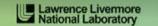
Building and running NPB-MZ-MPI on Leftraru

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What is the NPB-BT-MZ-MPI?

- •A benchmark from the NAS parallel benchmarks suite
- MPI version
- Implementation in Fortran
- Solves multiple, independent systems of block tridiagonal (BT) equations
- Represents workloads similar to many flow solver codes (3D Navier-Stokes equations)
- Probably not much unused optimization potential

We will use this application in most exercises during this workshop.



Properties of NPB-BT-MZ-MPI

- •The solution is done for multiple zones (MZ), in a repeated time-step loop
 - •After each time-step, the zones have to exchange boundary values
 - •Fine-grained parallelism within a zone
 - Coarse-grained parallelism between zones
 - Zones are not all equally sized and need to be distributed in a balanced way
- A larger problem size adds more zones
- Exploits multi-level parallelism
 - -Hybrid (OpenMP + MPI) implementation

Suitable testing application for a wide range of tools and analysis types!



First step: Use Intel tools

Add to your .bashrc: module load intel impi

Re-login to apply changes.



Second step: Building the benchmark

•Copy to your home:

```
$ cp -r /home/courses/instructor06/NPB3.3-MZ-MPI ~
This is the original source code, with no modifications.
```

•Create default config/make.def:

```
$ cd NPB3.3-MZ-MPI
$ cp config/make.def.template config/make.def
```

•Edit config/make.def to use mpiifort as compiler:

```
F77 = mpiifort
```

■ Issue make command (typing only make will give you a help text):

```
$ make bt-mz CLASS=B NPROCS=32
```



Third step: Run the application

```
Change to bin and copy job script from
/home/courses/instructor06/jobscript.slurm:
#!/bin/bash
#SBATCH -- job-name=example
#SBATCH --partition=slims
#SBATCH --exclusive=user
#SBATCH --ntasks per socket=8
#SBATCH -n 32
#SBATCH --output=out.txt
#SBATCH --error=out.txt
. /etc/profile
. /etc/profile.d/modules.sh
srun ./bt-mz.B.32
Run the job with sbatch <your_script>
```



Useful commands

Check your personal job queue: squeue

Cancel a job:
scancel < job id>

Print contents of output file:
cat out.txt

Follow the output, while job is running:
tail -F out.txt

Done!

You have successfully built and run the benchmark.





