Introducing Flux

A scalable resource manager for LC clusters

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What is flux?

- Flux is a modular, fully hierarchical resource manager and job scheduler.
- Modular development model allows a rich and consistent API which makes it easy to launch flux instances from within scripts.
- Fully hierarchical means that every flux 'job step' can be a full flux instance with the ability to schedule more job steps on its resources.
- Flux can be used now on LC systems.

Flux uses a new model for scheduling



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Usability: Submitting a Batch Job

- Slurm
 - sbatch -N2 -n4 -t 2:00 sleep 120
- Flux CLI
 - flux mini submit -N2 -n4 -t 2m sleep 120

Flux API

https://github.com/flux-framework/Tutorials/tree/master/2020-ECP



Scalability: Running Many Jobs

Slurm / CLI

- find ./ -exec sbatch -N1 tar -cf {}.tgz {}\;
 - Slow: requires acquiring a lock in Slurm, can timeout causing failures
 - Inefficient: uses 1 node for each task
- find ./ -exec srun -n1 tar -cf {}.tgz {}\;
 - Slow: spawns a process for every submission
 - Inefficient: is not a true scheduler

Flux API

```
flux start my_jobs.py
------
import flux, flux.job
from flux.job import JobspecV1

h = flux.Flux()
for f in os.listdir('.'):
    command = ["tar", "-cf", "{}.tgz".format(f), f]
    flux.job.submit(h, JobspecV1.from_command(command))
```

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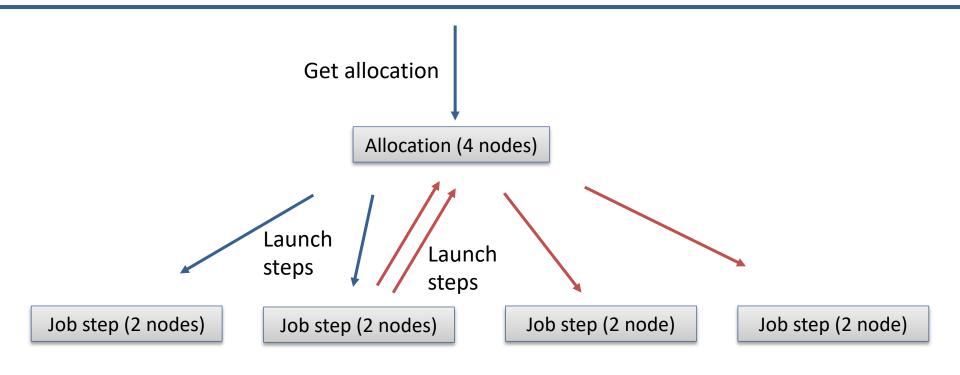
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Flux is fully hierarchical

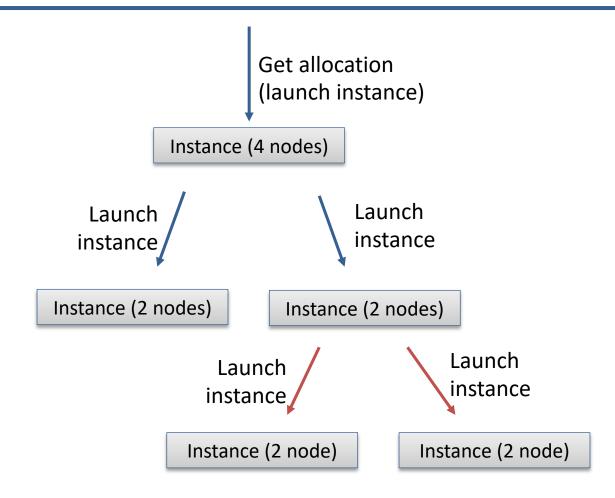


Flux is hierarchical: Launching steps in Slurm



Complex schedulers allow complex workflows

Flux is hierarchical: Launching instances in Flux



Complex schedulers allow complex workflows

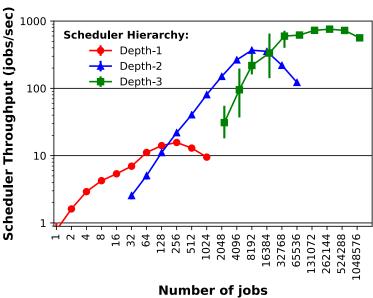


Scalability: Running Millions of Jobs

- Single Flux Instance— flux start my workflow.py
- Statically Partitioned Flux Instances

Flux Hierarchy

```
- flux-tree -N $num_nodes \
    -T ${num_nodes} \
    -J $num_jobs -- jobspec.yaml
- flux-tree -N $num_nodes \
    -T ${num_nodes}x${cores_per_noces}
    -J $num_jobs -- jobspec.yaml
```

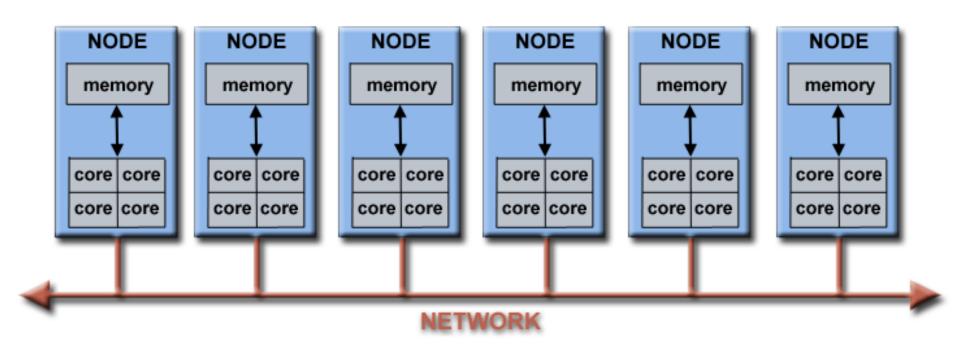


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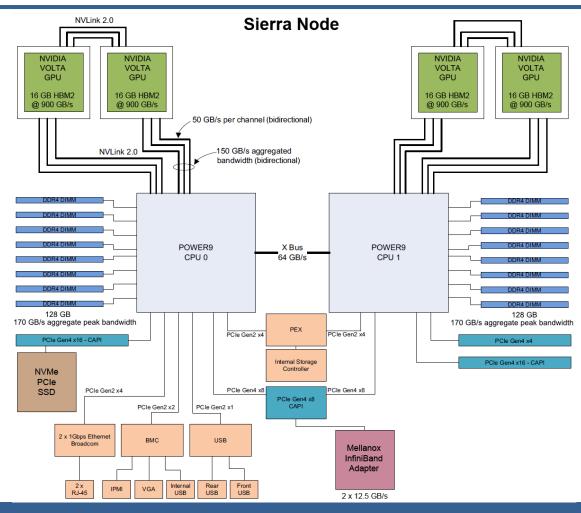
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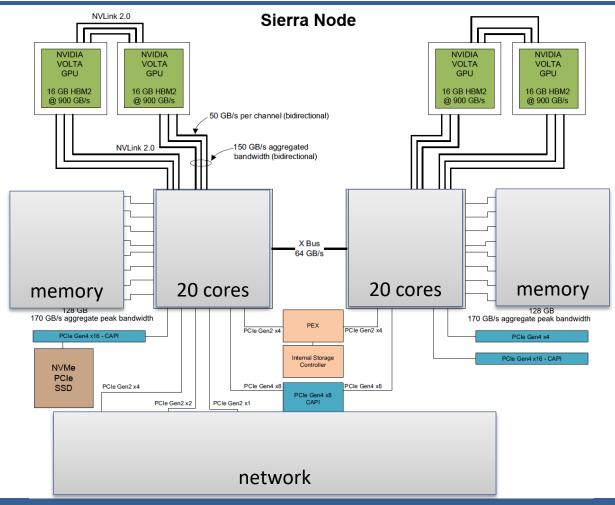


Traditional scheduling maps well to simple nodes



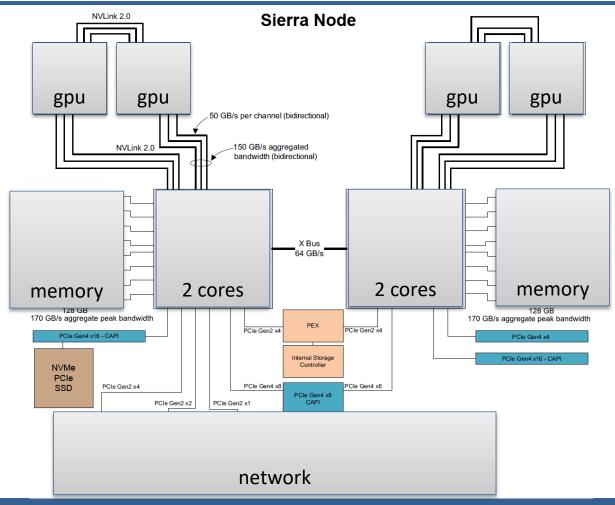
New systems are more complex and heterogeneous





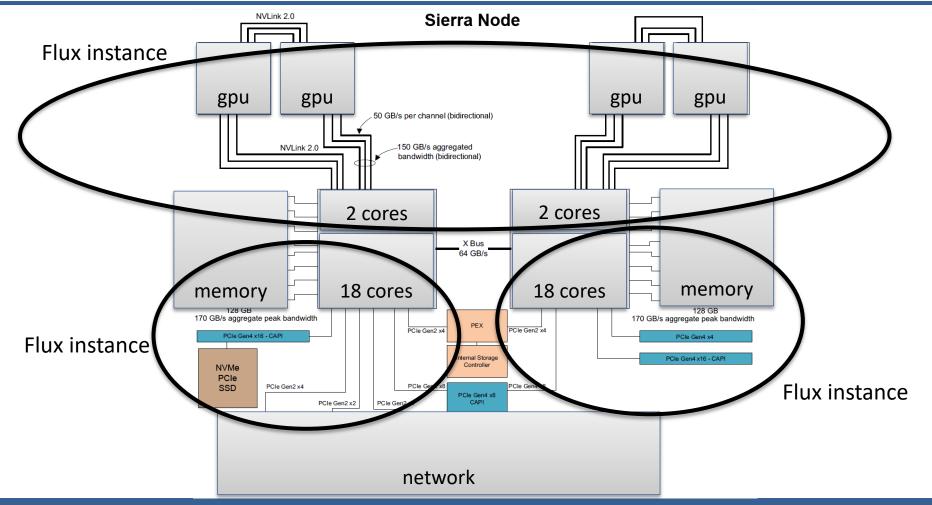
~10% of cycles on node





~90% of cycles on node

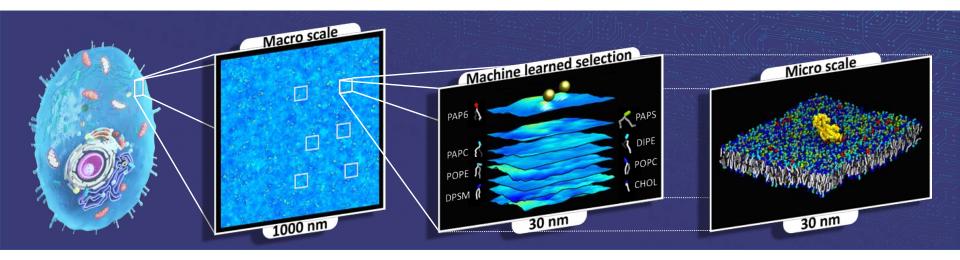




~100% of cycles on node



MuMMI implements a complex workflow to enable a new genre of multiscale simulation for cancer research



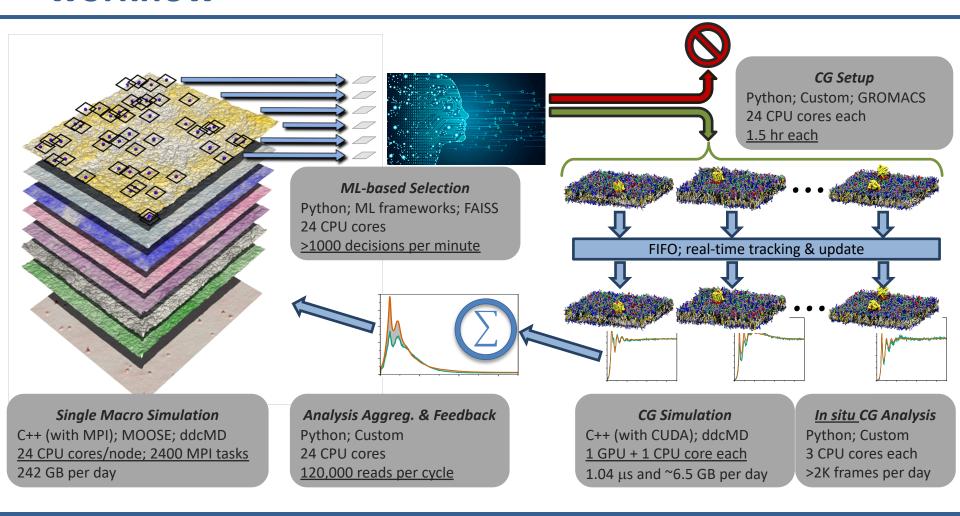
Multiscale Machine-Learned Modeling Infrastructure (MuMMI)

 Novel framework coupling multiple scales using a hypothesis driven selection process.

https://github.com/flux-framework/Tutorials/tree/master/2020-ECP (Di Natale)



MuMMI implements a complex and dynamic workflow

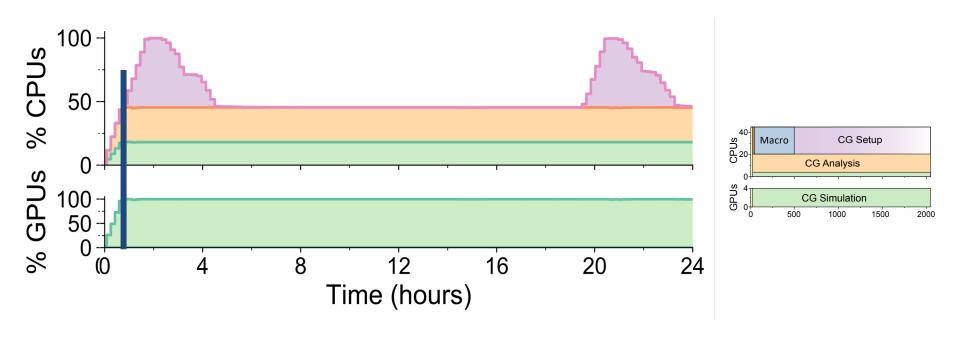


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The high through-put, low latency scheduling enables fast restarts and consistent utilization of all resources

- Hierarchical scheduling allows MuMMI to reach steady state in ~45 minutes (newer versions will reduce turnaround time)
- Depending on the scientific hypothesis MuMMI utilizes >95% of the available compute



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Using Flux: getting flux

Installed on LC systems

[day36@rzslic4:~]\$ ls /usr/global/tools/flux/\$SYS_TYPE/default/bin flux

- Install with spack: spack install flux-sched
- Build from source

git clone https://github.com/flux-framework/flux-core.git configure, make, make install

git clone https://github.com/flux-framework/flux-sched.git configure, make, make install

https://flux-framework.readthedocs.io/en/latest/quickstart.html#building-the-code

Using Flux: starting an instance

```
[day36@rzalastor2:~]$ salloc -N4 --exclusive salloc: Granted job allocation 220682 sh-4.2$ flux keygen
```

Saving /g/g0/day36/.flux/curve/client

Saving /g/g0/day36/.flux/curve/client_private

Saving /g/g0/day36/.flux/curve/server

Saving /g/g0/day36/.flux/curve/server_private

sh-4.2\$ srun -N4 -n4 --pty flux start

sh-4.2\$ flux mini run -n4 hostname

rzalastor16

rzalastor15

rzalastor17

rzalastor14

sh-4.2\$

https://flux-framework.readthedocs.io/en/latest/quickstart.html#starting-a-flux-instance

Using Flux: running a batch script

```
sh-4.2$ cat quickexample.sh
#!/bin/sh
flux mini batch -N 2 -n 2 --wrap << EOF
date
flux mini run -n 2 ~/hello/hello mpi
FOF
sh-4.2$ ./quickexample.sh
f4aDXvqSo
sh-4.2$ flux jobs -f completed, failed
   IOBID USER NAME
                        ST NTASKS NNODES RUNTIME RANKS
 f4aDXvqSo day36 batchscrip CD 2 2 4.302s [0-1]
  f5jrorGw day36 hostname CD 4 4 0.098s [0-3]
sh-4.2$ cat flux-f4aDXvqSo.out
Tue Dec 1 12:05:21 PST 2020
Hello from task 0 on rzalastor14!
MASTER: Number of MPI tasks is: 2
Hello from task 1 on rzalastor15!
sh-4.2$
```

https://flux-framework.readthedocs.io/en/latest/quickstart.html#launching-work-in-a-flux-session



Where to find out more

CLI

- https://flux-framework.readthedocs.io/en/latest/batch.html
- Man flux-mini, man flux-jobs, etc.

API / Workflow

- https://flux-framework.readthedocs.io/projects/flux-workflowexamples/en/latest/index.html
- https://github.com/flux-framework/Tutorials
- https://github.com/LLNL/maestrowf
 - https://lc.llnl.gov/confluence/display/MAESTRO/Maestro+Home
- Email <u>lc-hotline@llnl.gov</u> with questions, bugs, or to get in touch with the workflows team.

Questions?



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