
Introduction to Using Gordon

*2012 International Summer School on AstroComputing
San Diego, CA July 9-20*

*Robert Sinkovits
Gordon Applications Lead
San Diego Supercomputer Center*

An apology in advance

Those of you who are already experienced users of NSF, DoE, DoD, PRACE, etc. supercomputing resources may find much of the material being presented here to be very basic.

Our intention is to bring everyone up to speed early in the summer school so that the on-hands exercises will go as smoothly as possible

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Click the “SIGN IN” link

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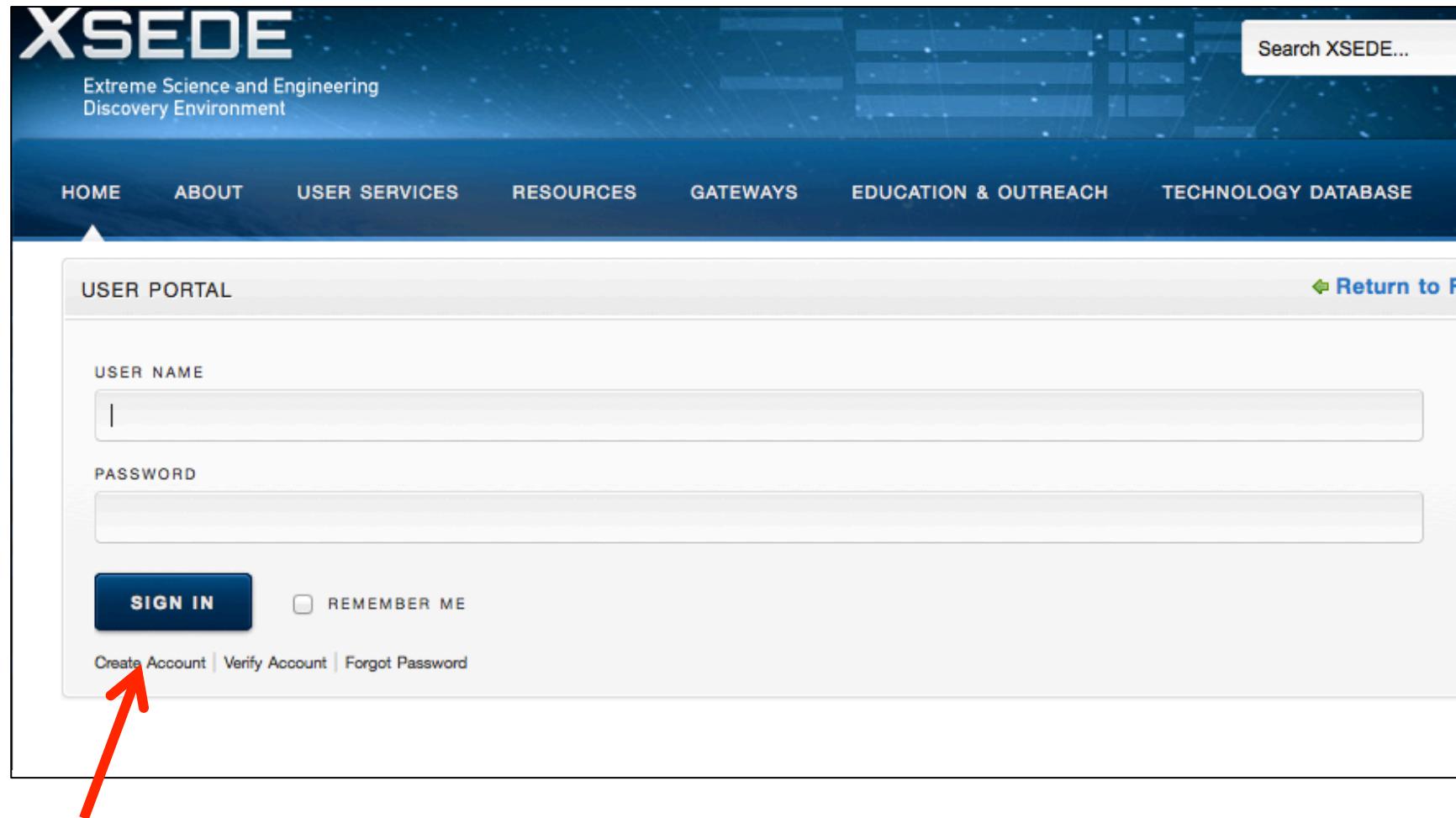
- See you in Chicago in July at the XSEDE 12 CONFERENCE
- Ready to work with XSEDE? READ THE GETTING STARTED GUIDE
- Log in to the XSEDE User Portal

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NICS simulations help supercomputing, software, support ... and social, too

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... click the “Create Account” link



... Fill out the information, create account and email your XSEDE portal name to rpwagner@sdsc.edu

The screenshot shows the XSEDE User Portal account creation page. At the top, there's a navigation bar with links for HOME, ABOUT, USER SERVICES, RESOURCES, GATEWAYS, EDUCATION & OUTREACH, and TECHNOLOGY DATABASE. On the right side of the header are a search bar labeled "Search XSEDE..." and a "SIGN IN" button. Below the header, a banner says "USER PORTAL" and includes a "Return to Full Page" link. The main content area is titled "Create an XSEDE User Portal account". It asks for personal information: FIRST NAME, MIDDLE NAME, LAST NAME, ORGANIZATION, DEPARTMENT, DEGREE, and DEGREE FIELD OF STUDY. The "FIRST NAME" field contains a single character, and the "DEGREE" dropdown menu has "Choose one" selected.

Why Gordon?



Designed for data and memory intensive applications that don't run well on traditional distributed memory machines

- Large shared memory requirements
- Serial or threaded (OpenMP, Pthreads)
- Limited scalability
- High performance data base applications
- Random I/O combined with very large data sets
- Large scratch files

Gordon is a national resource made possible through a grant from the National Science Foundation. One of three OCI Track 2D awards for innovative systems

Available to all U.S. academic researchers on a competitive basis and on a limited basis for-fee to most non-academic users



Design
Deployment
Support



Integrator



Funding
OCI #0910847



Processors
Motherboards
Flash drives



vSMP Foundation



3D Torus

Gordon Hardware overview

- **1024 dual-socket compute nodes**
 - 2 x Intel EM64T Xeon E5 (Sandy Bridge) processors
 - 64 GB DDR3-1333 memory
 - 80 GB local Flash memory
 - *64 TB total DRAM, 341 TFlop peak performance*
- **64 dual-socket I/O nodes**
 - 2 x Intel Westmere processors
 - 48 GB DDR3-1333 memory
 - 16 x 300 GB Intel 710 (Westmere) Solid State Drives (SSD)
 - *300 TB total flash memory*
- **Dual-rail 3D torus InfiniBand QDR (40 Gbit/s) network**
- **4 PB Lustre-based parallel file system**
 - *Capable of delivering up to 100 GB/s to Gordon*

A great Gordon application will ...

- Make use of the flash storage for scratch/staging
64 I/O nodes each w/ 16 x 300 GB SSDs (4.4 TB usable)
- Need the large, logical shared memory
~ 1 TB DRAM & 256 cores, w/ larger configurations possible
- Be a threaded app that scales to very large number of cores
- Require large physical memory per node
64 GB/node, 4 GB/core
- Be able to use the AVX instructions (8 flops/cycle/core)
- Need a high-bandwidth, low-latency inter-processor network

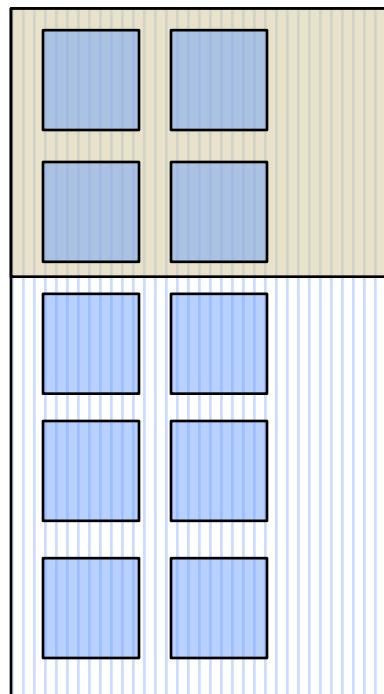
Foxglove Calculation using Gaussian 09 with vSMP - MP2 Energy Gradient Calculation



The Foxglove plant (Digitalis) is studied for its medicinal uses. *Digoxin*, an extract of the Foxglove, is used to treat a variety of conditions including diseases of the heart. There is some recent research that suggests it may also be a beneficial cancer treatment.



1 Compute node
= (16 cores/node)
64 GB/node)



Processor footprint - 4 nodes
64 threads

Time to solution:
43,000s

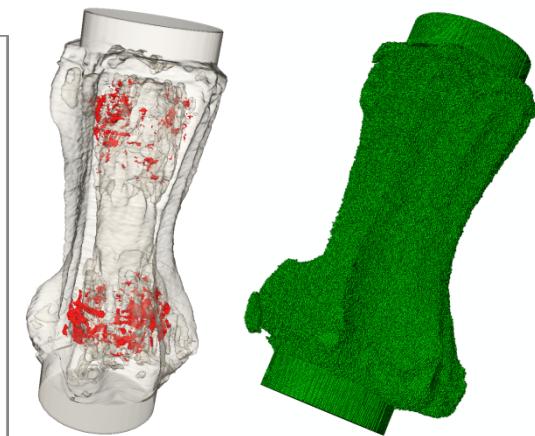
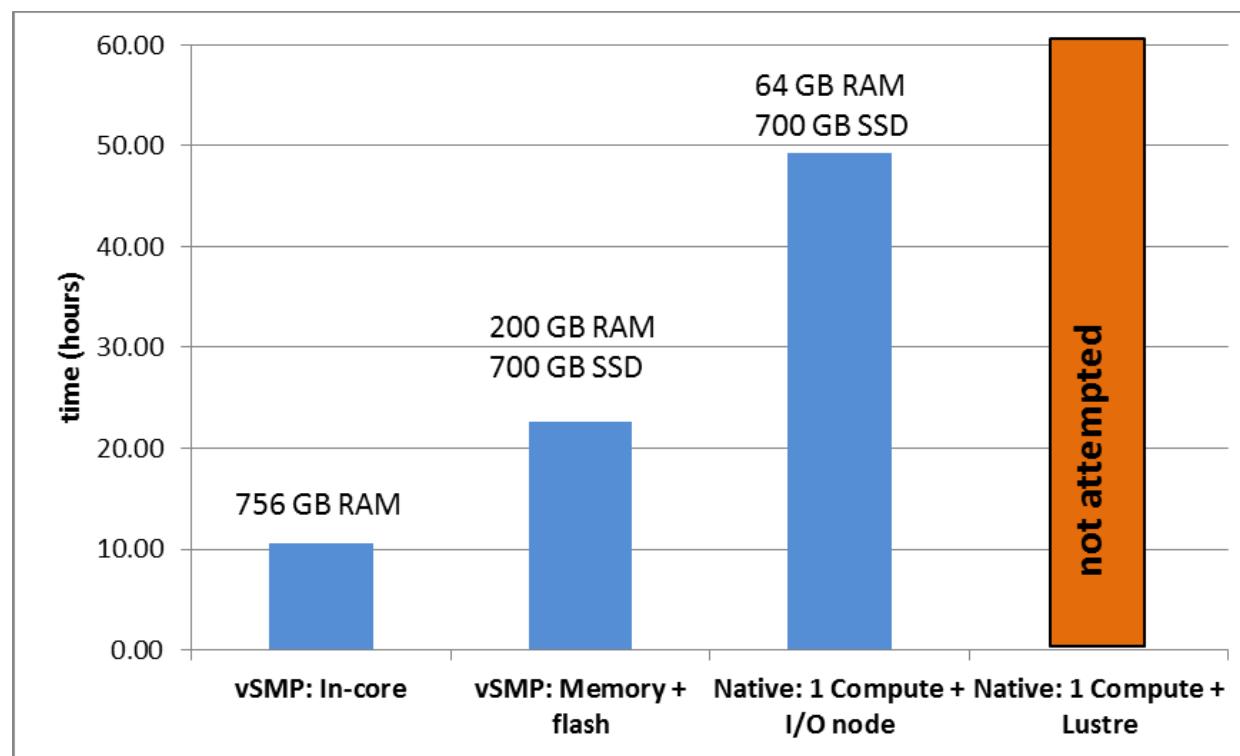
Memory footprint – 10 nodes
700 GB

V	M	F
C	T	L

Source: Jerry Greenberg, San Diego Supercomputer Center. January, 2012.

Axial compression of caudal rat vertebra using Abaqus and vSMP

The goal of the simulations is to analyze how small variances in boundary conditions effect high strain regions in the model. The research goal is to understand the response of trabecular bone to mechanical stimuli. This has relevance for paleontologists to infer habitual locomotion of ancient people and animals, and in treatment strategies for populations with fragile bones such as the elderly.



- 5 million quadratic, 8 noded elements
- Model created with custom Matlab application that converts 25^3 micro CT images into voxel-based finite element models

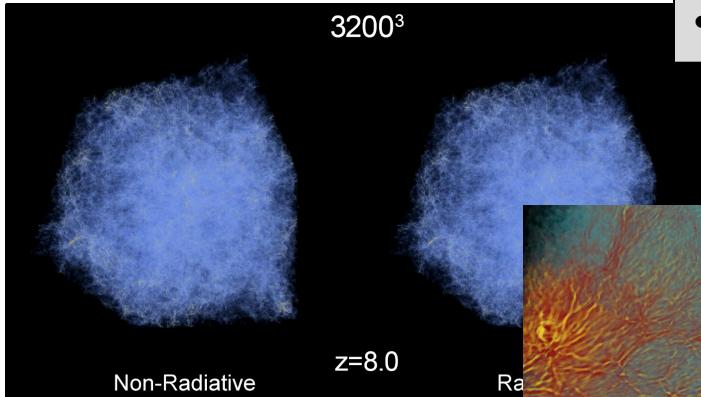
V	M	F
C	T	L

Source: Matthew Goff, Chris Hernandez. Cornell University. Used by permission. 2012

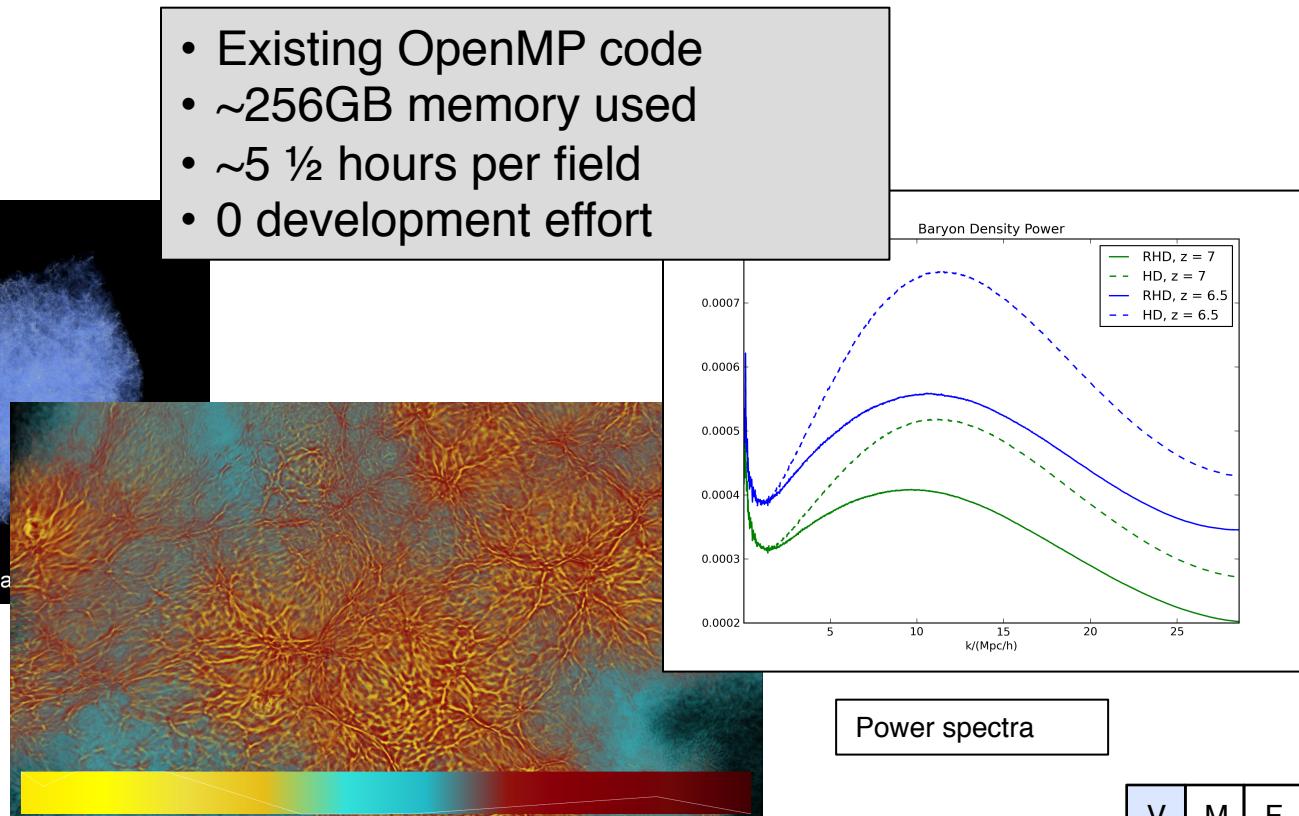
Cosmology simulation - matter power spectrum measurement using vSMP

Goal is to measure the effect of the light from the first stars on the evolution of the universe. To quantitatively compare the matter distribution of each simulation, we use radially binned 3D power spectra.

- 2 simulations
- 3200^3 uniform 3D grids
- 15k+ files each



- Existing OpenMP code
- ~256GB memory used
- ~5 ½ hours per field
- 0 development effort

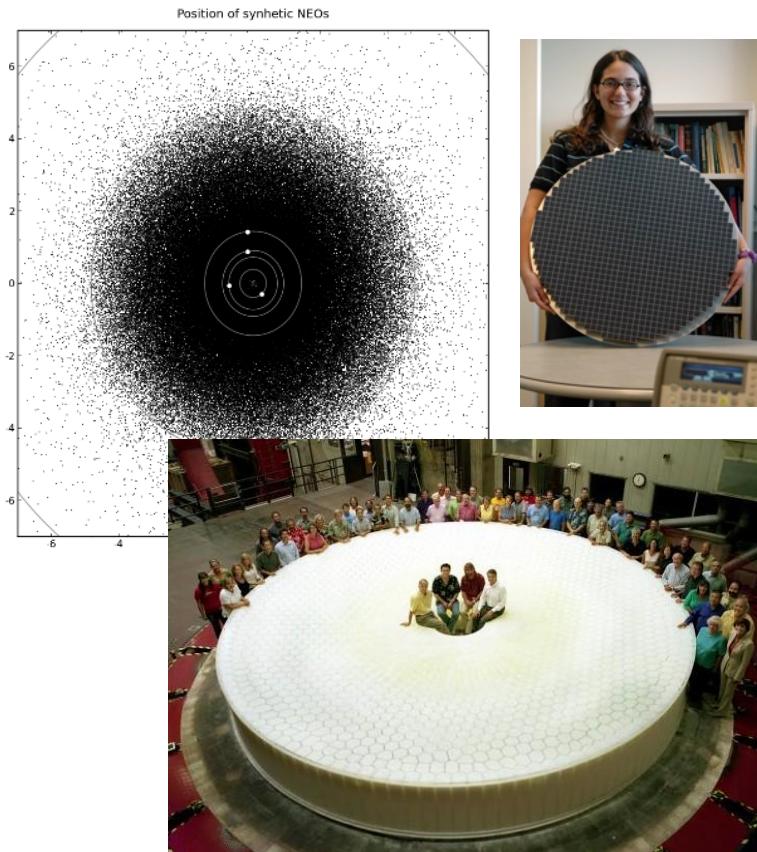


V	M	F
C	T	L

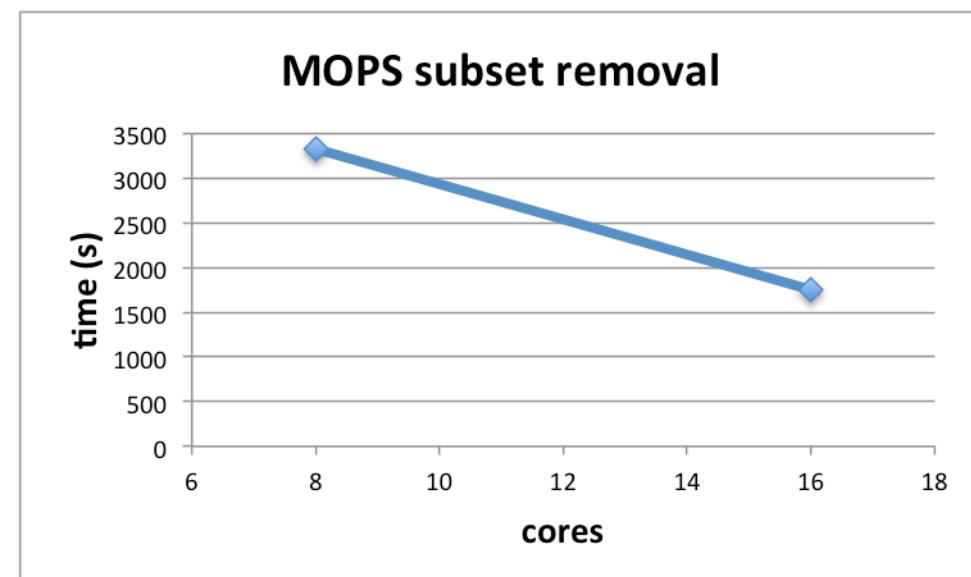
Source: Rick Wagner, Michael L. Norman. SDSC.

LSST – Moving Object Pipeline System

Images collected by the Large Synoptic Survey Telescope (LSST) will be processed using the Moving Object Pipeline System (MOPS). Detections from consecutive nights are grouped together into tracks that potentially represent small portions of the asteroids' sky-plane motion



Run time for subset removal algorithm scales almost linearly out to 16 cores

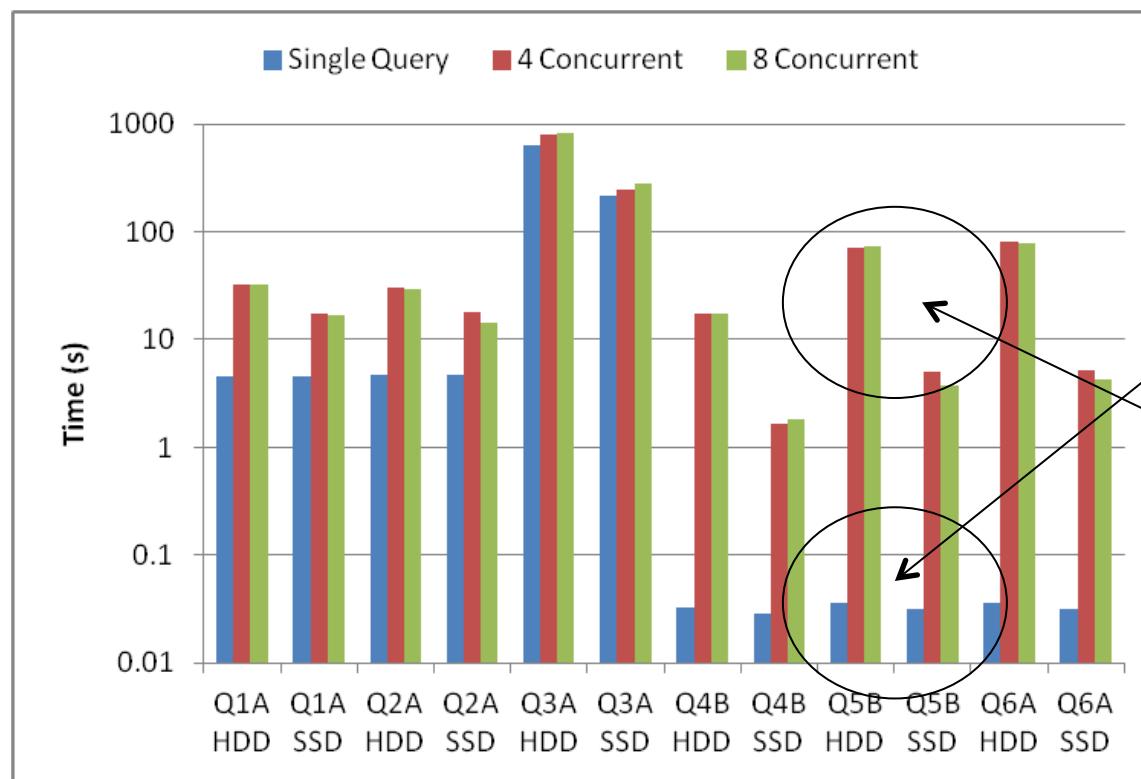


V	M	F
C	T	L

Source: Jonathan Myers, LSST Used by permission. 6/4/2012

PDB Query Comparisons, with DB2 Database on two Gordon I/O Nodes: One with HDD's, One with SSD's

The Protein Data Bank (PDB): Is the single worldwide repository of information about the 3D structures of large biological molecules. These are the molecules of life that are found in all organisms. Understanding the shape of a molecule helps to understand how it works.



- For single queries, HDD and SSD perform about the same.
- For concurrent queries, SSD's achieve big speedup.
- Q5B is > 10x, and performance varies by type of query

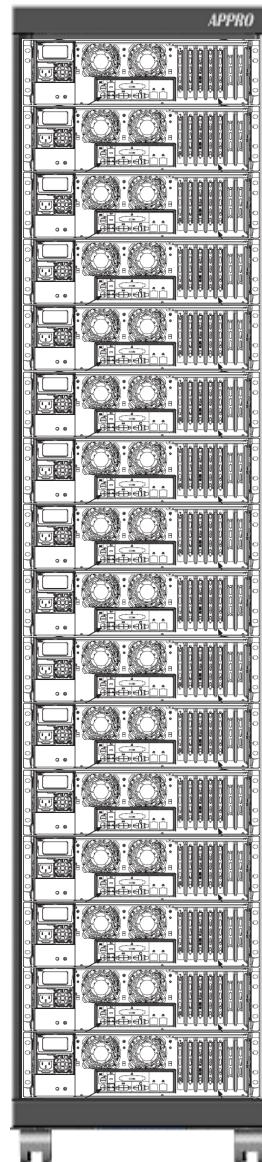
V	M	F
C	T	L

Source: Vishwinath Nandigam, San Diego Supercomputer Center. 2011

Gordon Rack Layout



16 compute node racks
4 I/O node racks
1 service rack



SDSC CN Rack SAN DIEGO SUPERCOMPUTER CENTER Service Nodes Rack

Compute node racks:
4 Appro subracks
64 blades

ION racks:
16 Gordon I/O nodes

Service rack:
4 login nodes
2 NFS servers
2 Scheduler nodes
2 management nodes

Based on Appro GreenBlade™ 8000 Series

designed for improved reliability and energy efficiency

Front View

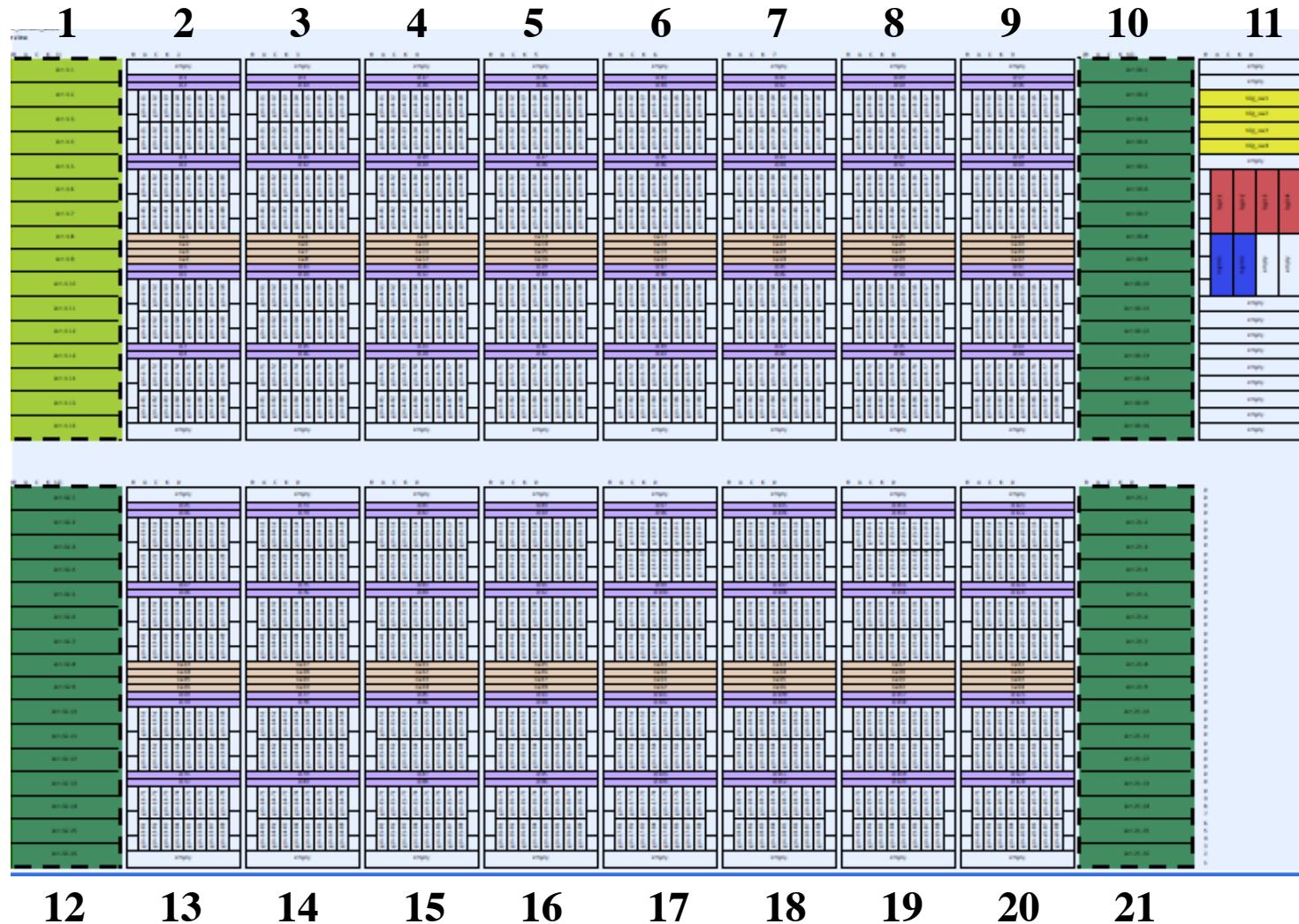


Rear View



- 8RU Subrack
- Supports 16x 2P Intel Sandy-Bridge Blades
- Support for up to six high-efficiency 1625W hot-swappable PS in N+1 configuration
- Support for dual-redundant platform management modules
- Supports six hot-swappable, redundant fan modules
- Shared reduces power consumption by up to 20W per blade over previous design

Gordon Rack Layout



Gordon naming conventions

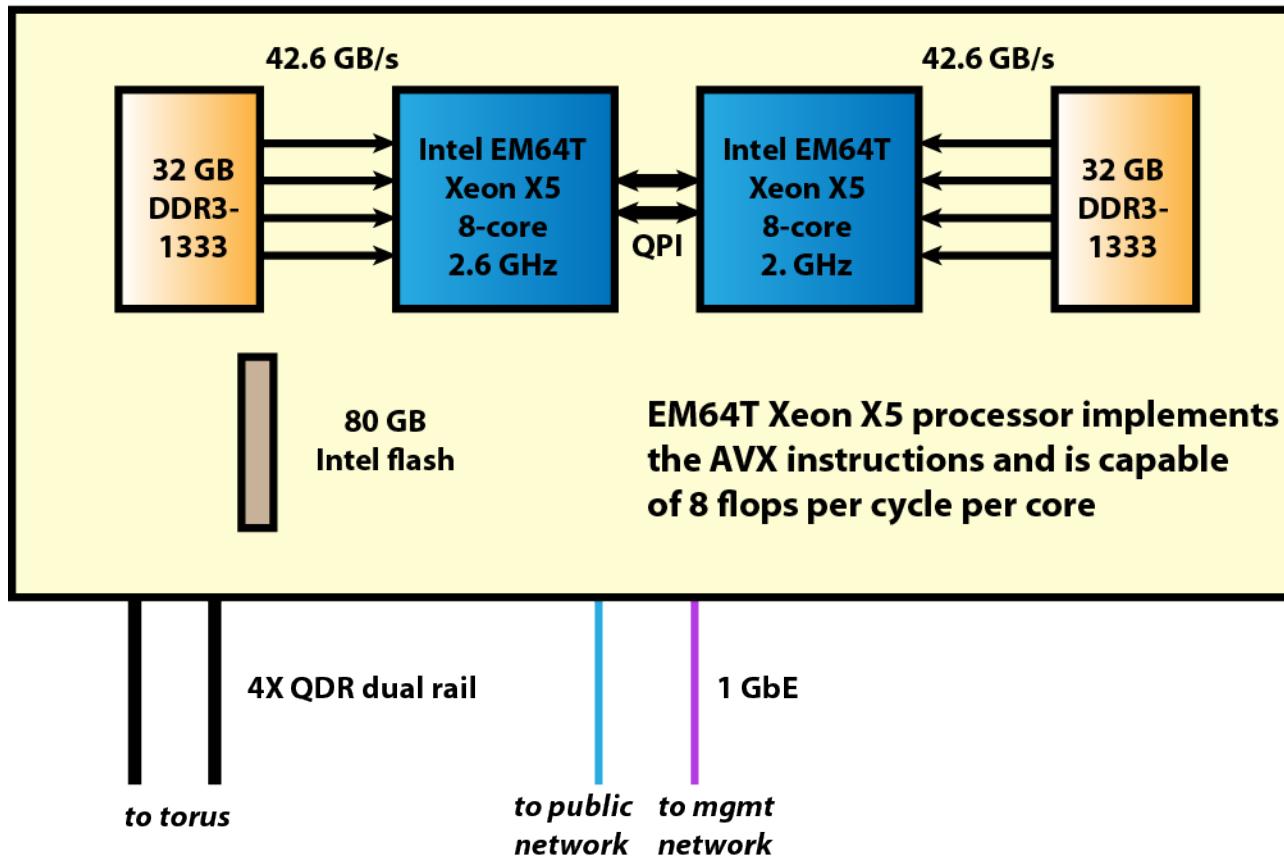
		empty	
		ib1	
		ib2	
	gcn-2-31	gcn-2-21	gcn-2-11
	gcn-2-32	gcn-2-22	gcn-2-12
	gcn-2-33	gcn-2-23	gcn-2-13
	gcn-2-34	gcn-2-24	gcn-2-14
	gcn-2-35	gcn-2-25	gcn-2-15
	gcn-2-36	gcn-2-26	gcn-2-16
	gcn-2-37	gcn-2-27	gcn-2-17
	gcn-2-38	gcn-2-28	gcn-2-18

Gordon Compute Node
rack 2, row 2, node 6



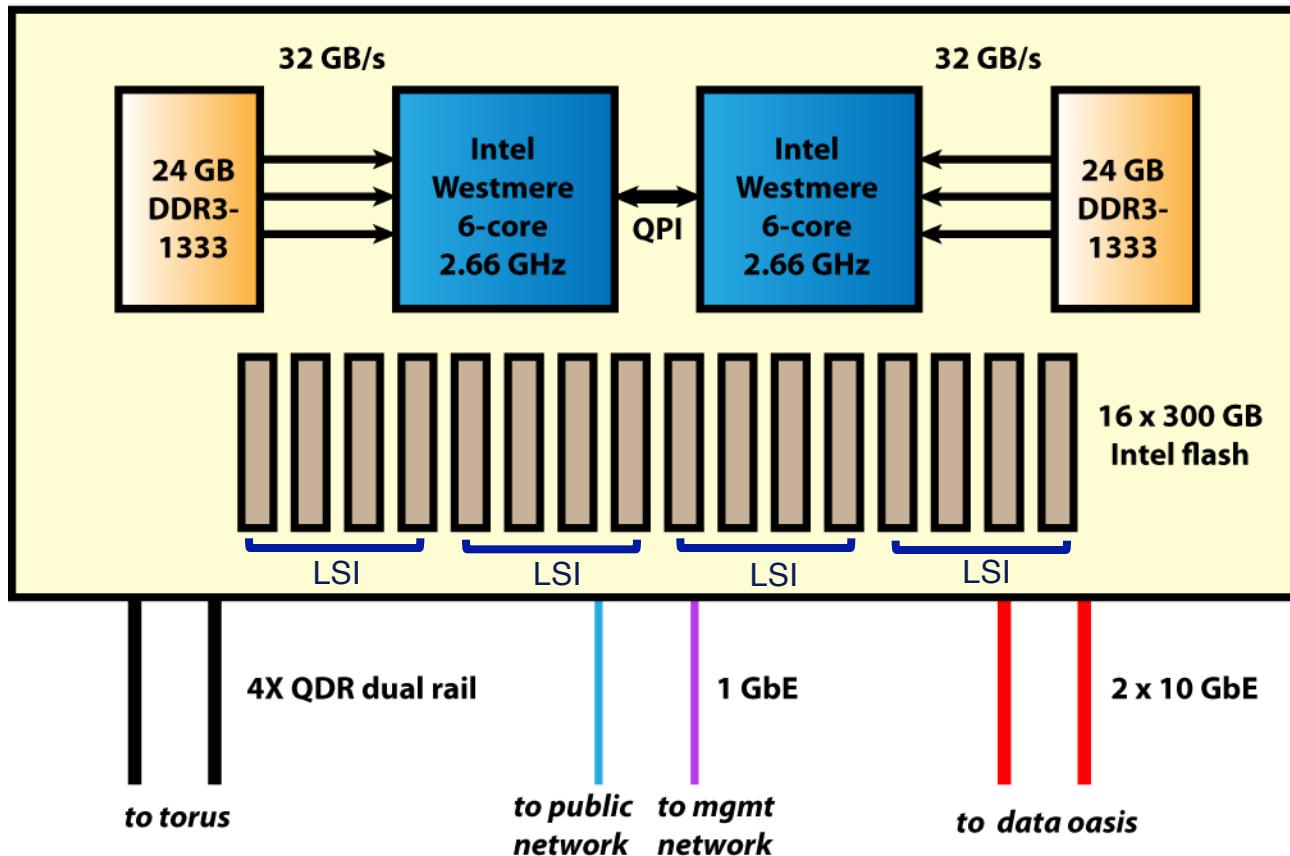
I/O Node
rack 1, node 3

Gordon compute node



For more information on AVX, see <http://software.intel.com/en-us/avx/>

Gordon I/O node

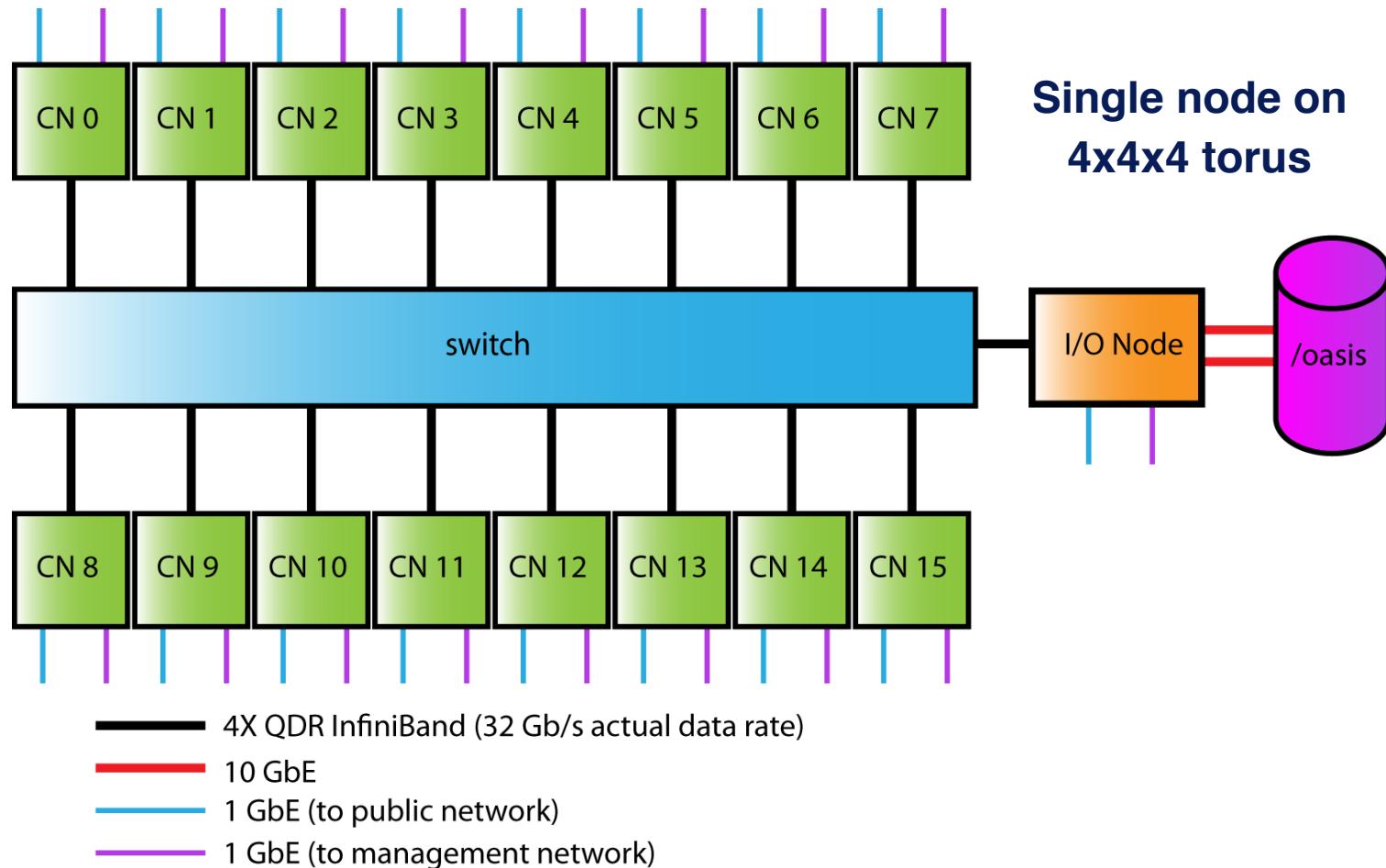


summary

48 GB DRAM
12 cores
2.66 GHz
4.8 TB flash

Bonded into single channel
~ 1.6 GB/s bandwidth

Simplified single rail view of Gordon connectivity showing routing between compute nodes on same switch, I/O node, and data oasis.



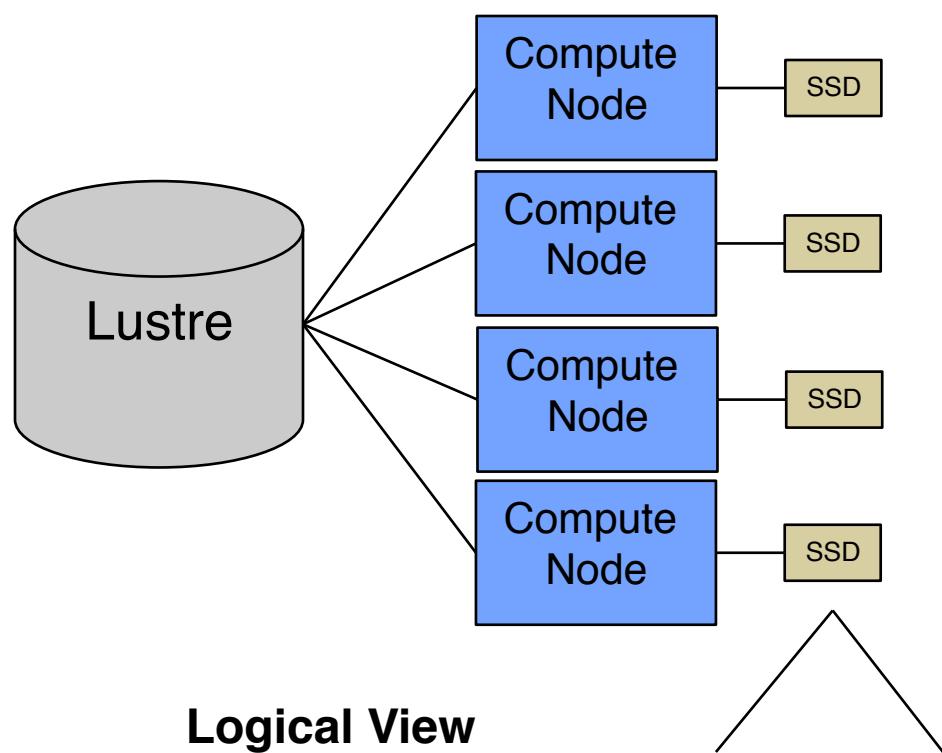
(Some) SSDs are a good fit for data-intensive computing



	Flash Drive	Typical HDD	Good for Data Intensive Apps
Latency	<.1 ms	10 ms	✓
Bandwidth (r/w)	270 / 210 MB/s	100-150 MB/s	✓
IOPS (r/w)	38,500 / 2000	100	✓
Power consumption (when doing r/w)	2-5 W	6-10 W	✓
Price/GB	\$3/GB	\$.50/GB	-
Endurance	2-10PB	N/A	✓
Total Cost of Ownership	Jury is still out.		

Exporting Flash

Model A: One SSD per Compute Node



- One 300 GB flash drive exported to each compute node appears as a local file system
- Lustre parallel file system is mounted identically on all nodes.
- Data is purged at the end of the run

Use cases:

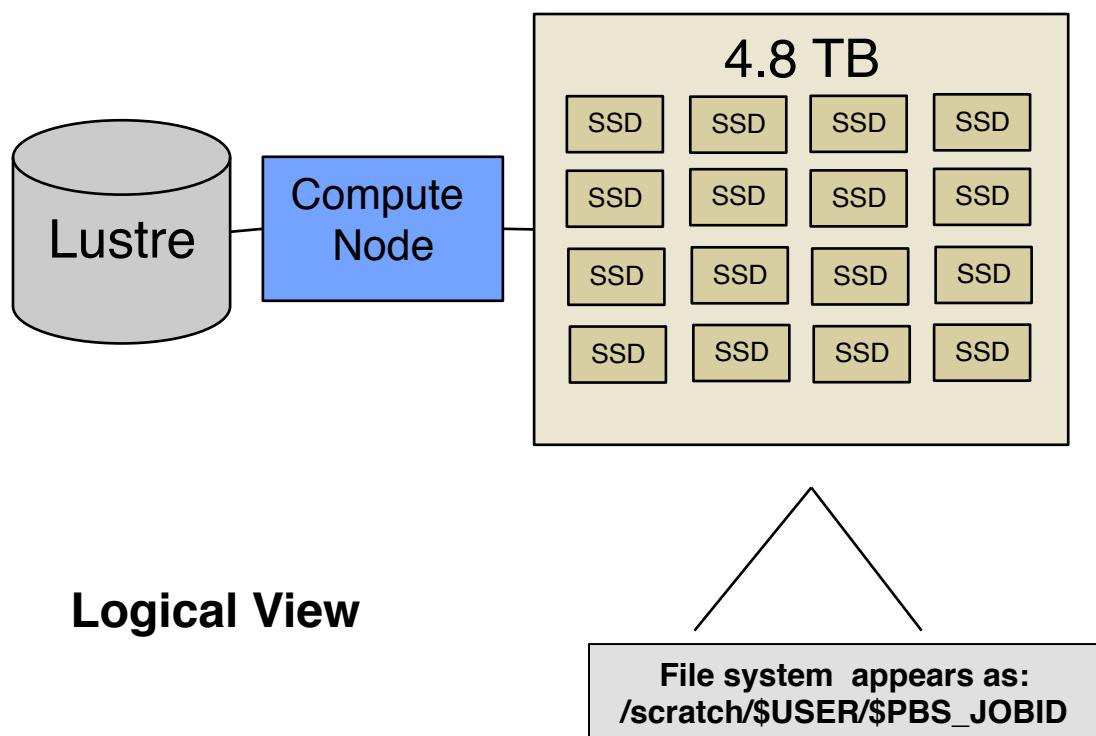
- Applications that need local, temporary scratch
- Gaussian
- Abaqus

This is what we'll be using for most of the summer school

File system appears as:
/scratch/\$USER/\$PBS_JOBID

Exporting Flash

Model B: 16 SSD's for 1 Compute Node

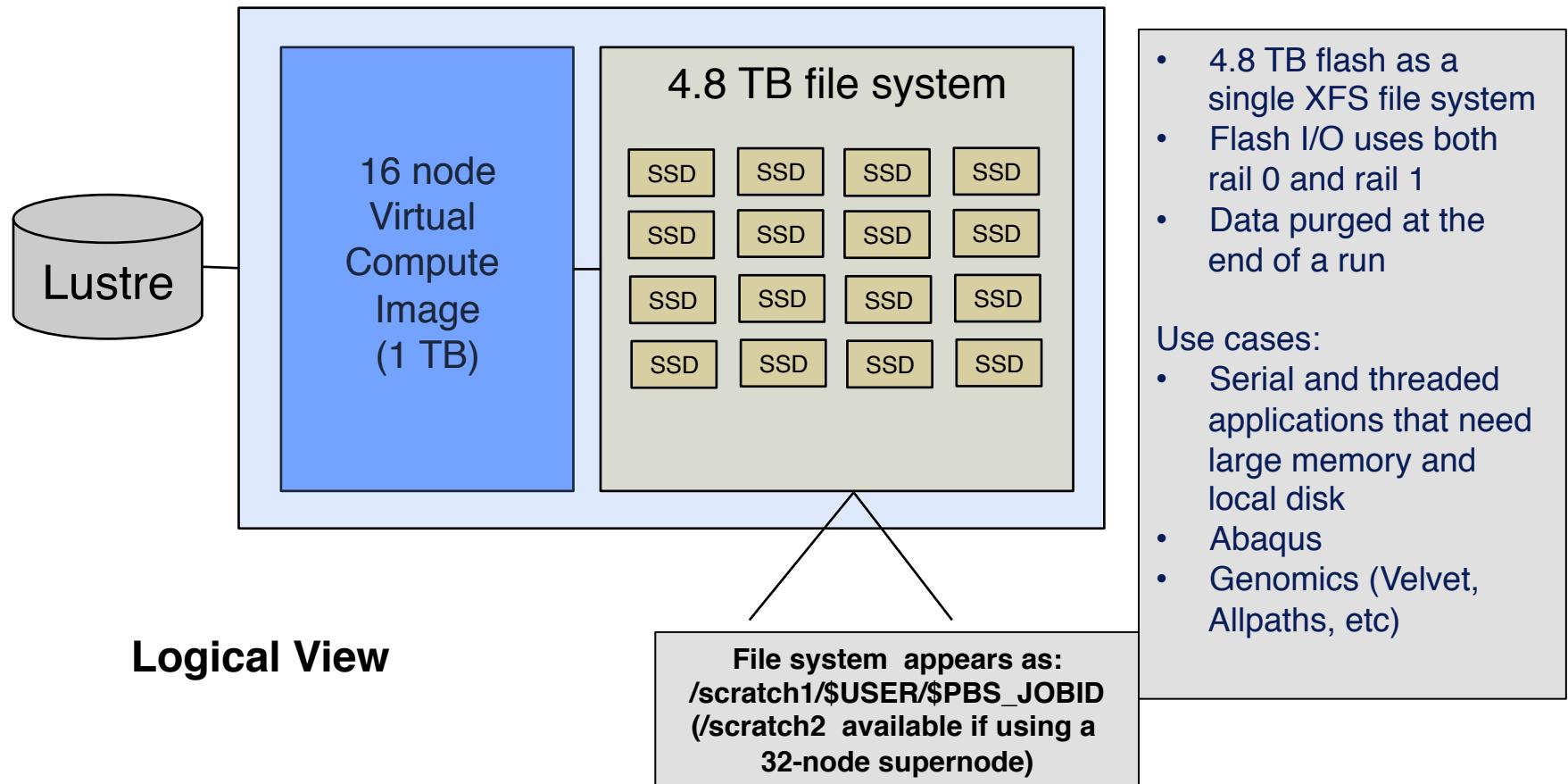


- 16 SSD's in a RAID0 appear as a single 4.8 TB file system to the compute node.
- Flash I/O and Lustre traffic uses Rail 1 of the torus.

Use cases:

- Database
- Data mining
- Gaussian
- Abaqus

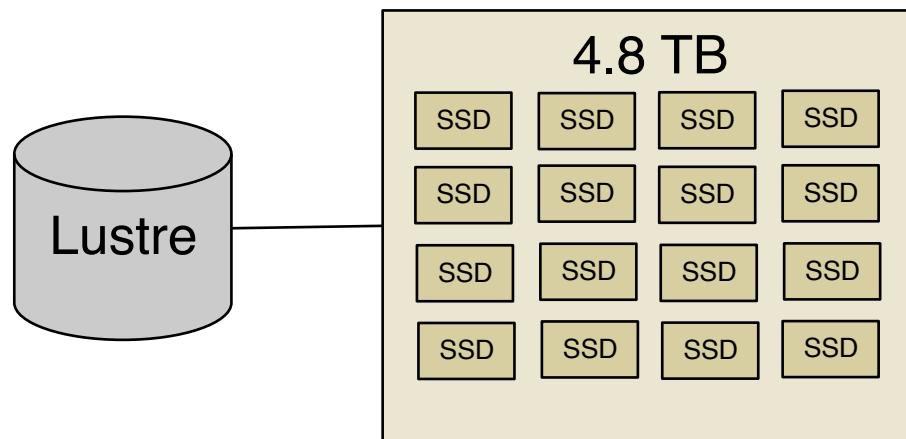
Exporting Flash Model C: 16 SSD's within a vSMP Supernode



Exporting Flash

Model D: Dedicated I/O node

******A new allocations model for data intensive computing ******



Logical View

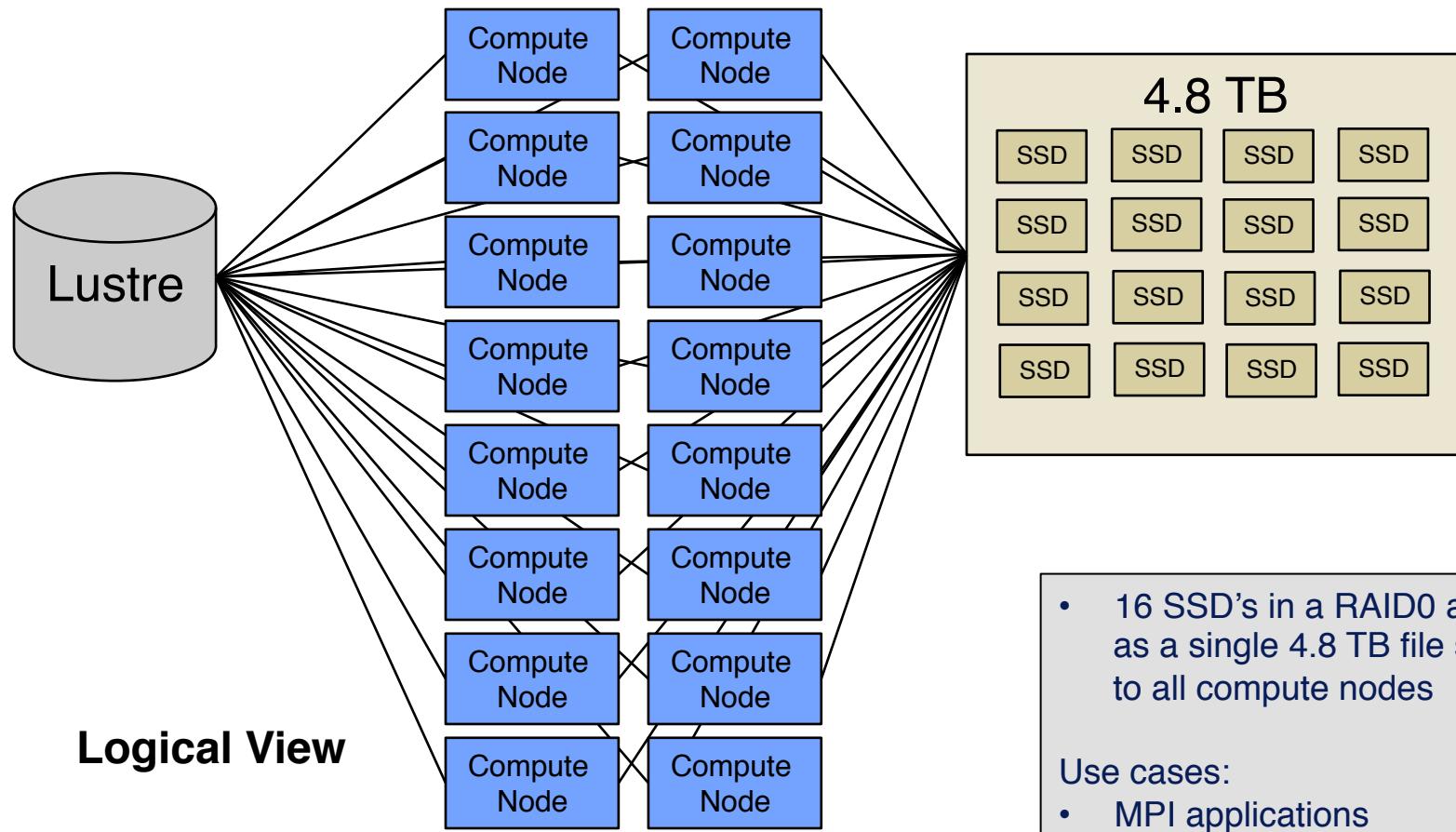
- I/O node is allocated for up to one year
- Users run applications directly on the I/O node
- Users may request dedicated compute nodes or access them through the scheduler
- Data is persistent

Use cases:

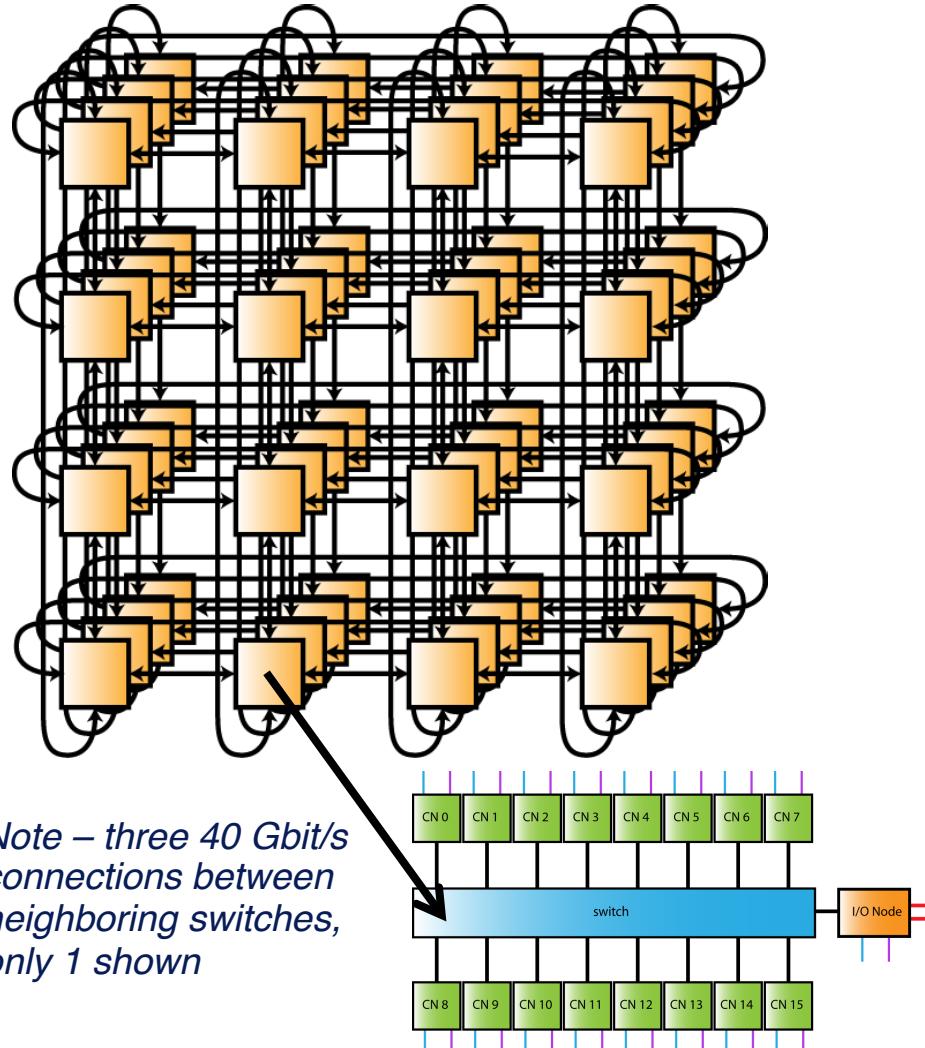
- Database
- Data mining

Exporting Flash

**Model E: 16 SSD's/ 16 compute node –
Flash mounted as OCFS parallel file system
(tested but not currently deployed)**



3D Torus Interconnect



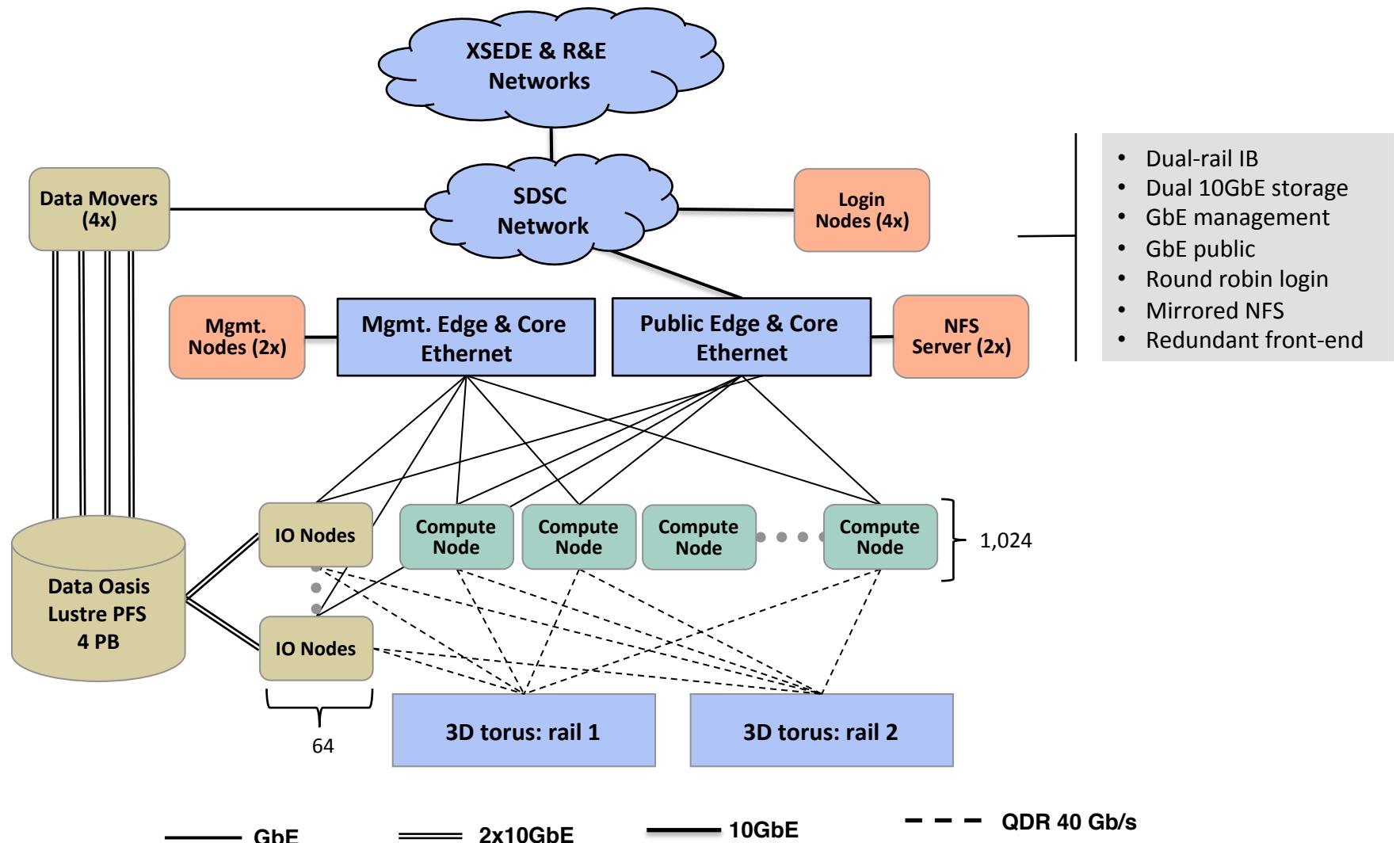
Gordon switches connected in dual rail 4x4x4 3D torus

Maximum of six hops to get from one node to furthest node in cluster

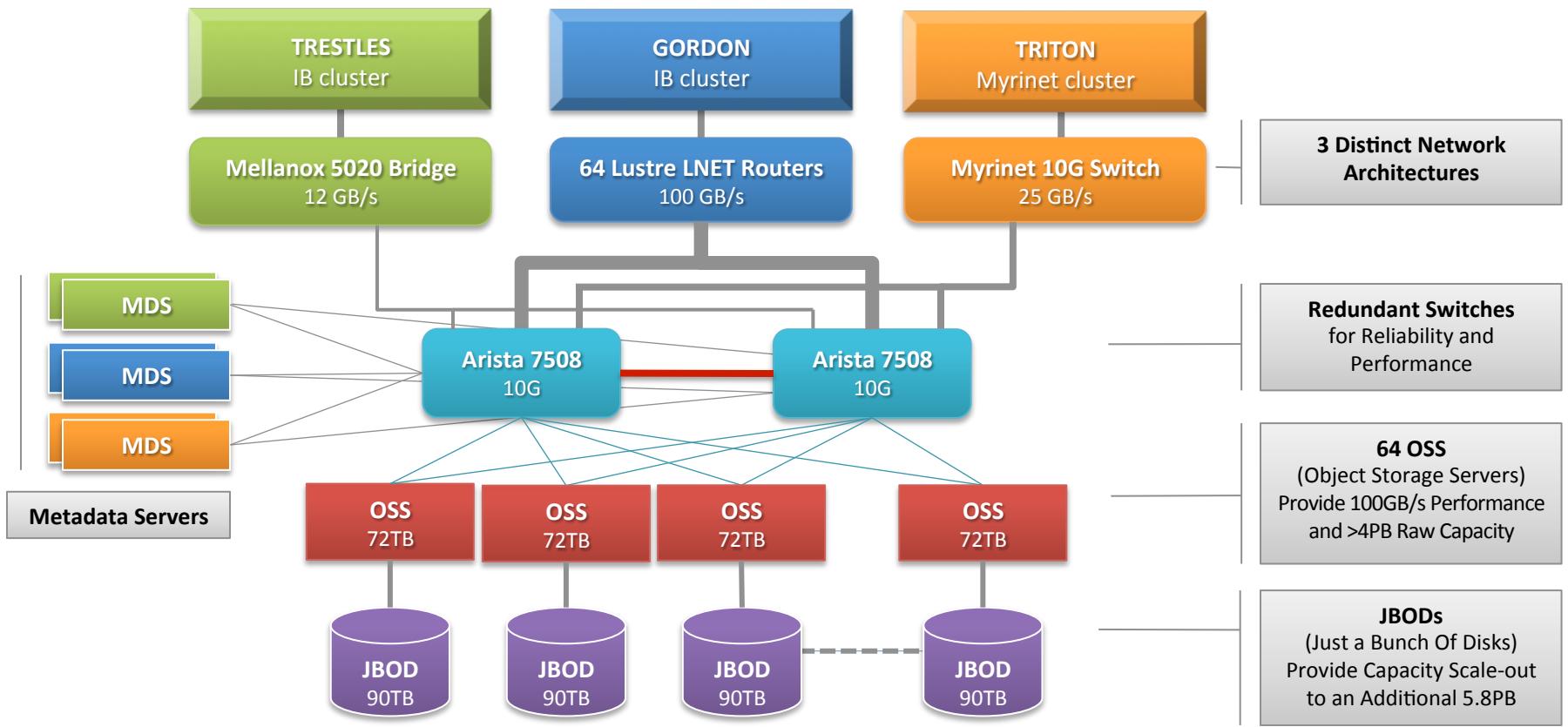
Fault tolerant, requires up to 40% fewer switches and 25-50% fewer cables than other topologies

Scheduler will be aware of torus geometry and assign nodes to jobs accordingly

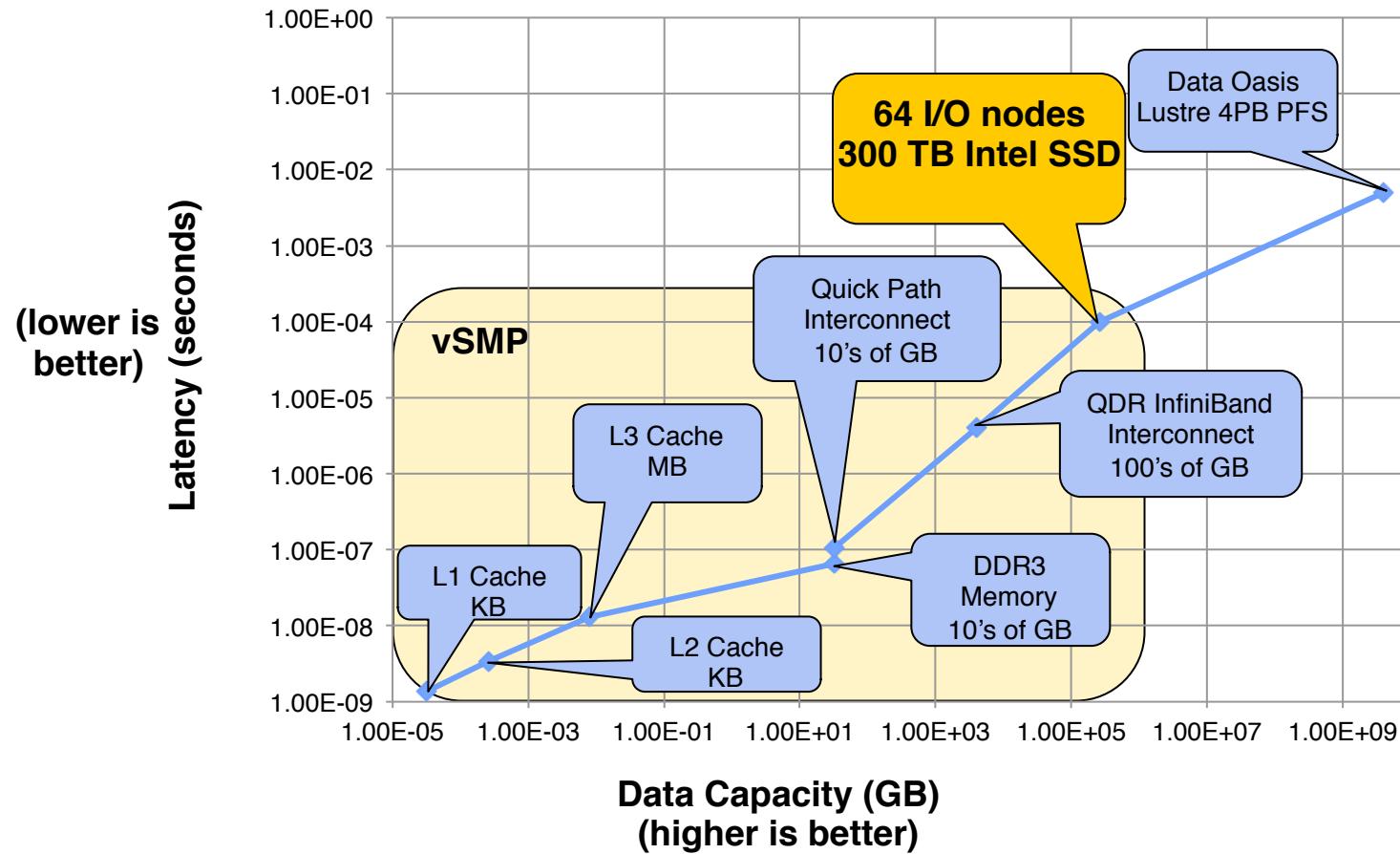
Gordon Network Architecture



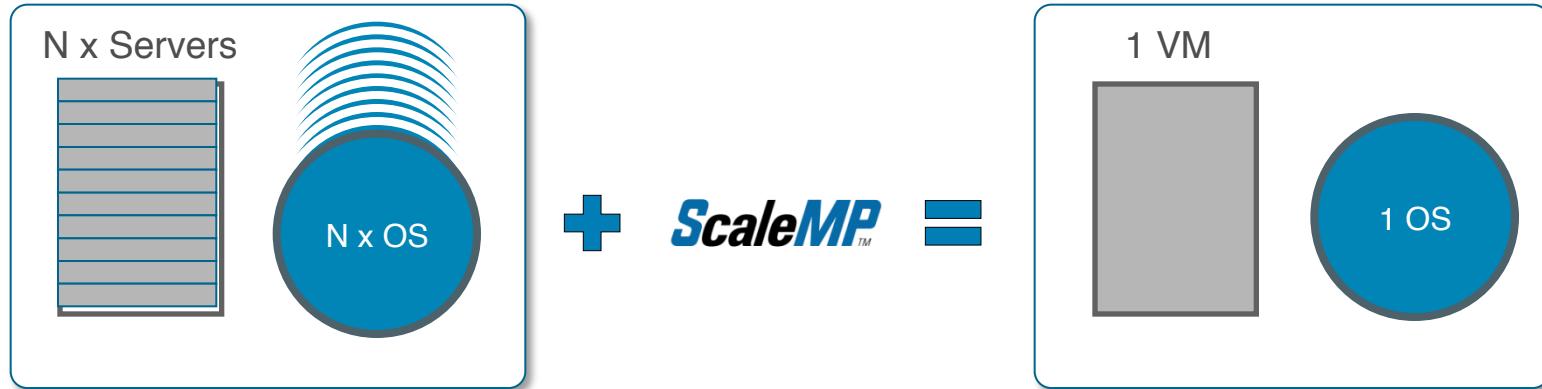
Data Oasis Heterogeneous Architecture Lustre-based Parallel File System



vSMP and Flash Bridge the Latency & Capacity Gap

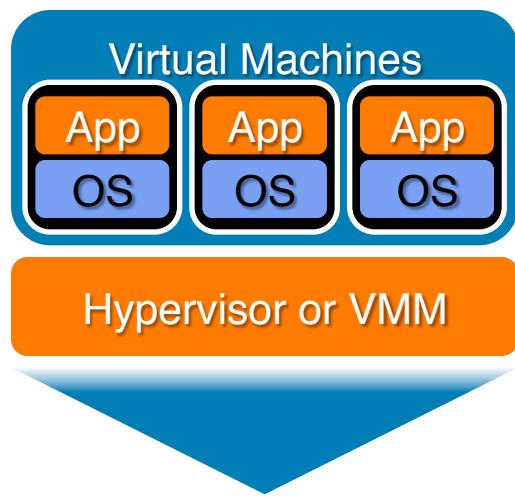


Introduction to vSMP

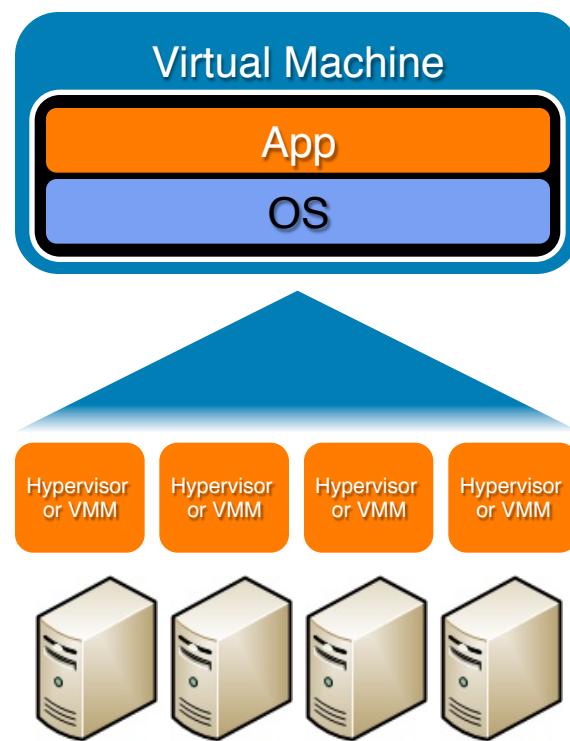


Virtualization **software** for **aggregating** multiple **off-the-shelf** systems into a single virtual machine, providing improved usability and higher performance

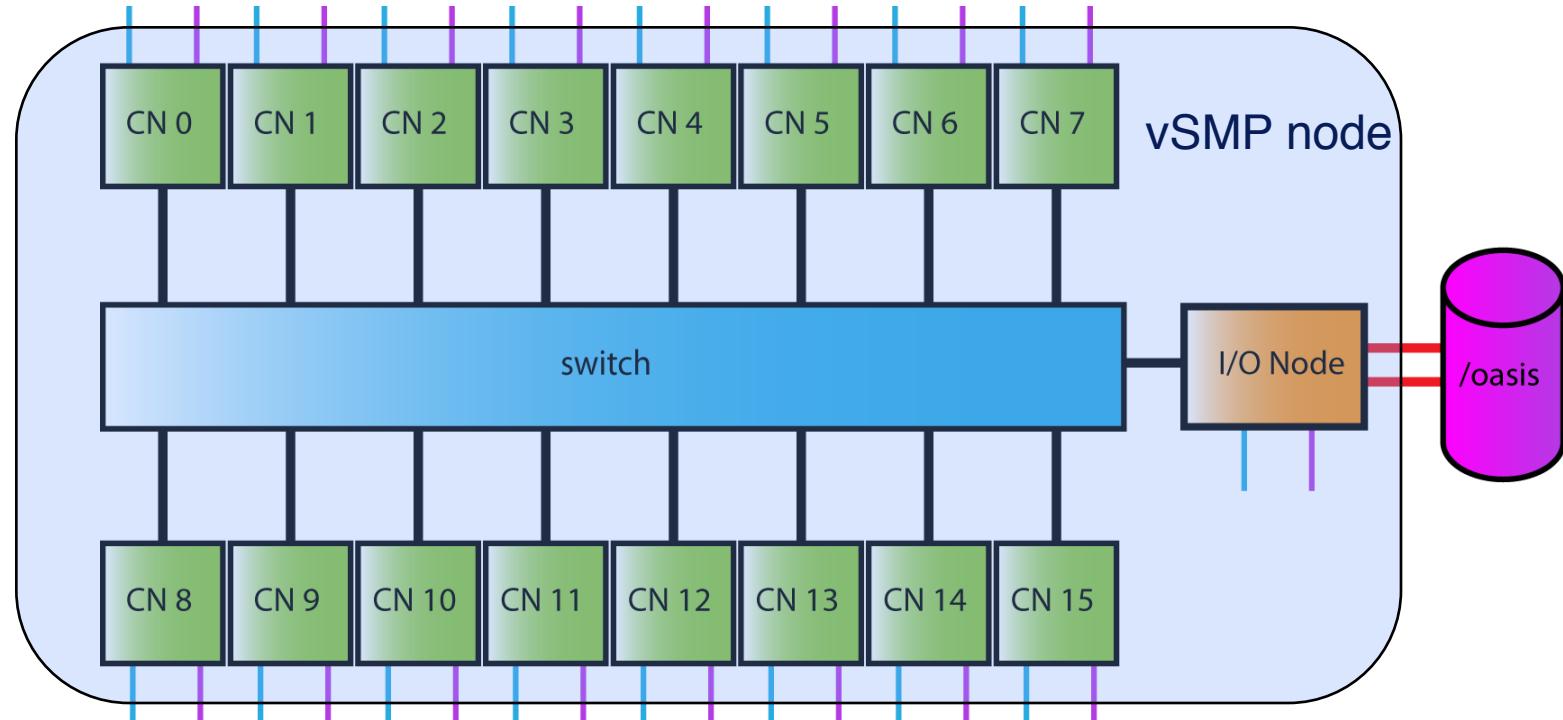
PARTITIONING



AGGREGATION

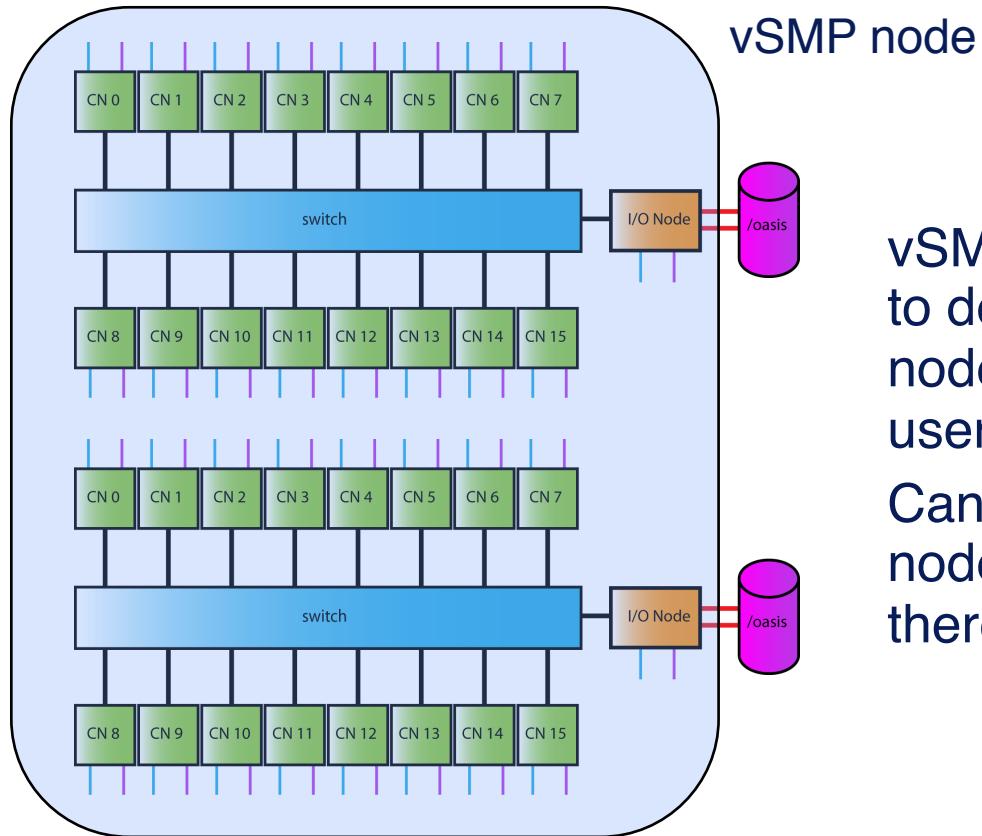


vSMP node configured from 16 compute nodes and one I/O node



To user, logically appears as a single, large SMP node

vSMP node configured from 32 compute nodes and two I/O node



vSMP software provides flexibility to deploy logical shared memory nodes in sizes demanded by users.

Can potentially configure vSMP nodes on the fly (but not quite there yet)

To user, logically appears as a single, large SMP node with ~2 TB memory (32×64 GB) and 512 compute cores (32×16)

Overview of a vSMP node

```
[diag@gcn-17-51 ~]$ vsmpversion
```

```
vSMP Foundation: 4.0.220.0 (Apr 03 2012 19:05:33)
```

```
System configuration:
```

```
Boards: 17
```

```
    16 x processor board
```

```
    1 x memory board
```

```
Processors: 32 (out of 34), Cores: 256 (out of 268)
```

```
    32 x Intel(R) Xeon(R) CPU E5-2670 0 @ 2.60GHz Stepping 06
```

```
    2 x Intel(R) Xeon(R) CPU X5650 @ 2.67GHz Stepping 02
```

```
Memory (MB): 957803 (out of 1096988), Cache: 100273, Private: 38912
```

```
    16 x 65491MB
```

```
    1 x 49132MB
```

```
Host bridge:
```

```
    16 x VID/DID=8086/3c00
```

```
    1 x VID/DID=8086/3406
```

```
Link Rate: 2 x 40Gb/s
```

```
Boot device: [HDD] ATA INTEL SSDSA2CW08
```

```
Serial number: 0
```

```
System key:
```

```
Supported until: -
```

vSMP node built from
16 compute nodes + 1 I/O nodes

Overview of a vSMP node

```
[diag@gcn-17-51 ~]$ grep processor /proc/cpuinfo | tail
processor      : 246
processor      : 247
processor      : 248
processor      : 249
processor      : 250
processor      : 251
processor      : 252
processor      : 253
processor      : 254
processor      : 255
[diag@gcn-17-51 ~]$ head /proc/meminfo
MemTotal:      967227852 kB
MemFree:       908057780 kB
Buffers:        170032 kB
Cached:        23341688 kB
SwapCached:      0 kB
Active:        44385452 kB
Inactive:      11914668 kB
Active(anon):   32792416 kB
Inactive(anon):  3464 kB
Active(file):   11593036 kB
[diag@gcn-17-51 ~]$
```

Logging in to Gordon

- Login to gordon.sdsc.edu
- You will be automatically directed to one of four identical login nodes gordon- $\text{In}[1-4]$.sdsc.edu
- Secure shell users should feel free to append their public RSA key to their `~/.ssh/authorized_keys` file to enable access from authorized hosts without having to enter their password (already done for some of you)
- Login nodes have same architecture as compute nodes
- Do not use login nodes for computationally intensive jobs

Gordon file systems

Gordon provides four different file systems that differ by capacity, performance, backup policy and persistence

- Home
- Oasis project
- Oasis scratch
- Flash scratch

When I/O performance and storage capacity are important, the lecturers will tell you which file system to use

Home file system

- \$HOME (/home/\$USER)
- Persistent storage. Use for source code, small data sets,
- Backed up nightly
- Quotas not currently enforced, but keep usage below 100 GB
- NFS filesystem, mounted on multiple machines, shared by many users. Not meant for I/O intensive jobs

```
[sinkovit@gordon-ln3 ~]$ echo $USER  
sinkovit
```

```
[sinkovit@gordon-ln3 ~]$ echo $HOME  
/home/sinkovit
```

Data Oasis (Lustre) parallel file systems

- Scratch: /oasis/scratch/\$USER/temp_project
- Project: /oasis/projects/nsf/[project]/\$USER
- Semi-persistent (scratch) to persistent (projects)
- High-performance, parallel file systems, capable of delivering 1.6 GB/s per object storage server
- Default allocation of 500 GB, shared among all users of an XSEDE allocations award. Multiple TB upon request
- Ideal for large sequential reads/writes, large files
- Poor performance for random I/O, large numbers of files

```
[sinkovit@gordon-ln3 ~]$ ls -d /oasis/scratch/sinkovit/temp_project  
/oasis/scratch/sinkovit/temp_project
```

```
[sinkovit@gordon-ln3 ~]$ ls -d /oasis/projects/nsf/sds136/sinkovit  
/oasis/projects/nsf/sds136/sinkovit
```

Flash file systems

- `/scratch/$USER/$PBS_JOBID`
- Created automatically upon start of Torque job
- Persistent only for duration of job
- High bandwidth, excellent random I/O performance
- Size depends on flash export model, 280 GB minimum
- Ideal for scratch space, staging repeatedly used data
- Be sure to account for copy-in/copy-out time

```
[sinkovit@gcn-13-28 sinkovit]$ df -h /scratch
Filesystem      Size  Used Avail Use% Mounted on
/dev/sdb1        280G   33M  280G   1% /scratch

[sinkovit@gcn-13-28 sinkovit]$ ls -ld /scratch/$USER/$PBS_JOBID
drwx----- 2 sinkovit root /scratch/sinkovit/157431.gordon-fe2.local
```

Using modules

The modules package provides for dynamic management of your Linux environment. Unless there is a compelling reason for you to run your own versions of common software, we strongly encourage you to use modules to load and run the versions that have been installed by our staff.

- Built using optimal compiler options (e.g. AVX support)
- Linked to appropriate libraries (e.g. MVAPICH2_IB)
- Environment variables automatically set correctly

module list lists currently loaded modules

module load [mod] adds module to your environment

module unload [mod] removes module from your environment

```
[user@gordon-ln1 ~]$ module list
1) binutils/2.22  2) intel/2011  3) mvapich2_ib/1.8a1p1

[user@gordon-ln1 ~]$ module load amber mopac
[user@gordon-ln1 ~]$ module list
 1) binutils/2.22  3) mvapich2_ib/1.8a1p1  5) amber/11
 2) intel/2011      4) mopac/2009

[user@gordon-ln1 ~]$ module unload amber mopac
[user@gordon-ln1 ~]$ module list
 1) binutils/2.22  2) intel/2011  3) mvapich2_ib/1.8a1p1
```

module avail lists the modules that are available. Default is the version that will be loaded if no version number provided

```
[user@gordon-ln1 ~]$ module avail

----- /opt/modulefiles/mpi/.intel -----
mpich2_ib/1.0.7(default) openmpi_ib/1.4.1(default)
mvapich2_ib 1.8a1p1(default)

----- /opt/modulefiles/applications/.intel -----
atlas/3.9.45(default)      hdf5/1.8.3(default)      petsc/3.2.p3(default)
fftw/2.1.5                 lapack/3.3.1(default)    scalapack/2.0.1(default)
[ --- additional lines not shown ---]

----- /opt/modulefiles/applications -----
R/2.13.1(default)          ddt/3.1(default)        mopac/2009(default)
amber/11(default)          fsa/1.15.2(default)    namd/2.6
[ --- additional lines not shown ---]

----- /opt/modulefiles/compilers -----
gnu/4.1.2(default)         gnu/4.6.1   intel/2011(default) pgi/11.9(default)
```

Unloading a module makes dependent modules unavailable.
For example, note that MPI libraries and numerical libraries
are no longer listed after unloading intel module

```
[user@gordon-ln1 ~]$ module unload intel
Unloading compiler-dependent module mvapich2_ib/1.8a1p1

[user@gordon-ln3 ~]$ module avail

----- /opt/modulefiles/applications -----
R/2.13.1(default)          ddt/3.1(default)          mopac/2009(default)
amber/11(default)          fsa/1.15.2(default)        namd/2.6
[ --- additional lines not shown --- ]

----- /opt/modulefiles/compilers -----
gnu/4.1.2(default)   gnu/4.6.1    intel/2011(default) pgi/11.9(default)
```

module display [mod] gives information about a module

```
[user@gordon-ln1 ~]$ module display binutils
-----
/opt/modulefiles/applications/binutils/2.22:

prepend-path      PATH /opt/binutils/bin
prepend-path      LD_LIBRARY_PATH /opt/binutils/lib
prepend-path      LD_LIBRARY_PATH /opt/binutils/lib64
-----
```

```
[user@gordon-ln1 ~]$ module display gaussian
-----
/opt/modulefiles/applications/gaussian/09.C.01:

prepend-path      PATH /opt/gaussian/g09
setenv            GAUSS_EXEDIR /opt/gaussian/g09
setenv            G09_BASIS /opt/gaussian/g09/basis
-----
```

module swap [mod1] [mod2] replaces mod1 with mod2

module purge unloads all modules

```
[user@gordon-ln1 ~]$ module list
1) binutils/2.22      2) intel/2011          3) mvapich2_ib/1.8a1p1

[user@gordon-ln1 ~]$ module swap mvapich2_ib openmpi_ib
[user@gordon-ln1 ~]$ module list
1) binutils/2.22      2) intel/2011          3) openmpi_ib/1.4.1

[user@gordon-ln1 ~]$ module purge
[user@gordon-ln1 ~]$ module list
No Modulefiles Currently Loaded.-
```

One last word about modules ...

The modules command is a shell function that is defined at login time and locate will not find it in the usual locations (e.g. /bin, /usr/bin, /sbin)

If you ever get the error “*module: command not found*”, then run the following (you may need to add this to your shell scripts that execute module commands)

```
source /etc/profile.d/modules.sh
```

Gordon software

- Numerical libraries: ATLAS, FFTW, GSL, LAPACK, PETSc, SPRNG, ScaLAPACK, SuperLU, MKL
- Partitioning and frameworks: ParMETIS, Trilinos
- File formats/libraries: netCDF, HDF4, HDF5
- Molecular dynamics / chemistry: AMBER, GAMESS, Gaussian, LAMMPS, NAMD, Gromacs, MOPAC, NWChem
- Visualization: Visit, ParaView (coming soon)
- Languages: R, Octave
- Standard compilers: GNU, Intel, PGI
- MPI: MPICH2, MVAPICH2, OpenMPI

http://www.sdsc.edu/us/resources/gordon/gordon_software_packages.html

Gordon software suggestions

- **Preferred compiler:** Intel
... but if you really need gnu for compatibility, load 4.6.1 or later to get AVX support
- **Preferred MPI:** MVAPICH2 (module mvapich2_ib)
... but if you're building an MPI application for vSMP, then use ScaleMP's specially tuned version of MPICH2
- **Preferred numerical library:** MKL
Tuned for best performance on Intel hardware; contains threaded BLAS, LAPACK, ScaLAPACK, FFTs, etc. Very easy to use, just link with -mkl
... of course, don't even think about writing your own!

Compiling code for Gordon (Intel)

Intel compilers are the default on Gordon and generally give the best performance

- OpenMP enabled using -openmp
- Use -fast or -xHOST to get AVX support
-fast = -xHOST -O3 -ipo -no-prec-div -static
- MKL library linked with -mkl

	Serial	MPI	OpenMP	Hybrid
Fortran	ifort	mpif90	ifort -openmp	mpif90 -openmp
C	icc	mpicc	icc -openmp	mpicc -openmp
C++	icpc	mpicxx	icpc -openmp	mpicxx -openmp

Compiling code for Gordon (PGI)

PGI compilers may give good performance on Gordon and also provide AVX support

- OpenMP enabled using -mp
- Use -fast to get AVX support

	Serial	MPI	OpenMP	Hybrid
Fortran	pgf90	mpif90	pgf90 -mp	mpif90 -mp
C	pgcc	mpicc	pgcc -mp	mpicc -mp
C++	pgCC	mpicxx	pgCC -mp	mpicxx -mp

Compiling code for Gordon (GNU)

GNU compilers are provided mainly for compatibility

- OpenMP enabled using -fopenmp
- Must use version 4.6 or later for AVX support
module load gnu/4.6.1, then compile with -mavx

	Serial	MPI	OpenMP	Hybrid
Fortran	gfortran	mpif90	gfortran -fopenmp	mpif90 -fopenmp
C	gcc	mpicc	gcc -fopenmp	mpicc -fopenmp
C++	g++	mpicxx	g++ -fopenmp	mpicxx -fopenmp

A few comments about compilers

- Always specify an optimization level. In some cases the default is -O0 and performance will be terrible
- Note that the OpenMP options are similar, but vary Intel (-openmp), PGI (-mp), GNU (-fopenmp)
- mpicc, mpif90, and mpicxx are wrappers, not compilers
They call the appropriate serial compilers under the hood
- For production runs, turn off all features that may impact performance (e.g. debug, bounds checking, profiling)

Torque

Torque is a freely available, open-source* distributed resource manager based on the PBS project (from an end user point of view Torque is virtually identical to PBS)

- **qsub** submits jobs to the queue
- **qstat** gets job and queue information
- **qdel** deletes a queued or running job
- **qalter** modifies job attributes
(limited capabilities for regular users)

See www.clusterresources.com/torquedocs
for more than you'll ever want to know about Torque

qsub submits a job to the queue. Torque options can be placed into a job file or specified on command line

```
[user2@gordon-ln1 ~]$ qsub job_script
[user2@gordon-ln1 ~]$ cat job_script

#!/bin/bash                         <-- File is a bash script
#PBS -q normal                      <-- Running in normal queue
#PBS -N jobname                     <-- Name of job as listed by qstat
#PBS -l nodes=1:ppn=16:native       <-- One node, 16 cores/node, native property
#PBS -l walltime=0:10:00             <-- 10 minutes wall time
#PBS -o jobname.out                  <-- stdout file
#PBS -e jobname.err                  <-- stderr file
#PBS -v Catalina_res_bind=xxx       <-- Reservation number
#PBS -V                             <-- Export environment variables
#PBS -M username@sdsc.edu           <-- Address for notification
#PBS -m abe                          <-- Notify on abort, begin and end
#PBS -A sds136                        <-- Account to be charged

# run jobs, move files, general bash commands, etc.
mpirun_rsh -hostfile $PBS_NODEFILE -np 16 ./hello_world
```

qsub can also be used to get an interactive node

```
[user2@gordon-ln1 ~]$ qsub -I -X -l nodes=1:ppn=16:native,walltime=10:00:00  
-q normal -v Catalina_res_bind=xxx,QOS=0 -A sds136  
  
-I                                <-- Interactive job  
-X                                <-- X11 forwarding  
-l nodes=1:ppn=16:native,           <-- 1 node, 16 cores/node, native property  
    walltime=10:00:00                10 hour wall clock  
-q normal                          <-- Normal queue  
-v Catalina_res_bind=xxx,QOS=0    <-- Reservation number  
-A sds136                           <-- Account to be charged  
  
[user2@gordon-ln1 ~]$ cat .bash_profile  
alias ggn='qsub -I -X -l nodes=1:ppn=16:native,walltime=10:00:00 -q normal  
-v Catalina_res_bind=xxx,QOS=0 -A sds136'
```

What happens after you are assigned a node

Users are normally prohibited from directly accessing a node, but an exception is granted after it has been assigned by Torque.

This can be particularly useful if you ever need to monitor a non-interactive job (e.g. using top to determine number of processes, threads, CPU utilization, memory footprint)

```
# Before assignment of node
[sinkovit@gordon-ln4 ~]$ ssh gcn-15-73 cat /etc/security/access.conf
Connection closed by 10.5.102.210

# After assignment of node
[sinkovit@gordon-ln4 ~]$ ssh -q gcn-15-73 cat /etc/security/access.conf
-:ALL EXCEPT root diag sinkovit:ALL
```

qstat gets job and queue information

```
[user2@gordon-ln1 ~]$ qstat -a
```

Job ID	Username	Queue	Jobname	NDS	Req'd			Elap	
					Memory	Time	S	Time	
1.gordon	user1	normal	g09	1	1gb	26:00	R	23:07	
2.gordon	user2	normal	task1	32	--	48:00	R	32:15	
3.gordon	user2	normal	task2	2	1gb	24:00	H	--	
4.gordon	user3	normal	exp17	1	--	12:00	C	01:32	
5.gordon	user3	normal	stats1	8	--	12:00	Q	--	
6.gordon	user3	normal	stats2	8	--	12:00	E	15:27	

```
[user2@gordon-ln1 ~]$ qstat -a -u user2
```

Job ID	Username	Queue	Jobname	NDS	Req'd			Elap	
					Memory	Time	S	Time	
2.gordon	user2	normal	task1	32	--	48:00	R	32:15	
3.gordon	user2	normal	task2	2	1gb	24:00	H	--	

qstat can also provide detailed information about a job

```
[user2@gordon-ln1 ~]$ qstat -f 57016
Job Id: 57016.gordon-fe2.local
  Job_Name = mg6
  Job_Owner = user@gordon-ln2.local
  resources_used.cput = 11:44:57
  resources_used.mem = 266012kb
  [ -- additional lines not shown -- ]

[sinkovit@gordon-ln3 ~]$ qstat -n 57016
                                         Req'd   Req'd   Elap
Job ID    Username Queue    Jobname NDS   Memory Time S Time
-----  -----
57016      user     normal   mg6      1     1gb    3:00 R  1:57
gcn-15-56/15+gcn-15-56/14+gcn-15-56/13+gcn-15-56/12+gcn-15-56/11
+gcn-15-56/10+gcn-15-56/9+gcn-15-56/8+gcn-15-56/7+gcn-15-56/6
+gcn-15-56/5+gcn-15-56/4+gcn-15-56/3+gcn-15-56/2+gcn-15-56/1+gcn-15-56/0
```

qdel deletes a queued, held, or running job

```
[user2@gordon-ln1 ~]$ qstat -a -u user2
                                         Req'd   Req'd   Elap
Job ID    Username Queue     Jobname NDS   Memory Time S Time
-----  -----
2.gordon  user2    normal   task1    32      --   48:00 R 32:15
3.gordon  user2    normal   task2     2       1gb  24:00 H  --
```

```
[user2@gordon-ln1 ~]$ qdel 2
```

```
[user2@gordon-ln1 ~]$ qstat -a -u user2
                                         Req'd   Req'd   Elap
Job ID    Username Queue     Jobname NDS   Memory Time S Time
-----  -----
3.gordon  user2    normal   task2     2       1gb  24:00 H  --
```

qalter modifies job attributes. Regular users have limited capabilities (e.g. reduce wall time for a running job)

```
[user4@gordon-ln1 ~]$ qstat -a 8
                                         Req'd   Req'd   Elap
Job ID    Username Queue    Jobname NDS   Memory Time S Time
-----  -----
8.gordon user4     normal   task1    32    --  10:00 R  6:15
```

```
[user4@gordon-ln1 ~]$ qalter -l walltime=9:00 8
```

```
[user4@gordon-ln1 ~]$ qstat -a 8
                                         Req'd   Req'd   Elap
Job ID    Username Queue    Jobname NDS   Memory Time S Time
-----  -----
8.gordon user2     normal   task1    32    --  09:00 R  6:15
```

http://hipacc.ucsc.edu/ISSAC2012_Program.html - Sinkovitz

- Login to gordon (ssh –X username@gordon.sdsc.edu)
- Verify that \$HOME and /oasis/scratch/\$USER/temp_project exist
- Add **contents** of ~sinkovit/ASTRO/add_to_profile to your .bash_profile
cat ~sinkovit/ASTRO/add_to_profile >> .bash_profile
(be sure to ‘source .bash_profile’)
- Launch an interactive job (gastro alias from add_to_profile)
- Verify that X-forwarding works (e.g. ‘xclock &’)
- While interactive job is running, connect directly to your compute node (gcn-x-xx) from login node
- Verify that /scratch/\$USER/\$PBS_JOBID exists
- Kill interactive job (CTRL-D or exit)
- Experiment with module commands (load, unload, avail, display)
- Try the qstat command – list all jobs (-a), pick a random job and get detailed information (-f) or node list (-n)
- Copy ~sinkovit/ASTRO/HELLO to your home directory. Follow directions to build executables and launch batch jobs
- Check out Gordon user guide <http://www.sdsc.edu/us/resources/gordon/>