

# Scientific Software Management in Real Life Deployment of EasyBuild on a Large Scale System

HUST '16, November 13, 2016 | D. Alvarez\*<sup>1</sup> A. O'Cais<sup>1</sup> M. Geimer<sup>1</sup> K. Hoste<sup>2</sup> |

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# Scientific Software Management in Real Life

## Part I: Introduction

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## Managing scientific software

- HPC systems typically used by different kind of users.
- Very different software requirements
  - Different compilers
  - Different libraries
    - Different versions of these libraries
  - Different levels of HPC expertise
  - Different tools
- Different time plans

## Managing scientific software

- Burden for system administrators and user support teams.
  - May lead to relying on OS packages
    - ⇒ Can only be updated during a maintenance window
    - ⇒ Limited to the OS available packages
    - ⇒ Increased size of OS images

### OS packages examples:

- Software for general programming
    - Subversion, git, CMake, ...
  - Software to support components of the scientific software stack
    - X11, additional Python modules, ...
- 
- How to deal with different versions?
  - How to keep them reasonably up to date?

## Scientific software from a user view

- Software often provided via environment modules.
  - Shell-independent way to modify a user's environment
  - Can be organized in various ways (flat, hierarchical, ...)
    - Though sometimes difficult to implement
- Creating and maintaining consistent module views is tedious and error-prone.

### Solution:

Various tools exist to help with software installations and automatic module file creation.

## EasyBuild & Lmod @ JURECA

- EasyBuild is a software installation framework
  - <http://hpcugent.github.io/easybuild/>
  - Already provides lots of useful functionality
  - Compartmentalized structure: framework, easyblocks, easyconfigs
  - Some features we require *were* still missing
- Lmod is a modern environment modules tools
  - <https://github.com/TACC/Lmod>
  - Powerful support for hierarchy of modules
- How do we use and *extend* these tools to support our users effectively and efficiently?

# Scientific Software Management in Real Life Part II: System details & requirements

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## JURECA system characteristics

- 1.8 + 0.44 PFlops, #57 in Top500 (June'16)
- 1872 compute nodes (Haswell)
- 75 compute nodes with NVIDIA K80 GPUs
- 12 visualization nodes, each with two NVIDIA K40 GPUs
- Mellanox EDR InfiniBand with fat tree topology
- Any guess on user requirements?



## User requirements

“I want it all, I want it all, I want it all, and I want it now” 🎵🎵🎵

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- And of course:
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  - Compatibility
  - A simple user view



# Scientific Software Management in Real Life Part III: Designing the User View

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    - Have to choose particular compiler and MPI combinations before seeing any other package
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    - Visible software is compatible
  - Hierarchy of compilers and MPI runtimes
    - Modules available are shown in a staged fashion
    - Intuitive
    - Visible software is compatible



## Designing the User View: Lmod as modules tool

- Lmod was designed with module hierarchies in mind
  - `module spider` and `module key`
  - Module families (`family("compiler")` or `family("mpi")`)
- Lmod also has other interesting features
  - Good support for hidden modules (`--show-hidden`)
  - Cache
  - Properties

# Scientific Software Management in Real Life Part IV: EasyBuild

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```
----- INSTALL.SH -----  
#!/bin/bash  
  
pip install "$1" &  
easy_install "$1" &  
brew install "$1" &  
npm install "$1" &  
yum install "$1" & dnf install "$1" &  
docker run "$1" &  
pkg install "$1" &  
apt-get install "$1" &  
sudo apt-get install "$1" &  
steamcmd +app_update "$1" validate &  
git clone https://github.com/"$1"/"$1" &  
cd "$1"; ./configure; make; make install &  
curl "$1" | bash &
```

Source: <http://xkcd.com/1654/>

## Why EasyBuild in JURECA?

- Designed exactly for this use case
- Production ready
- Easily configurable
- Nice integration with Lmod and different Module Naming Schemes
- Active and dynamic project
- Support for over 1000 packages

## Shortcomings

- 1 Was based on *monolithic* toolchains
  - Unnecessary redundancy in package builds.
    - E.g., CMake built with many different toolchains
- 2 Each of the X11 libraries (and other auxiliary libraries) had its own module
  - Swamps default module view with many libraries and their dependencies
- 3 Software that only compiles with GCC couldn't be visible in non-GCC toolchains
- 4 Cryptic toolchain names led to confusion and support issues.

## Implemented, user-driven enhancements I

- Enhanced dependency resolution **1**
  - *Minimal toolchains*
  - Software built with compiler  $x$  version  $y$  and MPI  $z$  version  $w$  can use libraries built just with a toolchain containing compiler  $x$  version  $y$ .
  - Toolchain hierarchy: dummy  $\Rightarrow$  compiler  $\Rightarrow$  MPI  $\Rightarrow$  Math libraries
- Common base compiler (GCC<sub>core</sub>) for toolchains **3 2**
  - Enables base layer for compilers, tools and auxiliary libraries
  - Toolchain hierarchy: dummy  $\Rightarrow$  *GCC<sub>core</sub>*  $\Rightarrow$  compiler  $\Rightarrow$  MPI  $\Rightarrow$  Math libraries

## Implemented, user-driven enhancements II

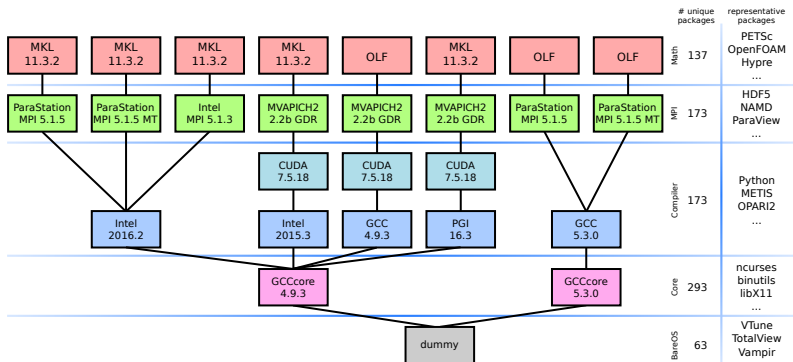
- Support for hidden modules [2](#)
  - Eliminates clutter
  - Supported in various ways (command line options, environment variables, easyconfig parameters)
  - Can hide GCCcore
- Custom module naming schemes [1](#) [4](#)
  - Flat
  - Hierarchical
  - Toolchain based
- Naming scheme-independent software installation directories [4](#)
- Performance improvements
- Refactoring of support for MPICH-based MPI libraries



# Scientific Software Management in Real Life Part V: Current state in JURECA

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# [Old] current state (Stage 2016a)



## User View and Hidden Modules

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  - Binary tools (VTune, Advisor, TotalView, . . .)

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- After loading a compiler:
  - MPI runtimes (ParaStationMPI, MVAPICH2, IntelMPI)
  - Packages built with `gcccore`
  - Packages compiled with the chosen compiler
- After loading an MPI runtime:
  - Packages compiled with the chosen compiler and MPI runtime
- Not all packages available for a given combination are visible:
  - There are almost 400 hidden packages in total!

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- 1 module per extension is excessive  
⇒ Bundles

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  - PyCUDA (6 extensions)
  - numba (2 extensions)



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- R (365 extensions)
- Perl (217 extensions)
- *X.Org* (229 extensions)

## Finding Software

- 1 important option and 3 commands:
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  - `module [--show-hidden]` available
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  - `module [--show-hidden] spider something[/version]`
    - Crawls the module tree looking for modules with `something` on their name
    - Tells what it finds and how to get to it
  - `module key something`
    - Crawls the module tree looking for modules with `<something>` on their description
    - Tells which modules have been found
    - Might need to use `spider` afterwards to find how to get them
    - Useful for looking for the contents of a bundle (ie: `numpy`)

## Upgrading and Retiring Software

- Stage concept:
  - Software deployment area for a given timeframe
  - A simple directory
  - Default stage upgraded every 6 months
  - There is a development stage to test software
  - Tested software is added to our *Golden* repository, and then added to the current production stage
  - Close to seamless transitions between stages during maintenance windows
  - Development and old stages are available but not visible by default

## Ensuring Consistency and Quality

- Software team
  - Allowed to install software in the development stage
  - Can test different compilation options, dependencies, functionality, etc
  - Anybody in the team can modify any other installation
- Software manager
  - Only account allowed to install software in the production stages
  - Supervises quality standards on easyconfigs before adding them to the *Golden* repository
    - Correct dependencies for the production stage
    - Proper programming in easyconfigs and patches (lack of hardcoded paths, use of EB provided variables)
  - Manages the whole infrastructure



## Divergence from Upstream EasyBuild I

- Divergence motivated by
  - Use of latest versions available at deployment time
  - Re-positioning of packages in the toolchain hierarchy
- Most differences are minimal:
  - Different versions of software
  - Different versions of dependencies
  - Different toolchains

EasyConfigs used in JURECA

<b>EB upstream EasyConfigs</b>	47
<b>JSC EasyConfigs</b>	777

## Divergence from Upstream EasyBuild II

- Toolchains divergence

Toolchains used in JURECA.

	EB upstream TCs	JSC TCs
<b>Comp.</b>	3	0
<b>Comp.+MPI</b>	3	3
<b>Comp.+MPI+Math</b>	3	3

## Divergence from Upstream EasyBuild III

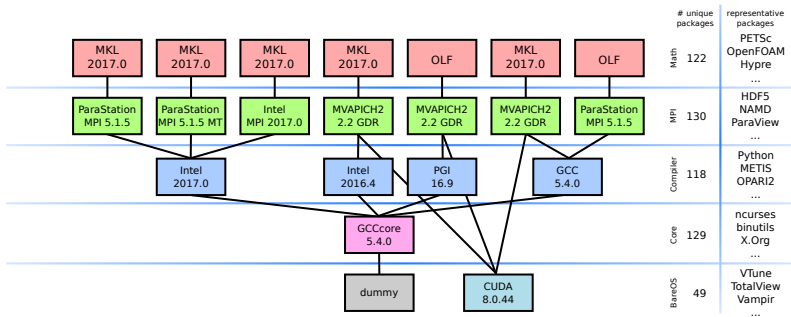
- EasyBlocks divergence

EasyBlocks used in JURECA.

<b>EB upstream EasyBlocks</b>	$\pm 65$
<b>JSC tweaked EasyBlocks</b>	5
<b>JSC merged EasyBlocks</b>	5
<b>JSC private EasyBlocks</b>	4

# Demo

# [New] current state (Stage 2016b)



# Scientific Software Management in Real Life Part VI: Porting to Other Clusters

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## Porting to Other Clusters

- Besides JURECA, JSC also has JUROPA3 and JUAMS
  - Similarities with JURECA: x86\_64, InfiniBand, Red Hat based OS
  - Differences: Different microarchitecture, different OSes, mix of Xeon Phi and GPUs
- Minimal changes needed to reuse JURECA's setup:
  - Fix erroneous easyconfigs
  - Provide new versions in EasyBuild of obsolete OS packages

Software in JUAMS and JUROPA3.

---

<b>Total packages in JUAMS</b>	671
<b>Total packages in JUROPA</b>	658
<b>Ad-hoc packages in both</b>	15

# Scientific Software Management in Real Life Part VII: Future Work

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## Future

- Automatic upgrades
  - Of dependency versions
  - Of software versions
- Default module sets
  - Preselected packages for users that don't care about compilers and MPI runtimes
- Linking with `-rpath` (experimental in EasyBuild 3.0)
- Tracking module usage with XALT
- Reshuffling packages
- “Fat” easyconfigs

# Scientific Software Management in Real Life

## Part VIII: Conclusions

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## Conclusions

- EasyBuild enables to deploy and manage a tremendous amount of software, using a small team
- Active project that grows everyday
- Effort needed to
  - Minimize SW replication
  - Provide latests and greatest (mismatch between our stage switch and EasyBuild releases)
  - Provide a meaningful user view
- EasyBuild enables easy porting to similar systems
- Still room for improvement

*Thank you for listening!*  
You can meet more EasyBuild folks at:

## 2<sup>nd</sup> EasyBuild User Meeting

Jülich Supercomputing Centre (Germany), February 8-10  
[https://github.com/hpcugent/easybuild/wiki/  
2nd-EasyBuild-User-Meeting](https://github.com/hpcugent/easybuild/wiki/2nd-EasyBuild-User-Meeting)

## FOSDEM'17 HPC, Big Data, and Data Science Devroom

Brussels (Belgium), February 4  
<https://hpc-bigdata-fosdem17.github.io/>