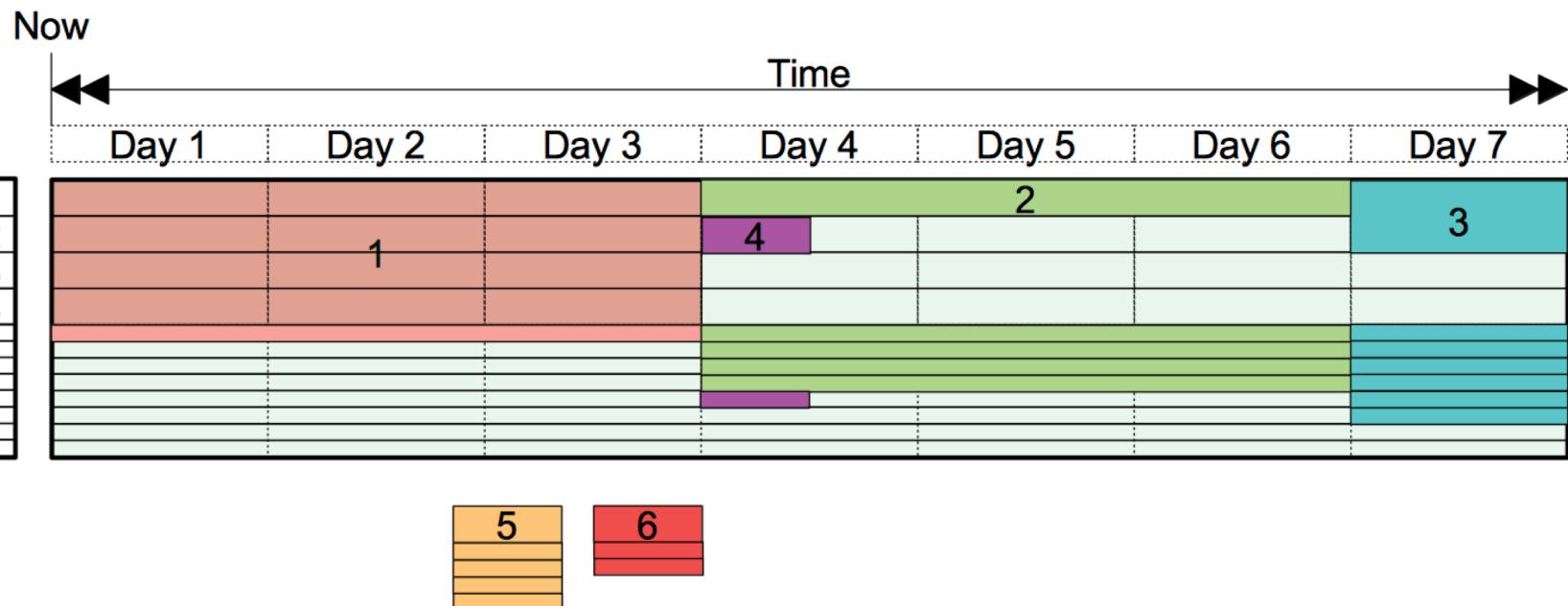
# Scheduling Cores and Memory



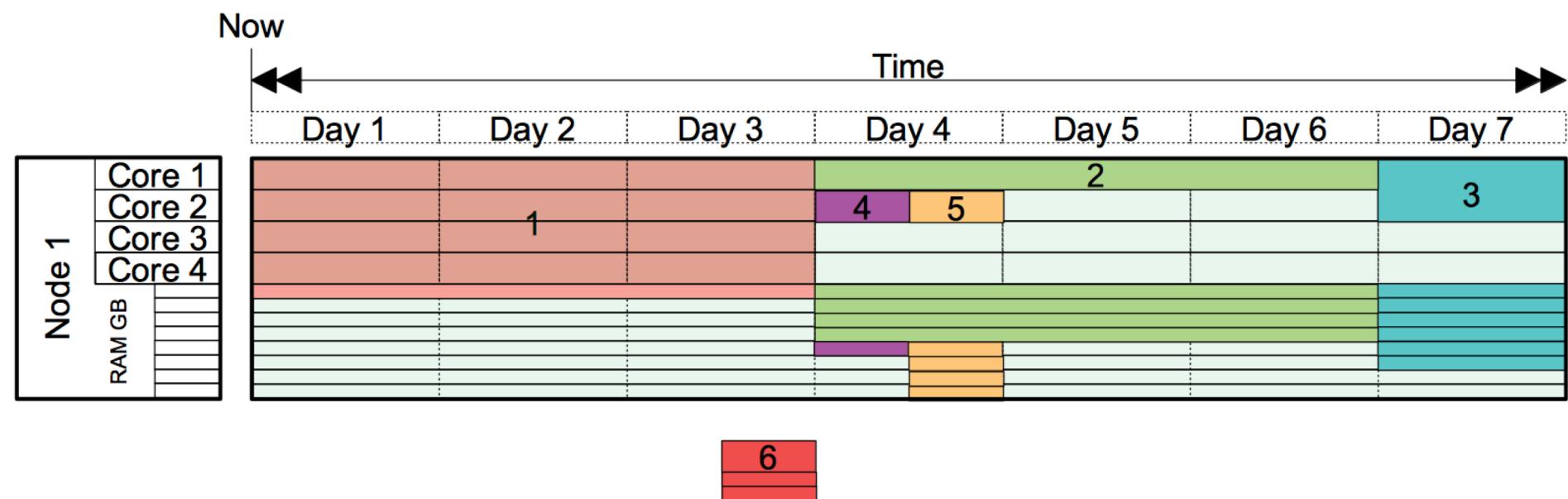
# Scheduling Cores and Memory



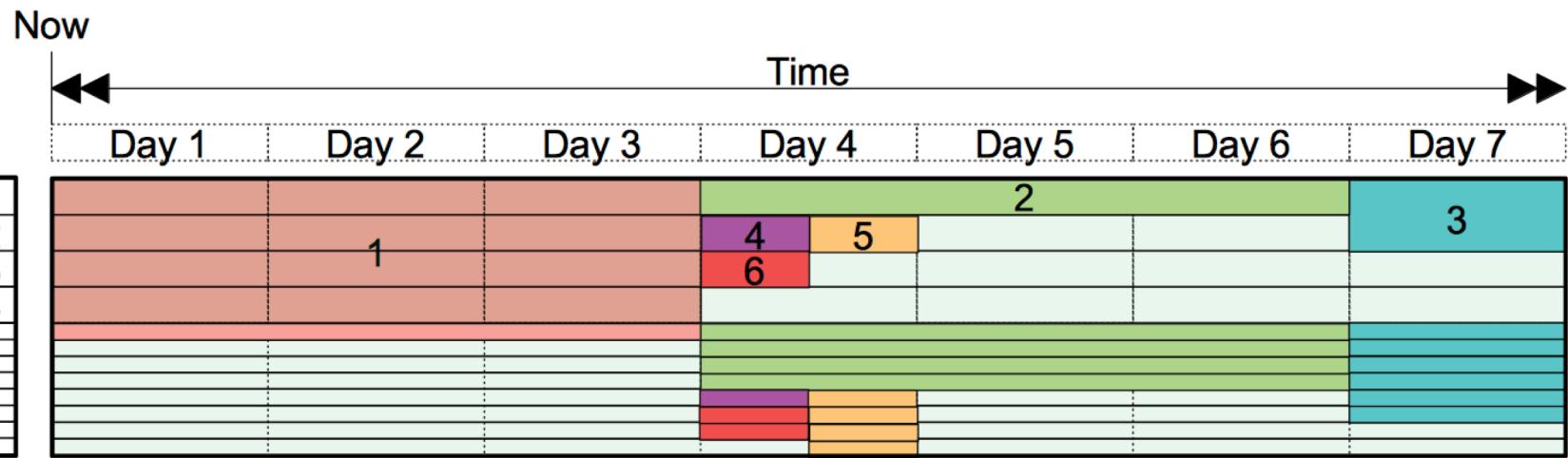
# Scheduling Cores and Memory



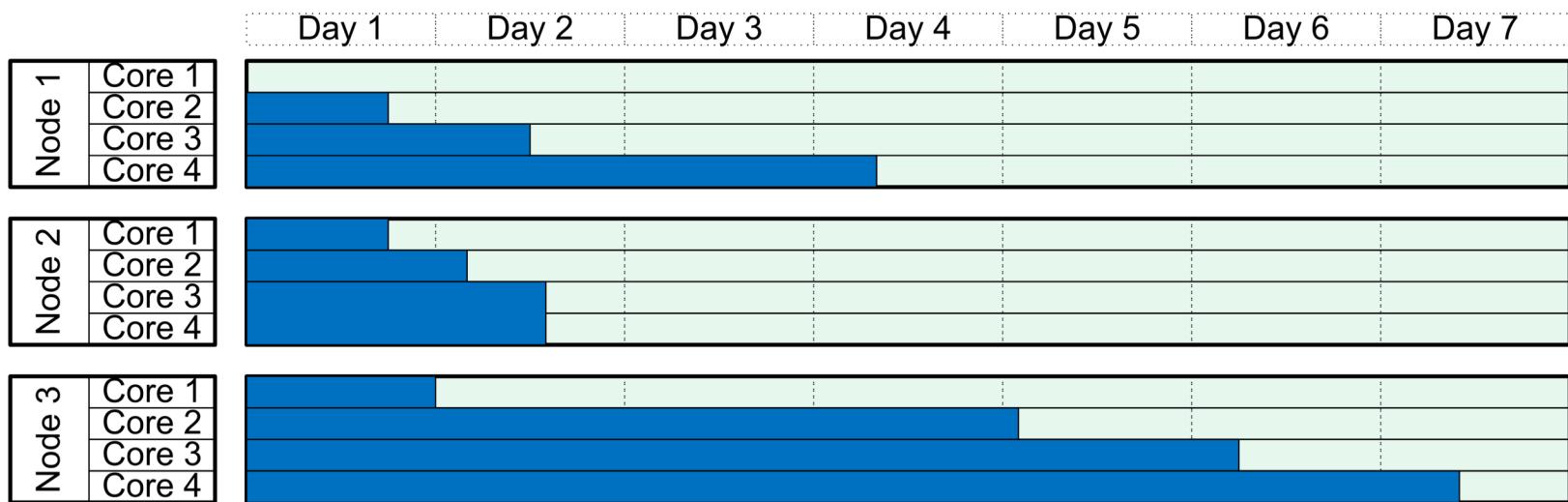
# Scheduling Cores and Memory



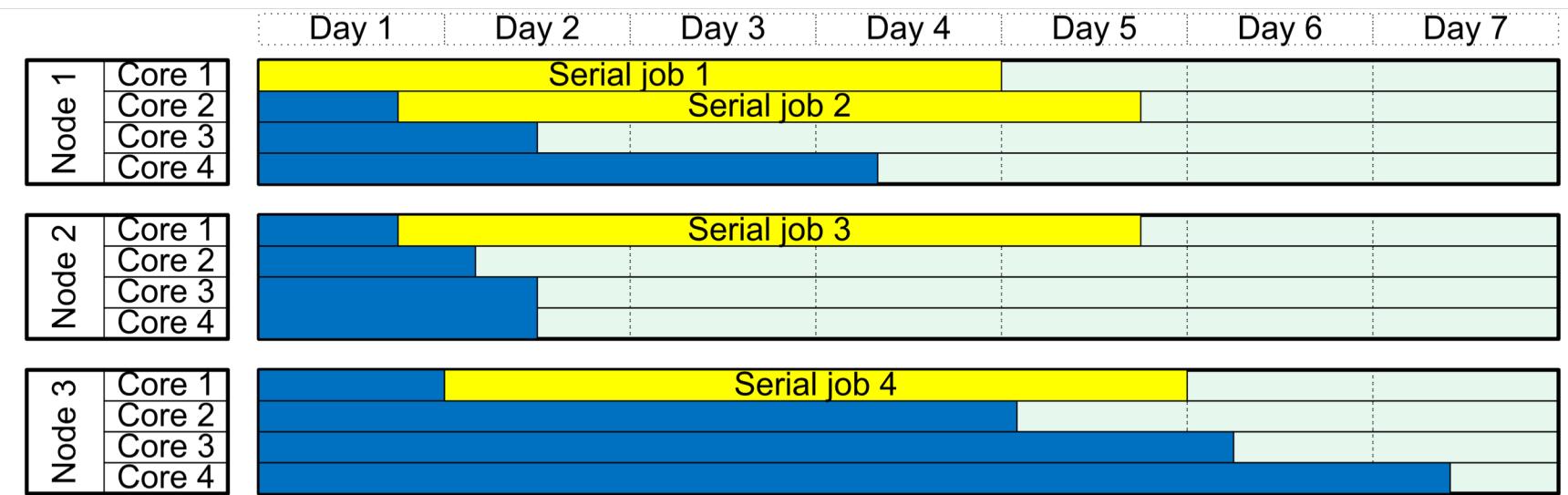
# Scheduling Cores and Memory



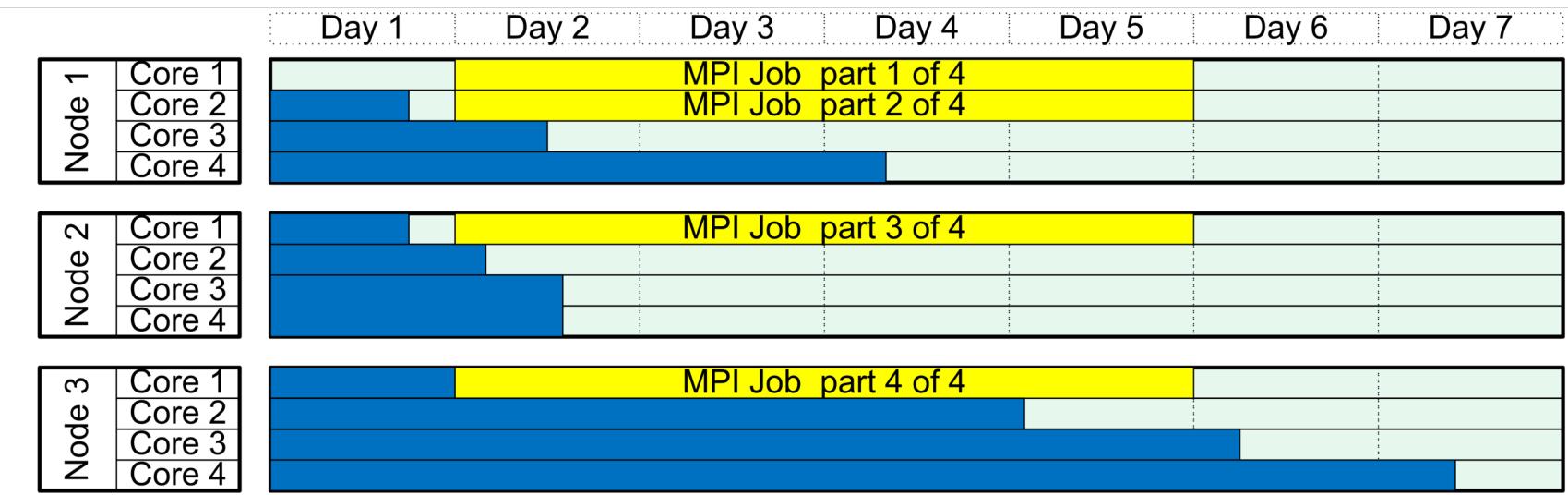
# Visualizing Multinode cluster



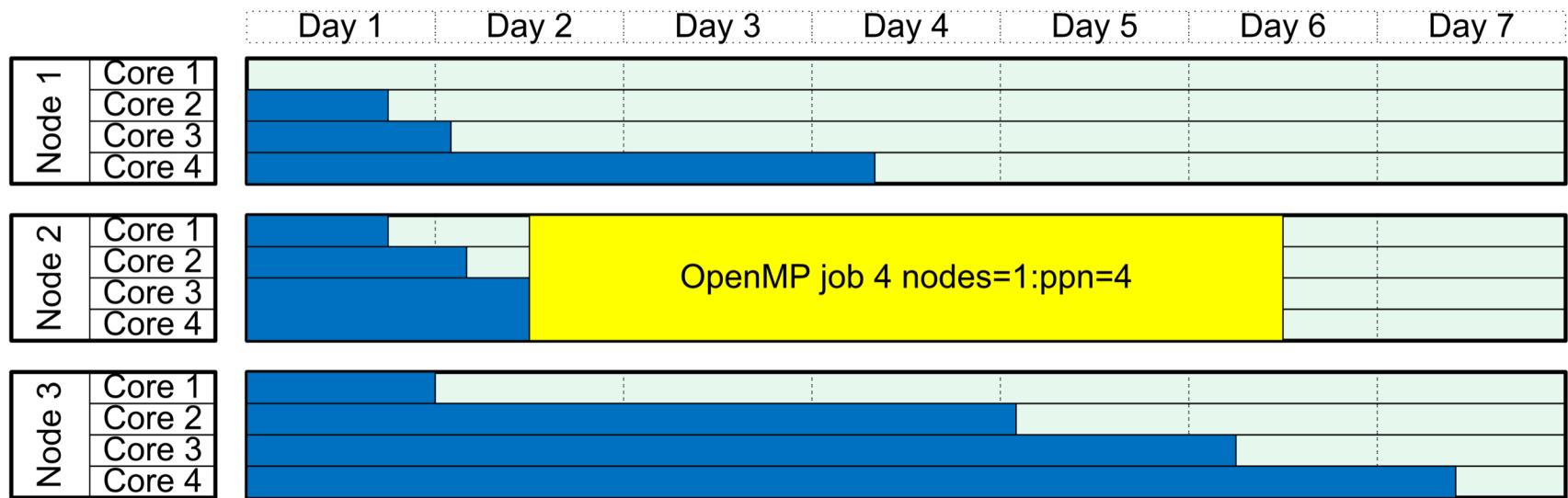
# Many Serial Jobs



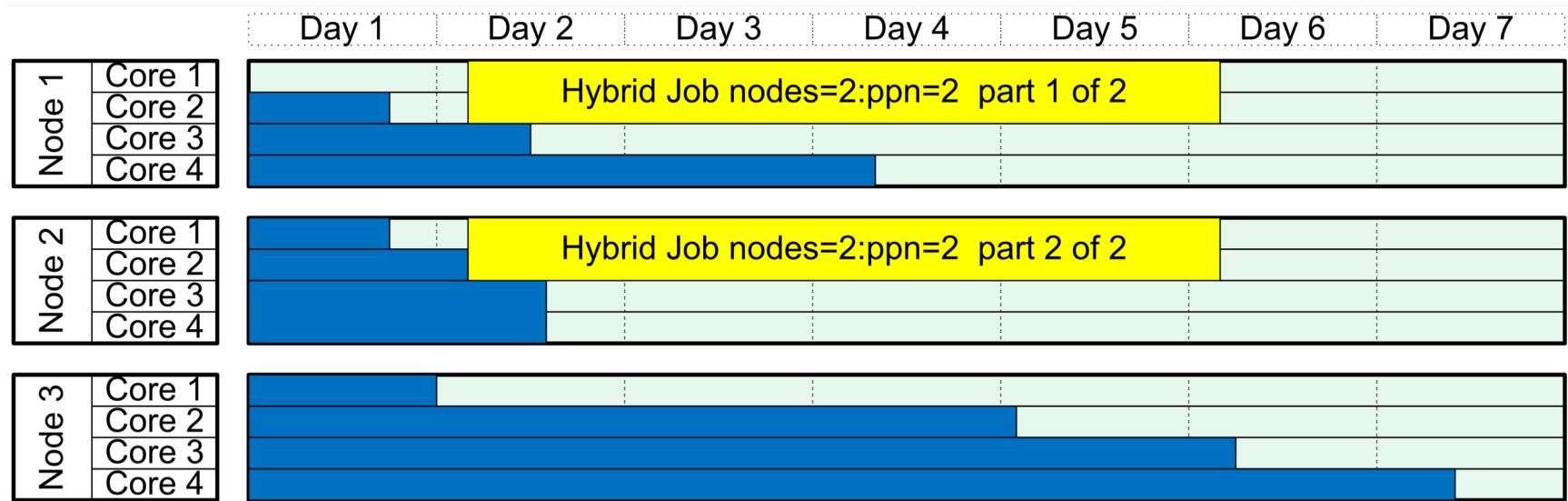
# MPI job



# Single node multi-core job (OpenMP, Gaussian, Threads)



# Hybrid Job

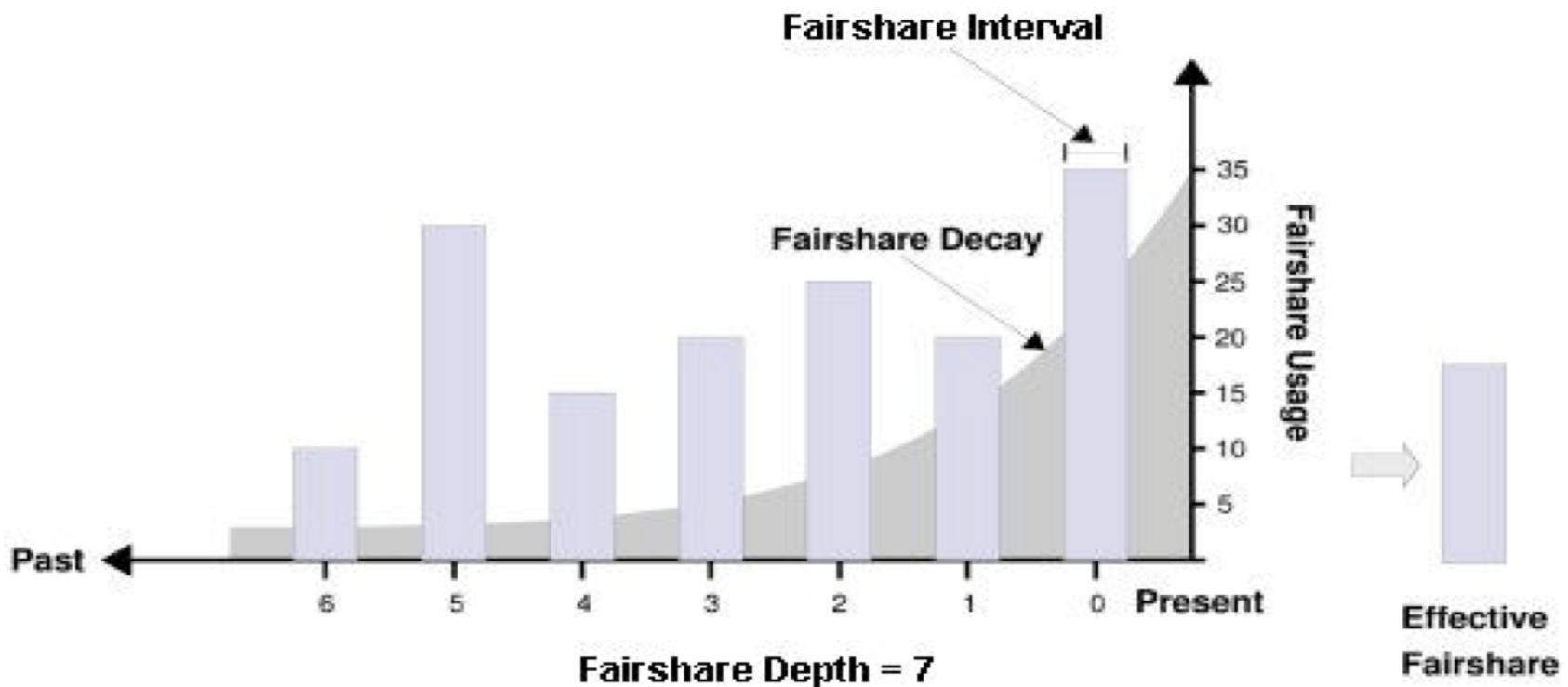


# Maximum job walltime partition limit

- A high maximum walltime is not necessarily a good thing, clusters that allow high walltime jobs take longer for jobs to start to run, and are less “fair”.
- There are advantages to running shorter jobs, such as how quickly your job can be started.
- The longer and larger a job is the greater the chances of experiencing hardware failure, minimize this through check pointing.
- Part of the resources of a cluster is dedicated for shorter jobs.
- Part of CC clusters are dedicated to whole node parallel jobs, other jobs with a short walltime of under 12 hours can run in this part at a reduced priority compared to whole node parallel jobs.

Partition name	Maximum walltime
*_b1	3 hours
*_b2	12 hours
*_b3	1 day
*_b4	3 days
*_b4	7 days
*_b6	28 days

# Fairshare



- Fair share usage is weighted by when the usage occurred recent usage is more important than usage at the end of the period

# Group's Status: “sshare -l”

Account	User	RawShares	NormShares	RawUsage	NormUsage	EffectvUsage	FairShare	LevelFS
<hr/>								
root			0.000000	639083114320110		1.000000		
no_rac_cpu		1320	0.043194	404703982221822	0.633257	0.633257		0.068209
ras_basic_cpu		1320	0.999243	404703982221822	0.633257	1.000000		0.999243
cc-debug_cpu		1	0.000236	1273287234	0.000002	0.000003		75.104409
cc-debug_cpu	kamil	1	0.004386	0	0.000000	0.000000	0.026537	inf
def-kamil_cpu		1	0.000236	0	0.000000	0.000000		inf
def-kamil_cpu	kamil	1	1.000000	0	0.000000	0.000000	0.486678	inf
no_rac_gpu		65	0.002127	6883285083841	0.010771	0.010771		0.197479
ras_basic_gpu		65	0.984848	6883285083841	0.010771	1.000000		0.984848
cc-debug_gpu		1	0.000236	12668	0.000000	0.000000		128389.386733
cc-debug_gpu	kamil	1	0.004386	0	0.000000	0.000000	0.508693	inf
def-kamil_gpu		1	0.000236	0	0.000000	0.000000		inf
def-kamil_gpu	kamil	1	1.000000	0	0.000000	0.000000	0.973463	inf

# Jobs by partition

## squeue -p <partitionname>

JOBID	PARTITION	NAME	USER	ST	TIME	NODES	NODELIST(R)
535639	cpubase_b	AE17631.	kamil	PD	0:00	1	(Resource)
591830	cpubase_b	bz.sh	erming	PD	0:00	1	(Resource)
615762	cpubase_b	AE21380.	kamil	PD	0:00	1	(Resource)
401219	cpubase_b	CTD095.s	john	PD	0:00	1	(Resource)
491576	cpubase_b	gen3x1s8	judy	R	2-08:04:59	1	cdr747
535638	cpubase_b	AE17594.	kamil	R	1-11:46:03	1	cdr101
491574	cpubase_b	gen3x1s6	masao	R	4-20:06:44	1	cdr79
491575	cpubase_b	gen3x1s7	masao	R	4-20:06:44	1	cdr85

# Priority sprio

JOBID	PRIORITY	AGE	FAIRSHARE	PARTITION	TRES
130976	<b>7088</b>	2500	0	625	cpu=2526,mem=1437
167003	<b>6150</b>	2500	0	1250	cpu=2008,mem=392
195802	<b>4996086</b>	2500	4991771	833	cpu=469,mem=45,gres/
195809	<b>4996086</b>	2500	4991771	833	cpu=469,mem=45,gres/
195810	<b>4996086</b>	2500	4991771	833	cpu=469,mem=45,gres/
205281	<b>8206</b>	2500	0	625	cpu=1875,mem=1800,gr
205290	<b>6408</b>	2500	0	625	cpu=1875,mem=2,gres/
544814	<b>23534</b>	1741	21571	208	cpu=13,mem=2
544815	<b>23534</b>	1741	21571	208	cpu=13,mem=2
617580	<b>24194</b>	373	22768	1042	cpu=10,mem=2
617581	<b>24194</b>	373	22768	1042	cpu=10,mem=2