

HPC Software – Modules, Libraries & Software JSC Training Course

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#### Outline

- 1 Modular Programming
- 2 Preinstalled HPC Software Packages at JSC
  - Navigating Modules
  - Mathematical Libraries
- 3 User Installations at JSC
- 4 Containers
- 5 Further Information



Software Implementation & Libraries

A **library** is a collection of resources.

In computer science: configuration data, documentation, help data, message templates, pre-written code and subroutines, classes, values or type specifications

- Don't reinvent the wheel: Recoding is time-consuming and error-prone
- Best practice for own code: Design interfaces and do different implementation separately (ideally separate files)



Compiling & Linking

#### Example:

 Code decomposition into an executable and an outsourced subprogram (C++: Link object files for different modules)

> g++ -g -c -o main.o main.cpp g++ -g -c -o alibrary.o alibrary.cpp g++ -g main.o alibrary.o -o main

(In real life: Use make or cmake)

- Copy .o and .h to separate directories (e.g. /base/lib/alibrary.o) and add
  - The -L option for the path to the library's object, the object code using -l
    - g++ -g -L/base/lib -o main main.o -lalibrary
  - The -I flag for include directories



Compiling & Linking

- For libraries that are not in standard directories, you need to set
   -I/base/include and -L/base/lib for compiling/linking
- Or define linux environment variables:
  - You either enter the *export*-commands on the linux prompt before compiling, or, copy them into to the .bashrc file in your home folder
- Usually you do not need -I or -L for libraries accessed using the module load command on our supercomputers



Installing Libraries from Source

- Library is not in the software module stack
- Common installation procedure:
  - make
    - ./configure -prefix=base make; make install
  - cmake

cmake -DCMAKE\_INSTALL\_PREFIX=base make; make install

Choose base as a directory you have write permissions





# Preinstalled HPC Software Packages EasyBuild



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**Navigating Modules** 

- Tools are available through "modules"
  - Allows to easily manage different versions of programs
  - Works by dynamic modification of a user's environment
- Module setup based on EasyBuild and Imod
  - "Staged", hierarchical setup
  - Automatically manages dependencies via "toolchains"
- Consistent setup on JURECA, JUSUF and JUWELS (Cluster & Booster)



**Toolchains: Underlying Compiler and MPI Libraries** 

Base: GCCcore

#### Compiler:

- Intel compiler
- GNU compiler
- NVIDIA (CUDA) compiler

#### MPI libraries:

- ParastationMPI
- Intel MPI
- OpenMPI

#### • Math libraries: e.g. FlexiBLAS



Toolchains: Underlying Compiler and MPI Libraries

Current software stage is 2024

- Base: GCCcore (12.3.0)
- Compiler:
  - Intel compiler (Intel 2023.2.1)
  - GNU compiler (GCC 12.3.0)
  - NVIDIA (CUDA) compiler (NVHPC 23.7)
- MPI libraries:
  - ParastationMPI (ParaStationMPI 5.9.2-1)
  - Intel MPI (Intel MPI 2021.10.0)
  - OpenMPI (OpenMPI 4.1.5)
- Math libraries: e.g. FlexiBLAS (MKL 2023.2.0, BLIS 0.9.0, OpenBLAS 0.3.23)



**Modules Environment** 

Available Compiler/MPI Combinations in Stages/2024

Compiler	MPI	Cuda available
GCC	ParaStationMPI	yes
GCC	OpenMPI	yes
NVHPC	ParaStationMPI	yes
NVHPC	OpenMPI	yes
Intel	ParaStationMPI	yes
Intel	IntelMPI	no
Intel	OpenMPI	yes



**Toolchains: Underlying Compiler and MPI Libraries** 

The most important module command

- module load <name> or ml <name>
- GCCcore is preloaded, which enables a lot of base software
- For HPC software you have to load a **compiler**, to expand the module tree

e.g. ml GCC (Default is 12.3.0)

- Then you load an MPI version
   e.g. ml ParaStationMPI (Default is 5.9.2-1)
- Then you can load any other math or application package e.g. ml PETSc/3.20.0



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#### **Modules Environment**

- ml spider name shows whether a library is available in the current stage and in which versions
   e.g. module spider petsc
   ⇒ PETSc/3.18.5, PETSc/3.20.0, ...
- ml spider name/version shows which environment you have to load before you can load that version
   e.g. module spider PETSc/3.20.0
   ⇒ Stages/2024 + GCC/12.3.0 + ParaStationMPI/5.9.2-1, ...
- Some packages are hidden. To see them use

ml -show-hidden spider name



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#### **Modules Environment**

- After loading compiler and MPI <u>ml avail</u> shows the software available with that combination
- ml avail name and ml help name will show you details about the name package
- Many libraries are available for more than one combination/toolchain
- Write e-mail to sc@fz-juelich.de if you want special versions or new software
  - No guarantee the software will be installed
  - alternative: Install packages in User space



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**Modules Environment** 

- For R, Python and Perl we use bundles
  - You might be looking for a software package that is part of a bigger module
- Use ml key software

ml key numpy will suggest SciPy-Stack

• You can use then ml spider to find out how to load the module



**Modules Environment** 

Stages

- The whole software stack of JURECA, JUSUF, JUWELS Cluster and Booster will be updated regularly
- Current stage is 2024
- Old stages are still accessible on JUWELS Cluster, but no guarantee
- To check availability in other stages first type ml use \$OTHERSTAGES



Mathematical Libraries: FlexiBlas

FlexiBlas: wrapper library, includes MKL, BLIS, OpenBLAS

- Linear Algebra Packages (LAPACK, ScaLAPACK, ...)
- Iterative Sparse Solvers, Trust Region Solver
- Vector Math Library
- Vector Statistical Library
- Fourier Transform Functions
- Trigonometric Transform Functions
- GMP routines, Poisson Library, ...



Mathematical Libraries: Sequential Packages

#### **Public domain Libraries**

- LAPACK (Linear Algebra PACKage)
- ARPACK (ARnoldi PACKage)
- GSL (Gnu Scientific Library)
- GMP (Gnu Multiple Precision Arithmetic Library)
- METIS (Serial Graph Partitioning and Fill-reducing Matrix Ordering)



Mathematical Libraries: Parallel Packages

- ScaLAPACK (Scalable Linear Algebra PACKage)
- ELPA (Eigenvalue SoLvers for Petaflop-Applications)
- FFTW (Fastest Fourier Transform of the West)
- MUMPS (MUltifrontal Massively Parallel sparse direct Solver)
- ParMETIS (Parallel Graph Partitioning)
- SCOTCH (Parallel Graph Partitioning)
- Hypre (high performance preconditioners)
- ARPACK (Parallel ARPACK)
- SPRNG (Scalable Parallel Random Number Generator)
- SUNDIALS (SUite of Nonlinear and DIfferential/ALgebraic equation Solvers)



# **GPU Library**

MAGMA, Matrix Algebra on GPU and Multicore Architectures

#### Parallel Systems

- PETSc, toolkit for partial differential equations
  - PETSc for Python (petsc4py)
- SLEPc Scalable Library for Eigenvalue Problem Computations Extension to PETSc for the computation of eigenvalues and eigenvectors



#### Licensed software

Matrix Laboratory

#### MATLAB for High-Performance Computing

- Licenses provided for non-commercial academic research or teaching
- Access via JuDoor
- www.fz-juelich.de/en/ias/jsc/services/ user-support/software-tools/matlab





# User Installations at JSC EasyBuild



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# EasyBuild

How it is used at JSC

- Used by the software team to create software stacks since 2014
  - Install in Production stage:
    - ml Stages/2024 ml Developers eb packages-1.2.3.eb

Allow users to install software on-top of the available modules

Install in User space:

ml Stages/2024 export USERINSTALLATIONS=/p/project/yourproject/user ml UserInstallations eb packages-1.2.3.eb



# Excerpt from PETSc easyconfig

https://github.com/easybuilders/JSC/blob/2024/Golden\_Repo/p/PETSc/ PETSc-3.20.0-gpsfbf-2023a.eb

04 name = "PETSc"
05 version = "3.20.0" # PETSc/3.20.0

#see Golden\_Repo: g=GCC, ps=ParaStationMPI, fbf=FlexiBlas
18 toolchain = 'name': 'gpsfbf', 'version': '2023a'

# where the installation files are downloaded
23 source\_urls = ['https://web.cels.anl.gov/...
24 sources = ['petsc-%s.tar.gz' % version]

#deps that are already present at stage 2023a <sup>35</sup> dependencies = [('METIS', '5.1.0'), ...

<sup>53</sup> configopts = '-with-large-file-io'



## Create new PETSc installation

PETSc-3.20.0-gpsfbf-2023a-myVersion.eb

- <sup>04</sup> name = "PETSc"
- 05 version = "3.20.0"
- <sup>06</sup> versionsuffix = '-myVersion' # PETSc/3.20.0-myVersion
- <sup>18</sup> toolchain = 'name': 'gpsfbf', 'version': '2023a'

```
# where the installation files are downloaded
23 source_urls = ['https://web.cels.anl.gov/...
24 sources = ['petsc-%s.tar.gz' % version]
```

```
#deps that are already present at stage 2023a
35 dependencies = [('METIS', '5.1.0'), ...
```

53 configopts = '-with-large-file-io'
54 configopts += '-something\_fancy '



### Create and run new PETSc installation

PETSc-3.20.0-gpsfbf-2023a-myVersion.eb

#### Install in User space:

ml Stages/2024 export USERINSTALLATIONS=/p/project/yourproject/user ml UserInstallations eb PETSc-3.20.0-gpsfbf-2023a-myVersion.eb

#### Load and Use of new Software:

export USERINSTALLATIONS=/p/project/yourproject/user ml Stages/2024 && ml GCC && ml ParaStationMPI ml PETSc/3.20.0-myVersion



# EasyBuild: Further Links

#### **Documentation:**

- EasyBuild Documentation https://docs.easybuild.io
- EasyBuild Tutorial https://easybuild.io/tutorial

#### Where do I find easyconfigs?

- JSC repository https://gitlab.jsc.fz-juelich.de/software-team/easybuild
- JSC public mirror https://github.com/easybuilders/JSC
- Upstream https://github.com/easybuilders/easybuild-easyconfigs





# Containers Apptainer



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### Containers

What they provide

- Containers package up pieces of software in a way that is portable and reproducible, they ...
  - manage different versions of programs
  - are more lightweight than virtual machines
  - provide the ability to build, ship, and run applications
- Some examples are Docker, Shifter, and Apptainer/Singularity
- They typically use so-called "images"
  - contain a file system including a minimal operating-system, the application, and some metadata
- We provide an up-to-date version of Apptainer
  - Formerly, we provided Singularity on the Systems
  - We have replaced Singularity by Apptainer, a fork maintained by the linux foundation
- To be granted access to the container runtime, you have to go to our user portal **JuDoor**



# **Apptainer Containers**

First steps

#### Download an image:

• Use the pull command to download pre-built images from an external resource like Docker Hub

apptainer pull centos.sif docker://centos:7

#### Call an executable:

 The shell command allows you to spawn a new shell within your container and interact with it

srun -N1 -p <part> -gres gpu:1 -pty apptainer shell -nv centos.sif

• To Slurm, Singularity is just another executable and can be called as such



## **Apptainer Containers**

First steps

#### **Container Build System:**

- JSC provides a build system that can build images on behalf of the user, based on a Docker- or Singularity-file
- For further information see https://apps.fz-juelich.de/jsc/hps/jureca/ container-runtime.html



#### Further information and support

http://www.fz-juelich.de/ias/jsc/jureca
http://www.fz-juelich.de/ias/jsc/juwels
http://www.fz-juelich.de/ias/jsc/jusuf

Supercomputer support: sc@fz-juelich.de

