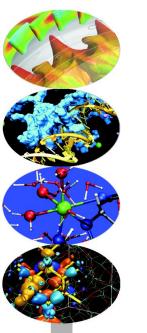


Introduction to CINECA HPC Environment



22nd Summer School on Parallel Computing

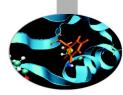
02 September 2013

m.cestari@cineca.it, i.baccarelli@cineca.it





Goals



You will learn:

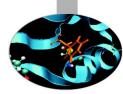
- The basic overview of CINECA HPC systems
- how to login to the school HPC system (PLX)
- basic concepts of the system architecture and user environment that directly affects your work
- how to explore and interact with the software installed on the system
- how to compile a parallel code





Contents

Systems overview (Fermi, Eurora, PLX)



A first step

login file transfer

Introduction to the environment

accounting disk systems module system

Programming environment

compilation compiling/linking issues

Production environment

job script PBS commands

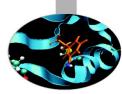
For further info...

useful links and documentation





Fermi characteristics



Model: IBM-BlueGene /Q

Architecture: 10 BGQ Frame with 2 MidPlanes each

Front-end Nodes OS: Red-Hat EL 6.2

Compute Node Kernel: lightweight Linux-like kernel

Processor Type: IBM PowerA2, 1.6 GHz

Computing Nodes: 10.240 with 16 cores each

Computing Cores: 163.840

RAM: 16GB / node; 1GB/core

Internal Network: Network interface

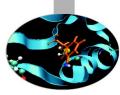
with 11 links ->5D Torus **Disk Space:** more than 2PB of scratch space **Peak Performance:** 2.1 PFlop/s

> ranked #7 in the top 500 supercomputer sites list (June 2012) #12 in the top 500 (June 2013)









EURORA characteristics

Model: Eurora prototype

Architecture: Linux Infiniband Cluster

Processors Type:

- Intel Xeon (Eight-Core SandyBridge) E5-2658 2.10 GHz (Compute)
- Intel Xeon (Eight-Core SandyBridge) E5-2687W 3.10 GHz (Compute)
- Intel Xeon (Esa-Core Westmere) E5645 2.4 GHz (Login)

Number of nodes: 64 Compute + 1 Login

Number of cores: 1024 (compute) + 12 (login)

Number of accelerators: 114 nVIDIA Tesla K20 (Kepler) + 14 Intel Xeon Phi (MIC)

RAM: 1.1 TB (16 GB/Compute node + 32GB/Fat node)

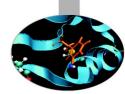
OS: CentOS release 6.3, 64 bit

ranked #1 in the Green 500 chart – The world's most energy-efficient supercomputers (June 2013), 3210 MFlops/s per Watt





PLX characteristics



Model: IBM iDataPlex DX360M3

Architecture: Linux Infiniband Cluster

Processor Type:

- Intel Xeon (Esa-Core Westmere) E5645 2.4 GHz (Compute)
- Intel Xeon (Quad-Core Nehalem) E5530 2.66 GHz (Service and Login)

Number of nodes: 274 Compute + 1 Login + 1 Service + 8 Fat + 6 RVN + 8 Storage + 2 Management

Number of cores: 3288 (Compute)

Number of GPUs: 548 nVIDIA Tesla M2070 + 20 nVIDIA Tesla M2070Q

RAM: 14 TB (48 GB/Compute node + 128GB/Fat node)

PLX system performance

Peak performance: 32 Tflops (3288 cores at 2.40GHz)

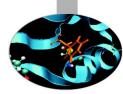
Peak performance: 565 TFlops SP or 283 TFlops DP (548 Nvidia M2070)

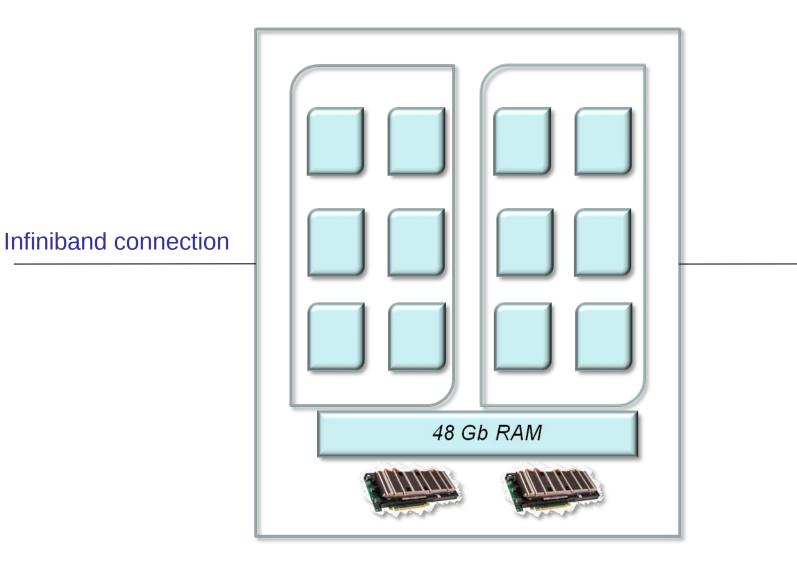






Compute nodes

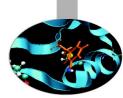








Access credentials



• You will be provided with the credentials (USERNAME a06trnXX) to log into PLX front-end, login.plx.cineca.it: open a shell (or a putty session on Windows) and establish a secure (ssh) connection to PLX front-end:

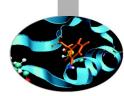
ssh a06trnXX@login.plx.cineca.it

This account will grant you access to PLX from September the 2nd to November the 4th. After such date the account will expire and the password will be reset.





PLX: how to log in



• After having established a ssh connection

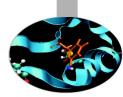
ssh <username>@login.plx.cineca.it

- Please check:
 - important messages that can be found in the message of the day
 - the user guide! <u>http://www.hpc.cineca.it/content/ibm-plx-gpu-user-guide-0</u>





PLX: file transfer

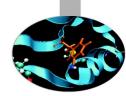


- **sftp / scp** (always available if sshd is running)
- \$ sftp -r <my_dir> <user>@login.plx.cineca.it:/path/to/ \$ scp -r <my_dir> <user>@login.plx.cineca.it:/path/to/
- **rsync**: allows incremental transfer
- \$ rsync -avzr --progress <my_dir> <user>@login.plx.cineca.it:
- **Filezilla**: recommended for Windows users https://filezilla-project.org/





Remark: who uses PLX?



8 nodes are dedicated to the Summer School via the "private" queue

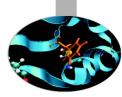
 But: PLX is a resource shared between different type users: academic, industrial, and special agreement users

➔ Please be responsible when you use it: if you crash the login node all the users will be affected





Work Environment



\$HOME:

- Permanent, backed-up, and local to PLX.
- Quota = 4GB.
- For source code or important input files.

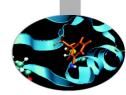
\$CINECA_SCRATCH:

- Large, parallel filesystem (GPFS).
- Temporary (files older than 30 days automatically deleted), no backup.
- No quota. Run your simulations and calculations here.

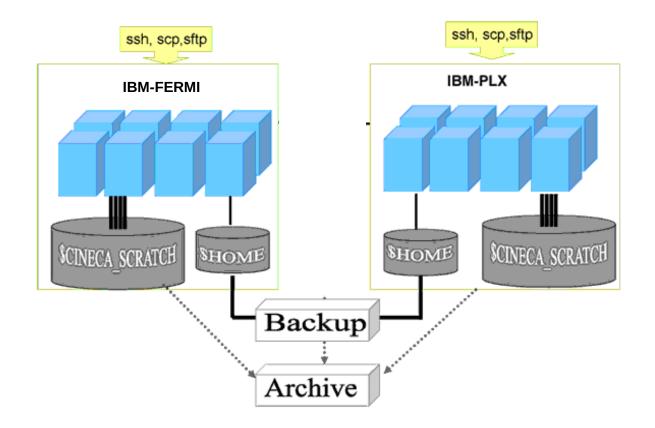




Disks and filesystems



Standard CINECA environment

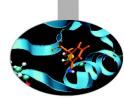


please use "cindata" command to get info on your disk occupation





Accounting: saldo



The accounting mechanism is based on the resources requested for the time of the batch job:

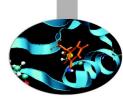
cost = no. of cores **requested** x job duration

In the CINECA system it is possible to have more than 1 budget ("account") The accounts available to your UNIX username can be found from the saldo command.

[a06trn50@node342 ~]\$ saldo -b							
account	start	end	total	localCluster	totConsumed	totConsumed	
			(local h)	Consumed(local h)	(local h)	%	
cin_visual	20120619	20131231	50000	21546	21546	43.1	
train_czss2013	20130523	20130622	20000	1	1	0.0	
train_sc22013	20130902	20131104	20000	0	0	0.0	
						•••	
						CINE	



module, my best friend



• CINECA's work environment is organized in modules, a set of installed libs, tools and applications available for all users.

 all the optional software on the system is made available through the "module" system

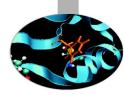
➔ provides a way to rationalize software and its env variables

- modules are divided in 3 *profiles*
 - → profile/base (stable and tested modules)
 - → profile/advanced (software not yet tested or not well optimized)
 - → profile/engineering (for industrial users)
- each profile is divided in 4 categories
 - → compilers (Intel, GNU, PGI, OpenMPI, IntelMPI, CUDA)
 - → libraries (e.g. LAPACK, BLAS, FFTW, ...)
 - → tools (e.g. Totalview, Valgrind, cmake, VNC, python,...)
 - → applications (software for chemistry, physics, ...)





Modules



 "loading" a module means that a series of (useful) shell environment variables wil be set

• e.g. after a module is loaded, an environment variable of the form "<MODULENAME>_HOME" is set

[ibaccare@node342 ~]\$ module load nwchem WARNING: nwchem/6.1.1 cannot be loaded due to missing prereq HINT: the following modules must be loaded first: IntelMPI/4.0--binary [ibaccare@node342 ~]\$ module load IntelMPI/4.0--binary [ibaccare@node342 ~]\$ module load nwchem [ibaccare@node342 ~]\$ echo \$NWCHEM_HOME /cineca/prod/applications/nwchem/6.1.1/IntelMPI_4.0-binary/

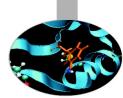
 you can make your life easier by loading the "autoload" module, which will load all the requested modules

[ibaccare@node342 ~]\$ module load autoload nwchem
auto-loading IntelMPI/4.0--binary
[ibaccare@node342 ~]\$ echo \$NWCHEM_HOME
/cineca/prod/applications/nwchem/6.1.1/IntelMPI-4.0-binary/





Module commands



> module available (or just "> module av")

Shows the full list of the modules available in the profile you have loaded, divided by: environment, libraries, compilers, tools, applications

> module (un)load <module_name>

(Un)loads a specific module

> module show <module_name>

Shows the environment variables set by a specific module

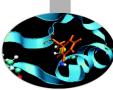
> module help <module_name> Gets all informations on how to use a specific module

> module purge Gets rid of all the loaded modules





Compiling on PLX



On PLX you can choose between three different compiler families: **gnu**, **intel** and **pgi**

You can take a look at the versions available with "*module av*" and then load the module you want. Defaults are: gnu 4.1.2, intel 11.1, pgi 11.1 *module load intel* # loads default intel compilers suite *module load intel/co-2011.6.233--binary* # loads specific compilers suite

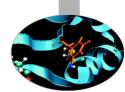
	GNU	INTEL	PGI
Fortran	gfortran	ifort	pgf77
С	gcc	icc	pgcc
C++	g++	ісрс	pgCC

Get a list of the compilers flags with the command *man*





Parallel compiling on PLX



Two families of MPI libraries are available: **openmpi** and **intelmpi**. They provide also the parallel compiler wrappers

There are different versions of openmpi, depending on which compiler has been used for creating them. Default is openmpi/1.4.5--gnu--4.1.2 *module load openmpi* # loads default openmpi compilers suite *module load openmpi/1.4.5--intel--11.1--binary* # loads specific compilers suite

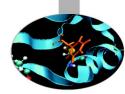
Warning: openmpi needs to be loaded after the corresponding basic compiler suite: you can use "autoload" to load both compilers at the same time

```
[ibaccare@node342 ~]$ module load openmpi
WARNING: openmpi/1.4.5-gnu-4.1.2 cannot be loaded due to missing prereq
HINT: the following modules must be loaded first: gnu/4.1.2
[ibaccare@node342 ~]$ module load autoload openmpi
### auto-loading modules gnu/4.1.2
```

If another type of compiler was previously loaded, you may get a "**conflict error**". Unload the previous module with "module unload"







Parallel compiling on PLX

	OPENMPI INTELMPI
Fortran	mpif90
С	mpicc
C++	mpiCC

Compiler flags are the same of the basic compiler (since they are basically MPI wrappers of those compilers)

\$ mpicc -show # to get the explicit list of arguments

OpenMP is provided with following compiler flags:

gnu: -fopenmp

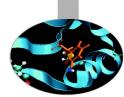
intel : -openmp

pgi: -mp





Static/Dynamic Linking



On PLX you can choose between dynamic and static linking (**dynamic** is the **default)**.

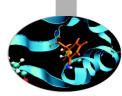
• Static linking means that the library references are resolved at **compile time**, so the necessary functions and variables are already contained in the executable produced. It means a bigger executable but no need for linking the library paths at runtime.

• **Dynamic linking** means that the library references are resolved at **run time**, so the executable searches for them in the paths provided. It means a lighter executable and no need to recompile the program after every library update, but environment variables have to be defined at runtime (i.e. LD_LIBRARY_PATH)

To enable static linking: -static (gnu), -Bstatic (intel, pgi)







Launching jobs

Now that we have our executable, it's time to learn how to prepare a job for its execution

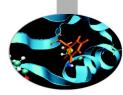
PLX uses PBS scheduler.

The job script scheme is:

#!/bin/bash
#PBS keywords
variables environment
execution line







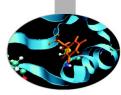
PBS keywords

#PBS -N jobname # name of the job
#PBS -o job.out # output file
#PBS -e job.err # error file
#PBS -l select=1:ncpus=8:mpiprocs=8:mem=24gb # resources
#PBS -l walltime=1:00:00 # hh:mm:ss, max 144 hours in queue "private"
#PBS -q private # chosen queue
#PBS -A <my_account> # name of the account: train_sc22013

select = number of chunk requested
ncpus = number of cpus per chunk requested
mpiprocs = number of mpi tasks per chunk
mem = RAM memory per chunk







PBS Keywords specific for the school

#PBS -A train_sc22013 # your "account" name
#PBS -q private # special queue used in this school

private is a particular queue defined on 8 compute nodes (8 cores each)

train_sc22013 is the cpu-hours budget that you need to set





PLX job script example

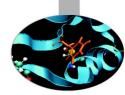
cd \$PBS_O_WORKDIR module load autoload openmpi module load somelibrary

mpirun ./myprogram < myinput





PBS commands



qsub

qsub <job_script>

Your job will be submitted to the PBS scheduler and executed when there will be nodes available (according to your priority and the queue you requested)

qstat

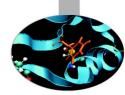
qstat

Shows the list of all your scheduled jobs, along with their status (idle, running, closing, ...) Also, shows you the job id required for other qstat options





PBS commands



qstat

qstat -f <job_id>

Provides a long list of informations for the job requested. In particular, if your job isn't running yet, you'll be notified about its estimated start time or, if you made an error on the job script, you will learn that the job won't ever start

qdel

qdel <job_id>

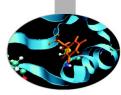
Removes the job from the scheduled jobs by killing it





.

PRIVATE queue



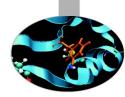
\$ qstat -Qf private

```
Queue: private
  queue type = Execution
  total jobs = 0
  resources_max.ncpus = 64
  resources_max.ngpus = 8
  resources_max.walltime = 144:00:00
  resources_default.ncpus = 8
  resources_default.ngpus = 0
  resources_default.place = free:shared
  acl group enable = True
  acl groups = -,a06trn00,+cin staff,+cinstaff,+corsi,+train scR2013
  enabled = True
  started = True
```





Documentation



Check out the User Guides on our website www.hpc.cineca.it

PLX:

Advanced topic http://www.hpc.cineca.it/sites/default/files/PBSProUserGuide10.0.pdf

