

STATENS
SERUM
INSTITUT



Danish
Diabetes
Academy

Environment Modules

Software and library versioning the easy way

Piotr Dworzynski, Pers Lab, NNF Center for Basic Metabolic Research
piotr@sund.ku.dk



The problem

Everyday struggle:

- Tool A requires R-3.2
- Tool B requires R-3.5 and Python 2.7
- Tool C-v1 requires R-3.4 and SomeLibrary v2.3
- Tool C-v2 requires R-3.5 and SomeLibrary v2.5

Environment modules address software:

- version dependencies
- version conflicts
- library conflicts
- reproducibility

Environment modules



Available at <http://modules.sourceforge.net/>

Ubuntu/RedHat/CentOS pkg: environment-modules

Manipulates environmental variables

- e.g. \$PATH, \$LD_LIBRARY_PATH, \$CPLUS_INCLUDE_PATH
- Handles dependencies and conflicts between modules



Basic use-case

By default R points to system R v3.5.0

```
$ which R
> /usr/local/bin/R
$ R
> R version 3.5.0 (2018-04-23) -- "Joy in Playing"
```

With module load command we make R point to a custom R installation

```
$ module load R/3.5.3
$ which R
> /tools/R/3.5.3/bin/R
$ R
> R version 3.5.3 (2019-03-11) -- "Great Truth"
```



Example setup

/tools

- overarching directory for software not coming from yum/apt

/tools/src

- downloaded software sources

/tools/modules/**XXX**/**YYY**

- environment modules live here
- **XXX** – name of the tool (R, anaconda, bwatools)
- **YYY** – plain text module file named as tool's version/flag (3.4, 3.5-openblas)

/tools/**XXX**/**YYY**/**ZZZ**

- software (tools, libraries) lives here
- **XXX** – name of the tool (R, anaconda, bwatools)
- **YYY** – directory named as tool's version/flag (3.4, 3.5-openblas)
- **ZZZ** – software executables/libs (--prefix-path would point here) e.g. /bin /lib /include



Anatomy of an environment module

Contents of example env module (text file; /tools/modules/R/3.5.3)

```
##Module1.0
module-whatis "R is a free software environment for statistical computing and graphics."

conflict R

prepend-path PATH /tools/R/3.5.3/bin/
prepend-path MANPATH /tools/R/3.5.3/share/man
prepend-path LD_LIBRARY_PATH /tools/R/3.5.3/lib64/
prepend-path LD_LIBRARY_PATH /tools/R/3.5.3/lib64/R/library/
prepend-path C_INCLUDE_PATH /tools/R/3.5.3/lib64/R/include/
prepend-path CPLUS_INCLUDE_PATH /tools/R/3.5.3/lib64/R/include/
prepend-path R_ROOT /tools/R/3.5.3/lib64

setenv RINSIDE_LIBRARY /tools/R/3.5.3/lib64/R/library
setenv RINSIDE_INCLUDE_DIR /tools/R/3.5.3/lib64/R/include
```



Use cases & Limitations

Use cases:

- Basic tool/library version management
- Data version management (e.g. reference genomes)
- Pipeline tool/library/data version management
- Ansible & env-modules powered software collections akin to [bio-ansible](https://bio-ansible.org/)

Limitations:

- Python/R packages management
- Not meant for very complex situations
- Supports only specific shells

Thanks!

STATENS
SERUM
INSTITUT



Danish
Diabetes
Academy



Learn more:

- [official webpage](#)
- [official documentation](#)
- [tutorial 1](#) (admin-magazine.com)
- [tutorial 2](#) (geoghegan.me)

Followup discussion:

A common software repository for NNF centers?

Ansible, EnvModules, Docker/Singularity images

Stay in touch:

- Piotr Dworzynski
- p.dworzynski@gmail.com
- piotr@sund.ku.dk