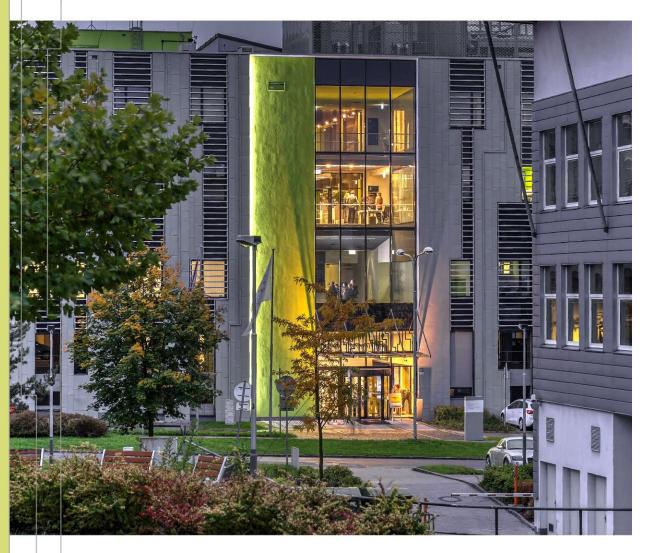
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INTRODUCTION TO HIGH PERFORMANCE COMPUTING

PART 3 HPC @ IT4INNOVATIONS BUILDING CODE ON THE CLUSTER

Jakub Beránek



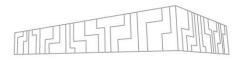
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OUTLINE

- Remote development
- Modules
- Available software on clusters
- Common toolchains (C/C++, Python)
- Containerization
- Gitlab, Cl

LOCAL VS REMOTE DEVELOPMENT



- Local code development is much easier
- **Remote IDE** VSCode/Clion/... offer remote development over SSH
 - The server compiles code on the cluster, but the UI runs on your PC
- sshfs treat the remote filesystem as a local one
- **git** synchronize code through a repository

BUILDING CODE AND DEPENDENCIES



- You must build your program and its **dependencies** for your target cluster
 - e.g. Karolina runs on Rocky Linux 8
- You DO NOT have admin privileges on the cluster
 - Standard package managers (like yum/apt) cannot be used

You have several options how to compile your code and its dependencies

- Use the available pre-installed modules
- Compile your code and all its dependencies from scratch
- Use <u>Apptainer containers</u>
- Use <u>Spack</u> (HPC package manager)

• .

All further command examples will assume execution on a login node

You can find more information <u>here</u>

USING LMOD MODULES

- Each IT4I cluster has its own set of pre-installed <u>modules</u> available for immediate use
- Module
 - Is a set of binaries, libraries, header files, ...
 - Has a set of modules that it depends on
 - Might have several available <u>versions</u> (Python/2.7.9 vs Python/3.6.1)
 - Might have a specific toolchain (GCC vs Intel toolchain)
- To use a module, you have to load it
 - Loading a module modifies environment variables (PATH, LD_LIBRARY_PATH)
 - This enables executing module binaries and linking to module libraries
- <u>Lmod</u> is used to load modules
- You can also create your <u>own modules</u> or <u>ask support</u> to install new modules for you
 - Modules are defined using EasyBuild
- If you find a module that is not working, contact support

AVAILABLE MODULES



- Language toolchains (Python, Java, C#, ...)
- C/C++ compilers (GCC, Clang, Intel C++ compiler, CUDA nvcc, ...)
- Communication libraries (MPI, GPI-2, ...)
- Parallel debuggers and profilers (Allinea Forge, VTune, PAPI, Scalasca, Score-P, Vampir, ...)
- Parallelized libraries (FFTW, PETSc, Trilinos, Octave, ...)
- Specialized software for chemistry, bioinformatics, physics, visualization, 3D rendering, ...
 - GROMACS, Gaussian, Molpro, NWChem, Orca, Phono3py, OpenFOAM, ParaView, ...

USING LMOD



show available modules
\$ ml av

load a module with its dependencies
\$ module load Python/3.6.8

list loaded modules

\$ module list Currently loaded modules: 1) GCC/6.3.0 2) Python/3.6.8 \$ python --version Python 3.6.8

unload all loaded modules
\$ ml purge
\$ python --version
Python 2.7.5

Useful hints

Always load specific versions of modules to avoid surprises

Х

- ml GCC/6.3.0
- ml GCC
- Module load order matters (because of conflicting dependencies)
 - ml A B might produce different results than ml B A
- Save module combinations that you commonly use into *collections*

\$ ml purge

- \$ ml GCC/6.3.0 Python/3.6.8 MPICH/3.2.1-GCC-6.3.0-2.27
- \$ ml save mpienv1 # save current modules under name mpienv1
 # _______
- \$ ml restore mpienv1 # restore modules from collection mpienv1
- Filtering modules
 - \$ ml spider <package>
 - ml command also provides tab completion
- ml command is case sensitive
- Match module toolchains (GCC vs Intel)
- Do not forget to load correct modules in your Slurm job script!

USING GPUS

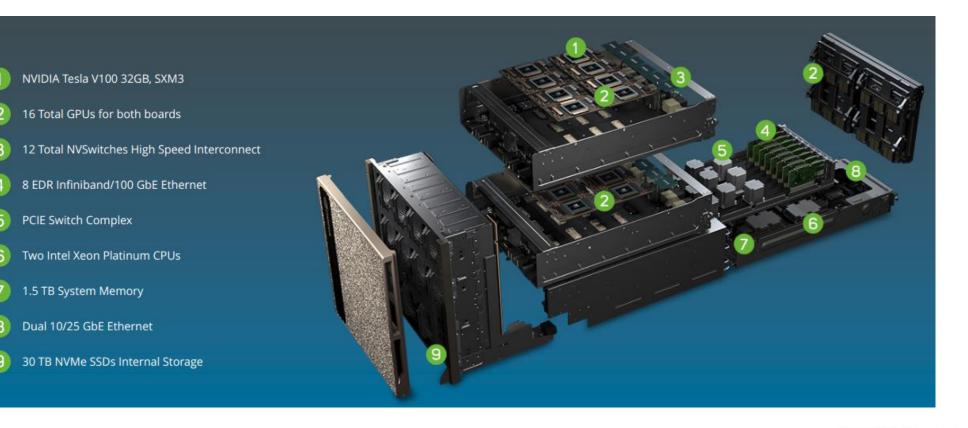
- Available clusters with GPUs:
 - Karolina: 72 nodes, 8 A100 (40 GiB) GPUs per node
 - Barbora: 8 nodes, 4 V100 (16 GiB) GPUs per node
 - DGX: 16 V100 (32 GiB) GPUs
- By default, Karolina jobs will allocate a single GPU
 - Check if your tool has support for Multi-GPU setups
- Use prepared modules:
 - \$ ml CUDA/12.2.0
 - \$ ml av nvhpc
- When loading multiple GPU modules, match their versions!
 - \$ ml CUDA/12.2.0 cuDNN/8.9.2.26-CUDA-12.2.0





DGX-2

- Has a dedicated PBS queue
 - Accessible from Barbora (qdgx queue)
- Check if your tool has direct support for it
 - Dask-DGX



COMPILING C/C++ PROGRAMS



- 1. Load necessary modules
 - Compiler (e.g. GCC/6.3.0)
 - Dependencies (e.g. MPICH/3.2.1-GCC-6.3.0-2.27)
 - Build system (e.g. CMake/3.16.2)
- 2. Build your program on a login node
 - Once your binary is built, it can be accessed by all cluster nodes using the shared filesystem
- 3. Adjust your PATH/LD_LIBRARY_PATH environment variables
 - PATH directories where binaries are located
 - LD_LIBRARY_PATH directories where shared dynamic libraries are located

C/C++ COMPILATION FLAGS AND TIPS (GCC)

- Make use of optimizations and available instruction sets
 - Karolina has AVX2 (256-bit vectorization)
 - Barbora has AVX-512 (512-bit vectorization)
- Useful flags
 - <u>Optimizations</u>: -02, -03
 - Benchmark what works best for your code
 - <u>Use native instruction set</u>: -march=native
 - Fast floating point math (at the cost of precision): -ffast-math
 - Link-time optimization: -flto
 - Profile-guided optimization: -fprofile-generate, -fprofile-use
 - Enable OpenMP: -fopenmp
- Tip: you can check generated assembly at godbolt.org

COMMON C/C++ BUILD SYSTEMS



• Makefile

- Simply run make in the project directory
- CMake
 - 1. Load CMake module
 - 2. Create build files inside a build directory
 - 3. Invoke Make (or other build systém, e.g. Ninja) to build the project
- CMake tip: use -DCMAKE_INSTALL_PREFIX=<dir> so that you can use make install

```
$ ml CMake/3.13.1
$ mkdir build
$ cd build
$ cmake -DCMAKE_BUILD_TYPE=Release ..
$ make -j16
```



MPI



- Choose desired MPI implementation and module
 - MPICH, OpenMPI, Intel MPI (impi)
 - Keep the same impl. and version for compilation and execution
- Compile using mpicc or mpicxx
- Run your program
 - \$ mpirun -n 2 <program>
- More information about MPI in a later section

PYTHON



- Works mostly out-of-the-box on all clusters
- Make sure to load the same Python version module
 - When setting up your environment
 - Inside Slurm jobs
- Avoid using system/user Python, use virtual environments instead
 - Puts all your dependencies inside a single directory
 - venv usage example

\$ python3 -m venv venv \$ source venv/bin/activate (venv) \$ pip install -U pip setuptools wheel (venv) \$ pip install <my-package>



PYTHON (PERFORMANCE)

IAI SUPERCOMPLITING

- Many useful cluster/HPC frameworks exist
 - Parallelize computation or put it on GPU with a few lines of codes
 - Distributing computation: <u>Dask</u>, <u>Ray</u>, <u>PySpark</u>, <u>HyperLoom</u>
 - GPU-acceleration: <u>RAPIDS</u> (cuDF, cupy), <u>numba</u>

import dask.dataframe as dd
df = dd.read_csv('2015-*-*.csv')
df.groupby(df.user_id).value.mean().compute()



- Python compute bound programs can be accelerated by <u>PyPy</u> or <u>Cython</u>
- Profile performance using <u>py-spy</u> or <u>Scalene</u>

CONTAINERIZATION USING APPTAINER

- Containers allow you to
 - Prepare your code and all dependencies
 - Distribute them easily in the form of an archive (image)
 - Execute them in a sandboxed environment
- Popular container solution is Docker
 - It cannot be used on IT4I clusters directly because of security issues
- You can use Apptainer instead
 - Preferred deployment method on DGX-2
 - Nvidia containers available at <u>NGC</u>

\$ ml apptainer \$ apptainer shell docker://centos:latest \$ apptainer build ubuntu.img docker://ubuntu:latest \$ apptainer shell -B /scratch ubuntu.img



GITLAB



- IT4I hosts a GitLab instance at https://code.it4i.cz
- Code storage, sharing and review (repositories, pull requests)

Vojtech Cima > mloc > Details

- Project management (issue tracker, wiki)
- Container repository
- Continuous integration

M Project ID: 494	ches 🖉 0 Tags 🗈 1.2 MB Files 🗔 1.2 MB	△ ~ ★ Star 0 ¥ Fork 0
naster ~	mloc / + •	History Find file Web IDE
ENH: Updated de Vojtech Cima aut	apendencies hored 1 month ago	7147ae53 🖞
🖻 README 🛛 🕂 MIT L		NTRIBUTING Set up CI/CD
Name	Last commit	Last updat
docs	ENH: Added MSR logo	2 months ag
🖨 examples	ENH: Added PBS backend	3 months ag
🖿 mloc	ENH: Updated dependencies	1 month ag
tests	[test] Added test for convolutiona	al neural net 2 years ag
🚸 .gitignore	First draft	2 years ag
🗼 Dockerfile	ENH: Updated dependencies	1 month ag
LICENSE	Added license	2 years ag
M# README.md	ENH: Added MSR logo	2 months ag
🔒 requirements.txt	ENH: Updated dependencies	1 month ag

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GITLAB CI (CONTINUOUS INTEGRATION)

- Pipelines = scripts executed after a push to a repository
 - IT4I has 5 shared runners that can run pipelines
- Check that your code was not broken by a commit
 - Correctness (unit tests)
 - Performance (benchmarks)
 - Code style, lints, ...
- Deploy built artifacts
- Configured with .gitlab-ci.yml

Status	Pipeline	Triggerer	Commit	Stages	
⊘ passed	#12563 latest	-	P routing_lib ↔ 4eeld985 ENH: PTDR base for wrapper	\odot	රී 00:08:02 ё 4 months ago
⊘ passed	#12544 latest	٩	₽ master - 848f9c28 ② Update .gitlab-ci.yml	\odot	ੴ 00:08:07 ё 4 months ago
⊘ passed	#12492	۹	₽ settings_Di 848f9c28 Opdate .gitlab-ci.yml	\odot	ੴ 00:07:57 Ё 4 months ago
(★) failed	#12491	۹	₽ settings_Di → 508be6f0 @ Update .gitlab-ci.yml	× -»	♂ 00:00:52 Ё 4 months ago

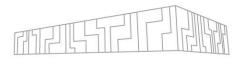
1	<pre>image: 'rust:latest'</pre>			
2	-			
3	stages:			
4	- build			
5	- test			
6				
7	variables:			
8	CARGO_HOME: \$CI_PROJECT_DIR/cargo			
9	RUSTUP_TOOLCHAIN: stable			
10				
11	before_script:			
12	- apt-get update -yq			
13	 apt-get installno-install-recommends -y libzmq3-dev 			
14				
15	build:routing:			
16	stage: build			
17	artifacts:			
18	paths:			
19	- build/			
20	expire_in: 6h			
21	script:			
22	- mkdir build			
23	- cd build			
24	- cmake -DCMAKE_BUILD_TYPE=Release			
25	- make -j4			

🗈 .gitlab-ci.yml 🖺 798 Bytes

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Gitlab CI documentation can be found <u>here</u>

FURTHER READING



- <u>Productivity tools</u> workshop
 - Git
 - EasyBuild
 - Gitlab Cl
 - Singularity
 - Lmod
 - kvm





Jakub Beránek jakub.beranek@vsb.cz

IT4Innovations National Supercomputing Center VSB – Technical University of Ostrava Studentská 6231/1B 708 00 Ostrava-Poruba, Czech Republic www.it4i.cz VSB TECHNICAL | IT4INNOVATIONS ||||| UNIVERSITY | NATIONAL SUPERCOMPUTING OF OSTRAVA | CENTER



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