



Production environment on FERMI



Introduction to the FERMI Blue Gene/Q, for users and developers

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PRODUCTION TIME!!

So let's say you have compiled your executable and you want to launch it...

The question is...**HOW TO DO THAT????**

Before you do that, let's take a look at your operational space...

This can be done by writing a small batch script that will be Submitted to a scheduler called **LoadLeveler**





WORK ENVIRONMENT



Once you're logged on FERMI or PLX, you are on your **home** space. It is best suited for **programming** environment (compilation, small debugging sessions...) Space available: 50 GB (FERMI) – 4 GB (PLX) Environment variable: \$HOME

Another space you can access to is your **scratch** space. It is best suited for **production** environment (launch your jobs from there) Space available: UNLIMITED (FERMI) – 32 TB (PLX) Environment variable: \$CINECA_SCRATCH WARNING: On PLX is active a **cleaning procedure**, that deletes your files older than 30 days!

Use the command "cindata" for a quick briefing about your space occupancy







MODULE PROFILES

- > module load <profile_name>
- Available modules ("module av profile"):
 - profile/base (default): it contains the application modules compiled for back-end nodes and ready to be used
 - profile/front-end: it contains the applications modules compiled for front-end nodes and ready to be used
 - profile/advanced. Testing profile. It contains the applicantions modules that have to be tested yet. Usable but not guaranteed





APPLICATION MODULES



>module available (or just "> module av") Shows the full list of the modules available in the profile you're into, divided by: environment, libraries, compilers, tools, applications

Below is the list of the application modules available on profile/base, updated to 17th march, 2013

------ /cineca/prod/modulefiles/base/applications -----abinit/6.12.3 crystal09/1.01 pluto/4.0 amber/12(default) crystal09/2.0.1 (default) qe/5.0bgq bigdft/1.6.0 dl_poly/4.03(default) siesta/3.1 cp2k/2.3(default) gromacs/4.5.5 (default) siesta/3.1-TS cpmd/3.15.3_hfx(default) lammps/20120816 vasp/5.2.12 cpmd/v3.15.3 namd/2.9 vasp/5.3.2





MODULE COMMANDS

> module load <module_name>

Loads a specific module

> module show <module_name>

Shows the environment variables set by a specific module

> module help <module_name>

Gets all informations about how to use a specific module







EXECUTION MODALITIES

- Via command line
 - >./myexe
- On Front-end nodes only

Via batch

 >Ilsubmit job.cmd

 On Front-end and Back-end nodes





EXECUTION: FRONT-END NODES

- Pre and Post processing
- Data transfer
- **Serial** execution (1 core)
- Executables compiled with serial FE compilers >front-end-gnu/4.4.6 >front-end-xl/1.0
- Command line execution (10 min)
- Batch execution (up to 6 h)







BATCH EXECUTABLE: FRONT-END NODES

USER EXECUTABLE

Variables initialization
Execution line
./myexe <options>





BATCH EXECUTABLE: FRONT-END



MODULE EXECUTABLE

 Shell interpreter path #!/bin/bash
 Load Leveler Scheduler Keywords (we'll check them later ;-))

 # @ # @

#@

Variables initialization module load profile/front-end module load <module_name> Execution line myexe <options>







BATCH EXECUTABLE: FRONT-END

LL KEYWORDS

- # @ job_name = serial.\$(jobid)
- # @ output = \$(job_name).out
- # @ error = \$(job_name).err
- # @ wall_clock_limit = 0:10:00 # h:m:s
 execution time up to 6 hours
- # @ class = serial
- # @ queue







EXECUTION: BACK-END NODES

Serial (WARNING: 64 compute nodes are still required) and Parallel execution *Executable compiled with serial and parallel **BE** compilers >bgq-gnu/4.4.6 >bgq-xl/1.0 NO command line execution **Batch** execution (from 64 compute nodes up to 2048 compute nodes, wall clock time up to 24 h) **Runjob** command >runjob <options> >man runjob





BATCH EXECUTABLE: BACK-END NODES



- **Shell** interpreter path
 - #!/bin/bash
- *Load Leveler Scheduler Keywords
 - # @ # @ # @

Variables inizialization
 Execution line
 >runjob <runjob_options> : ./myexe <myexe_options>







BATCH EXECUTABLE: BACK-END NODES

MODULE EXECUTABLE

- **Shell** interpreter path
 - #!/bin/bash

*Load Leveler Scheduler Keywords

- #@
- #@
- #@

Variables inizialization

module load <module_name>

Execution line

>runjob <runjob_options> : \$MODULE_HOME/bin/exe
<exe_options>







LL KEYWORDS

- # @ job_name = check
- # @ output = \$(job_name).\$(jobid).out
- # @ error = \$(job_name).\$(jobid).err
- # @ environment = COPY_ALL #export all variables from your submission shell
- # @ job_type = bluegene
- # @ wall_clock_limit = 10:00:00 #execution time h:m:s, up to 24h
- # @ **bg_size** = 64 # compute nodes number
- # @ notification = always|never|start|complete|error
- # @ notify_user = <email_address>
- # @ account_no = <budget_name> #saldo -b

@ queue

Highlighted are the mandatory keywords, the others are highly suggested







LL KEYWORDS - TOPOLOGY

#@ bg_shape = MD(A)xMD(B)xMD(C)xMD(D)
#midplanes number in the A,B,C,D dimensions
@ bg_rotate = true|false
@ bg_connectivity = torus|mesh|either|
Xa Xb Xc Xd #type of connectivity







LL KEYWORDS - BG_SIZE

- # @ bg_connectivity = Mesh #default
- # @ bg_size = number of compute nodes
- for requests <= 1midplane (512 compute nodes)
 bg_size = 64| 128| 256| 512</pre>
- for requests > 1midplane

 bg_size = (512)X2 | (512)X3 | (512)X4 |

 (512)X5 | (512)X6 | (512)X8 | (512)X10 | (512)X12 |

 (512)X16







EXECUTION LINE

Your executable is launched on the compute nodes via the command "runjob", that you can set in two ways:

- 1) Use ":" and provide executable infos how you're used to runjob : ./exe_name arg_1 arg_2
- 2) Use specific runjob flags
 --exe Path name for the executable to run runjob --exe ./exe_name
 - --args Arguments for the executable specified by --exe runjob --exe ./exe_name --args arg_1 --args arg_2





EXECUTION LINE: MPI TASKS SETTING

- --ranks-per-node (-p) Number of ranks (MPI task) per compute node. Valid values are 1, 2, 4, 8, 16, 32 and 64 (default=depending on the tasks requested) bg_size = 64
- runjob --ranks-per-node 1 : ./exe <options> #64 nodes used, 1 task per node runjob --ranks-per-node 4 : ./exe <options> #64 nodes used, 4 tasks per node
- --np (-n) Number of ranks (MPI task) in the entire job (default=max) bg_size = 64
- runjob --np 64 -- ranks-per-node 1: ./exe <options> #64 tasks, 1 per node runjob --np 256 -- ranks-per-node 4: ./exe <options> #256 tasks, 4 per node runjob --np 200 -- ranks-per-node 4: ./exe <options> #200 tasks, 4 per node until all tasks are allocated
- runjob --np 1 --ranks-per-node 1: ./exe <options> # serial job

Formula: np <= bg_size*ranks-per-node









- --envs Sets the environment variables for exporting them on the compute nodes
 - #MPI/OpenMP job (16 threads for each MPI task)
 runjob -n 64 --ranks-per-node 1 --envs OMP_NUM_THREADS = 16 : ./exe
- --exp-env Exports an environment variable from the current environment to the job
 - export OMP_NUM_THREADS = 16 runjob -n 64 --ranks-per-node 1 --exp-env OMP_NUM_THREADS : ./exe





LOADLEVELER COMMANDS



Your job script is ready! How to launch it?

llsubmit

Ilsubmit <job_script>

Your job will be submitted to the LL scheduler and executed when there will be nodes available (according to your priority)

llq

llq -u \$USER

Shows the list of all your scheduled jobs, along with their status (idle, running, closing,...)

Also, shows you the job id required for other llq options

IIq -s <job_id>

Provides information on why a selected list of jobs remain in the

NotQueued, Idle, or Deferred state.





LOADLEVELER COMMANDS



Ilq -l <job_id>
Provides a long list of informations for the job requested.
In particular you'll be notified about the bgsize you requested and the
real bgsize allocated:

BG Size Requested: 1024 BG Size Allocated: 1024 BG Shape Requested: BG Shape Allocated: 1x1x1x2 BG Connectivity Requested: Mesh BG Connectivity Allocated: Torus Torus Torus Torus

llcancel

llcancel <job_id>

Removes the job from the scheduler, killing it





JOB CLASSES



The class you're going into depends on the resources you asked:

debug: bg_size=64, wall_clock_time <= 00:30:00

longdebug: bg_size=64, wall_clock_time > 00:30:00 (up to 24h)

parallel: bg_size>64 (valid values: 128,256,512,1024,2048)

There are some classes that you can specify:

special: bg_size>64 (up to 512), @ class = special this class allows you to run in 16 I/O nodes racks with bigger jobs

keyproject: for bigger jobs (> 2048 nodes). You have to be an authorized user (write to **superc@cineca.it**)



MODULE «SUPERC»



>module load superc

jobtyp (provides useful information about job in the LL queues - user, tasks, times, ...)

- **For using**
- > jobtyp <job_id>

sstat/sstat2 (provides useful information about the system status - jobs in the LL queues, allocated nodes, Midplane status,...

- For using
- > sstat
- > sstat2

bgtop (draws a full-terminal display of nodeboards and jobs) >bgtop

loadHPC (calculates aggregate statistics of LL jobs) >loadHPC





ACCOUNTING

As an user, you have access to a limited number of CPU hours to spend. They are not assigned to users, but to **projects** and are shared between the users who are working on the same project (i.e. your research partners). Such projects are called **accounts** and are a different concept from your username.

You can check the state of your account with the command "saldo –b", which tells you how many CPU hours you have already consumed for each account you're assigned at

(a more detailed report is provided by "saldo –r").

[amarani0@fen08 ~]\$ saldo -b start total localCluster totConsumed totConsumed account end Consumed(local h) (local h) (local h) 30365762 cin staff 20110323 20200323 1000000000 30527993 3.1 cin totview 20130123 20130213 50000 0.0 0 0 train sc32013 87458 20130211 20130411 1250000 87458 7.0 train cnl12013 20130311 20130411 100000 0 0 0.0



SMT



It is possible to improve the efficiency of every single CPU by activating Simultaneous Multi Threading (SMT)

Each CPU is divided into threads that act as separated tasks, sharing the CPU resources to work simultaneously (with some loss because of latency)

On FERMI, you can activate 2 or 4 simultaneous threads per CPU, meaning for example that you can launch a job with 2048 or 4096 tasks asking only for 1024 cores!

This is achieved by asking for ranks-per-node = 32 (2*16) or ranks-per-node = 64 (4*16)





SUB-BLOCKING





There is, however, a technique that allows to launch multiple executables on a single 64 nodes allocation, partitioning it in sub-groups of nodes called **sub-blocks**

With sub-blocking, you can get advantage of the full number of resources you have to allocate, even with smaller or not very scalable applications. Nothing is wasted!





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HOW TO USE SUB-BLOCKING



Some environment variables have to be set for sub-blocking usage:

> module load subblock

You can find a complete jobscript in our LL User Guide (link at final slide)

Jobscript USER SECTION: export N_SUBBLOCK=4 ### No. of sub-block you're asking (2,4,8,16,32,64) export RANK_PER_NODE=16 ### No. of MPI tasks in each node. ### module load <your applications> export WDR=\$PWD export EXE_1=\$WDR/executable_1.exe export EXE_2=\$WDR/executable_2.exe

export EXECUTABLES="\$EXE_1 \$EXE_2 \$EXE_3 \$EXE_4"





USEFUL DOCUMENTATION



FERMI USER GUIDE:

http://www.hpc.cineca.it/content/ibm-fermi-user-guide http://www.hpc.cineca.it/content/batch-scheduler-loadleveler-0

Job command file keyword descriptions IBM

http://publib.boulder.ibm.com/infocenter/clresctr/vxrx/index.jsp? topic=/com.ibm.cluster.loadl.v5r1.load100.doc/am2ug_sbmbgjbs.htm

