DAY 4

# Day 4: Essential pieces of reproducibility - Conda, Jupyter, Galaxy, Blogs

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Conda and Bioconda

#### **BLACKDUCK** | Hub **Hub Detect**



















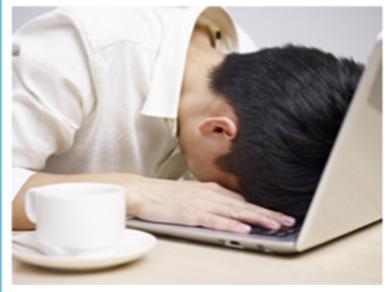
## Linux Package Managers: APK, DPKG, RPMs











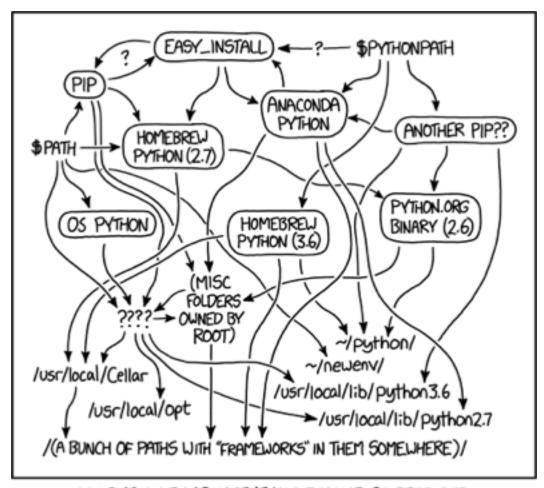
From: https://www.cad-notes.com/wpcontent/uploads/2015/11/frustratedcomputer-user.jpg

#### Conda

Package, dependency and environment management for any language

"Conda is an open source package management system and environment management system that runs on Windows, macOS and Linux. Conda quickly installs, runs and updates packages and their dependencies. Conda easily creates, saves, loads and switches between environments on your local computer. It was created for Python programs, but it can package and distribute software for any language."

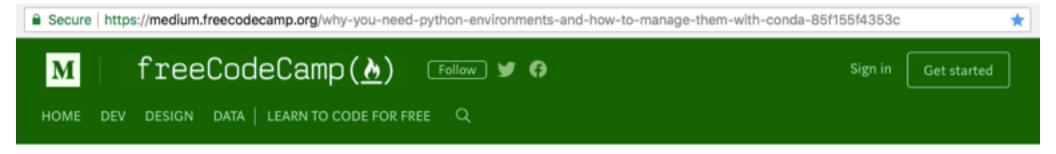
from: <a href="https://conda.io/docs/">https://conda.io/docs/</a>



MY PYTHON ENVIRONMENT HAS BECOME SO DEGRADED THAT MY LAPTOP HAS BEEN DECLARED A SUPERFUND SITE.

which python

## Recommended blog post

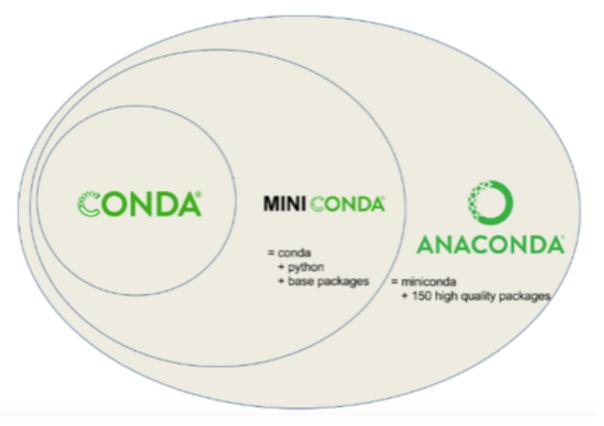




# Why you need Python environments and how to manage them with Conda



#### **Conda Distributions**



Slide from Galaxy Team and John Chilton

### Conda can be used to install:

- Python (with disclaimer)
- Python packages
- R
- Tools and packages for bioinformatics and data science
- √ Specific versions

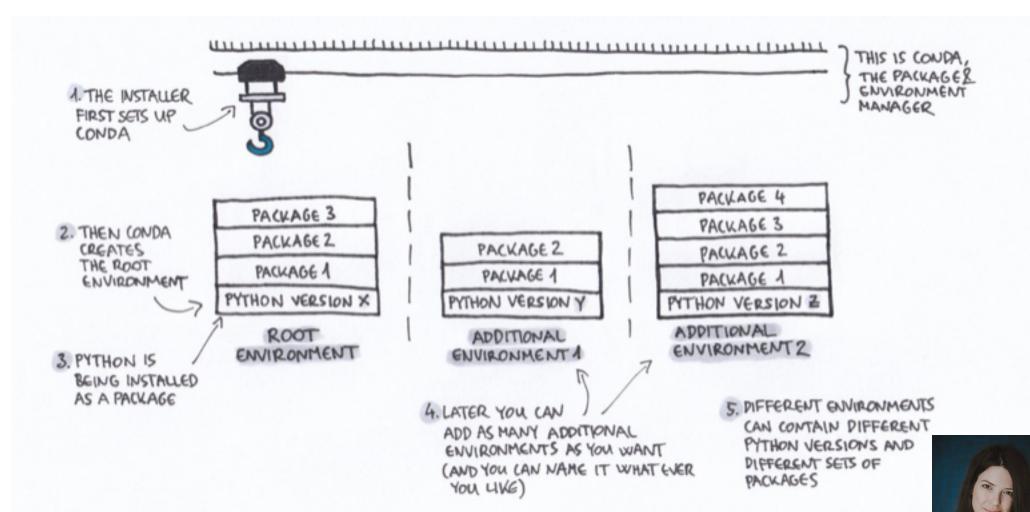
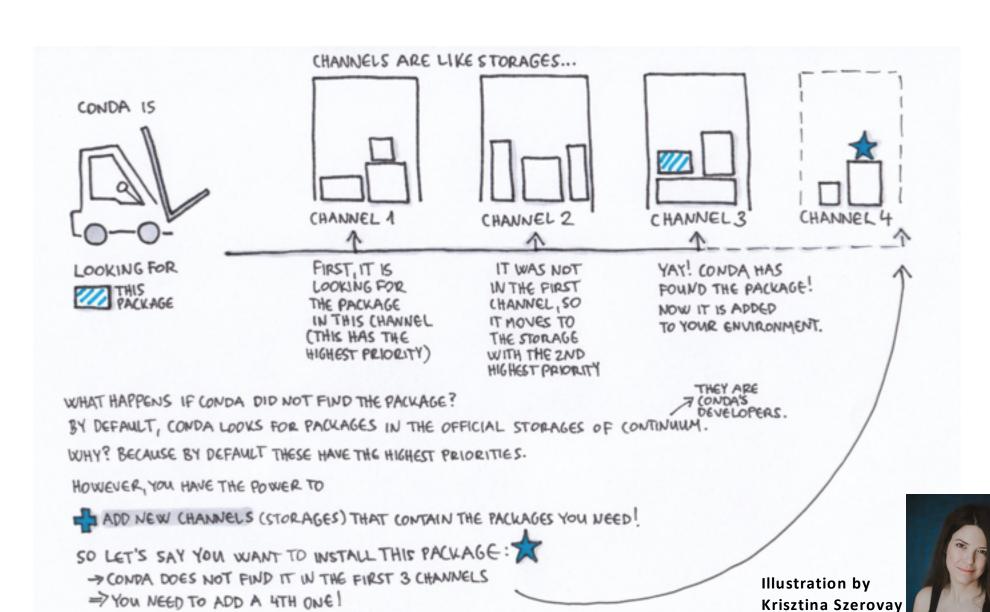
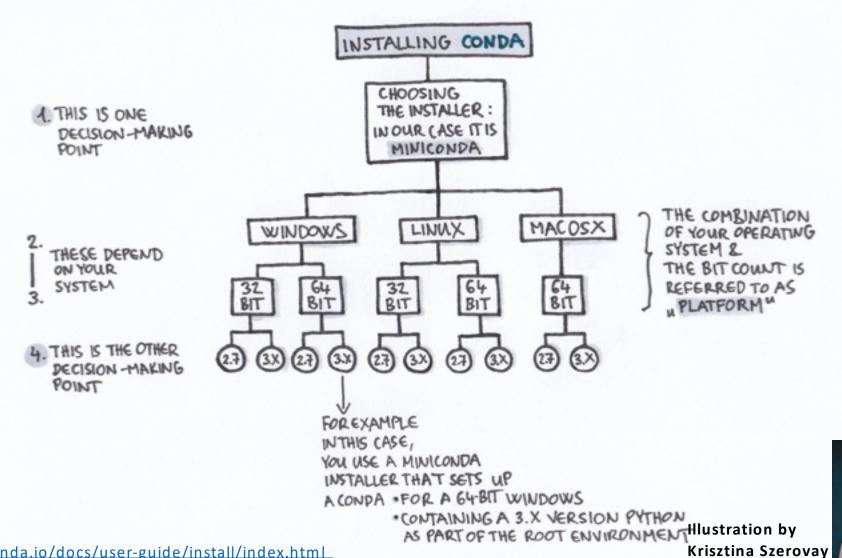


Illustration by

Krisztina Szerovay



Questions?





#### General conda commands

- conda --help
- conda list #all packages available in a given environment
- conda env list #lists all environments

#### Create and activate new environment

- conda create --name test
- source activate test
- conda install -c bioconda plink
- conda install -c bioconda minimap
- conda install -c bioconda bioawk
- conda install --channel r r-essentials=1.4
- conda install --channel r r=3.3.1
- source deactivate

# Looking for packages to install

**Newest version:** 

Google conda name\_of\_the\_package

Which versions of the package are available? conda search -f samtools

#### Power of bioconda

- conda search --channel bioconda
- conda search --channel bioconda | wc -l

# BIOCONDA

# Installing perl modules via conda

- conda create --name perly\_env
- source activate perly\_env
- conda install -c anaconda perl # Installing perl
- conda install -c bioconda perl-app-cpanminus # Installing CPAN for managing perl modules
- If -c bioconda does not work, try:

  conda install -c conda-forge perl-app-cpanminus
- cpanm Bio::Perl # Getting any module you like

Commands courtesy of Samarth Rangavittal

https://hcc-docs.unl.edu/display/HCCDOC/Installing+Perl+modules

# Questions?

