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Workload Management & Slurm

Presented by CSCNSI

What is Workload Management?

- We need a way to easily put our many compute nodes to use
- We need a tool that can match work that needs to be done with resources. that are available
- It needs to be smart enough to
 - Start multi-node jobs
 - Schedule jobs from a long list of waiting tasks according to policy
 - Know what resources are available
 - Know what jobs to send to what hardware
 - Know how to start a job on our hardware
- And generally a lot of other things, like keeping track of accounting information, etc.

Scheduling vs. Resource Management

Scheduler

- The "Brain" of the system
- Single, centralized daemon
- Obtains job and node data from Resource Manager
- Prioritizes jobs
- Decides:
 - What jobs will run
 - Where they will run
 - When they will run

Resource Manager



- The "Worker" of the system
- Server daemon plus per-node client daemons
- Gathers node information (CPUs, memory, disk, load)
- Answers queries for this information
- Spawns jobs when told to
- Can add/remove resources

Scheduling Cycle

Job Ordering

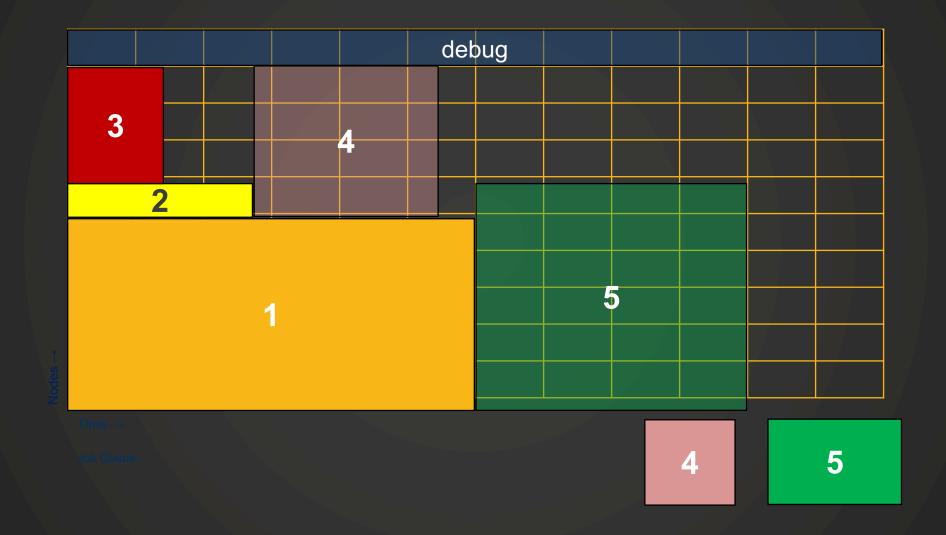
- Calculate job priorities and sort job list by priority
- Start all jobs which can start
- Make reservations for next 1024 jobs
- Use backfill policy to sort remaining jobs (1024)

Job Placement (for each job being started)

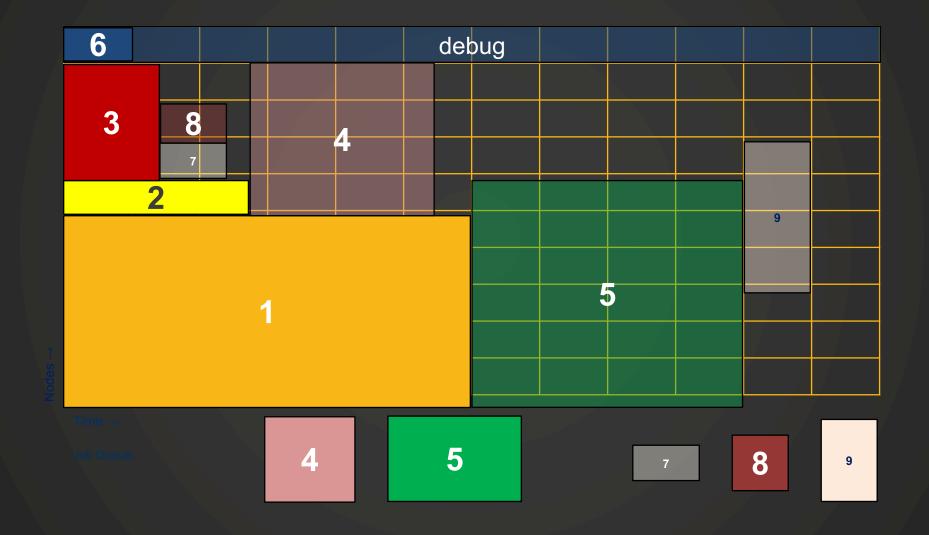
- Filter list of all nodes based on requirements
- Filter list of nodes based on eligibility
- Sort final node list based on node allocation policy
- Start job on top n nodes



Scheduling Example – Iteration #1



Scheduling Example – Iteration #2 (backfill)



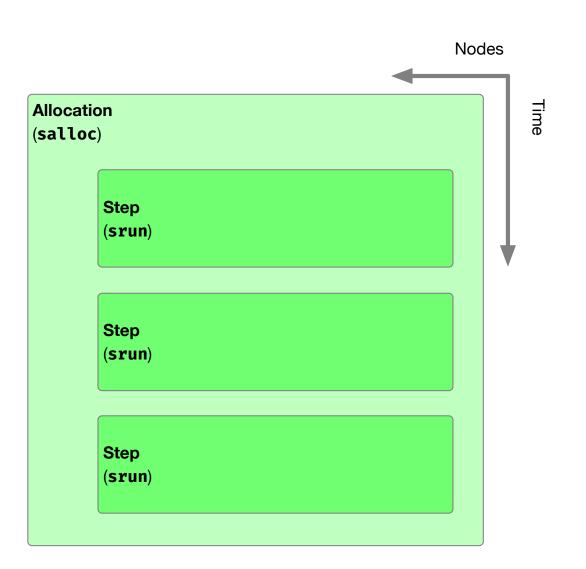
What is Slurm?

- Slurm is a Workload Manager for HPC workloads
- Originally "Simple Linux Utility for Resource Management"
 - Was spelled SLURM
- Since it now is also a scheduler, was rebranded *Slurm* (not an acronym)
- Originally written at LLNL
- Now commercially developed & supported by SchedMD
- Used on more than half of the top 10 of the top500



Structure of a Slurm job

- A job needs an "allocation"
 - Specifies needed resources
 - Memory
 - Nodes/CPUs
 - Specifies a max time
- Within an allocation multiple steps can execute
 - Steps, in total, must fit within allocation



Slurm interactive job

- An allocation is made with the salloc command
- An allocation is bound to a process
- **srun** started within that process create a step
- The job ends when the allocation process ends

```
[fe] $ salloc -n1 /bin/bash
salloc: Granted job allocation 423
[fe] $ srun --pty /bin/bash
[n01] $ ./do_work
[n01] $ exit
[fe] $ srun -pty ./do_more_work
(runs on n01)
[fe] $ exit
salloc: Relinquishing job allocation
423
[fe] $
```

Slurm batch job

- Jobs can be specified as a predefined script
- Special #SBATCH lines provide Slurm parameters
- Can submit multiples of the same task
 - Or can submit arrays of the same job, keyed by the
 \$SLURM_ARRAY_TASK_ID
- Submitted with:
 - sbatch <script>

```
#!/bin/bash
#
#SBATCH --job-name=test_emb_arr
#SBATCH --output=res_emb_arr.txt
#SBATCH --ntasks=1
#SBATCH --time=10:00
#SBATCH --mem-per-cpu=100 #
#SBATCH --array=1-8
srun ./my_program.exe
   $SLURM_ARRAY_TASK_ID
```

Common Slurm commands

Start an allocation	salloc
Run a step in an allocation	srun
Submit a batch script	sbatch
View the queue	squeue
Stop a running job	scancel
Attach to I/O of current task	sattach
View the cluster status	sinfo
View job accounting data	sacct
Control Slurm	scontrol

All of the commands have extensive options. See `man <slurm_command>`

Los Alamos National Laboratory Questions?