CONDA CONDA CHEAT SHEET

Take a conda test drive at bit.ly/tryconda

in the latest release of Anaconda, all of which are tested for compatibility. May not contain the newest versions.

NOTE: Environments install by default into the envs directory in your conda directory. You can specify a different path; see conda create --help for details.

For full documentation of any command, type the command followed by $--\mathtt{help}$,

such as conda create --help

TIP: Anaconda Navigator is a point-and-click way to manage packages and environments with conda. For example, with Navigator you can run Jupyter Notebooks or Spyder without using a terminal. If you have Anaconda, Navigator is already installed. Double-click the Navigator icon on your desktop

MANAGING CONDA & ANACONDA	
Verify conda is installed, check version number,	conda info
see basic information about conda	
Update conda to the current version	conda update conda
Update all packages in the environment to the versions	conda update anaconda

USING ENVIRONMENTS	
Get a list of all my environments, active	conda infoenvs
environment is shown with *	
List all packages and versions installed in active environment	conda list
Create an environment named bio-env in your home	conda createprefix -/bio-env biopython
directory and install the biopython package	

Acti	ivate the new environment to use it	LINUX, MAC: source activate ~/bio-env
		WINDOWS: activate ~/bio-env
Сге	ate a new environment named py34, specify	conda createprefix ~/py34 python=3.4 astroid
Pyt	hon version, install astroid package	
Mal	ke exact copy of an environment	conda createprefix ~/bioenvcopyclone bio-env
Dea	activate the current environment	LINUX, MAC: source deactivate
		WINDOWS: deactivate
List	the history of each change to the current environment	conda listrevisions
Res	tore the environment to a previous revision	conda installrevision 2
Del	ete an environment method 1	conda removeprefix ~/bioenvcopyall

Delete an environment method 2	LINUX, MAC: rm -rf -/bioenvcopy	
	WINDOWS: rmdir /s bioenvcopy	
Save environment to a text file	conda listexplicit > bio-env.txt	
Load environment from a text file	conda createprefix -/bio-envfile bio-env.txt	

FINDING CONDA PACKAGES	
Go to Anaconda Cloud in the browser	https://anaconda.org
and search by package name	
Use conda to search for a package (Numba)	conda search numba
View list of all packages associated with Anaconda	https://docs.continuum.io/anaconda/pkg-docs



INSTALLING PACKAGES	
Install a new package (Jupyter Notebook)	conda install jupyter
in the current environment	
Run an installed package (Jupyter Notebook)	jupyter-notebook
Install a new package (toolz) in a	conda installprefix -/bio-env toolz
different environment (~/bio-env)	
Update a package in the current environment	conda update toolz
Install a package (boltons) from a	conda install -c conda-forge boltons
specific channel (conda-forge)	
Install a package directly from PyPI into the	pip install boltons
current active environment using pip	
Remove one or more packages (toolz, boltons)	conda removeprefix ~/bio-env toolz boltons
from a specific environment (~/bio-env)	
NOTE: If you do not include the path of the environment	

active environment.	
MANAGING MULTIPLE VERSIONS OF PYTHON	
Install different version of Python in a	conda createprefix -/py34 python=3.4
new environment named py34	
Switch to the new environment that	LINUX, MAC: source activate -/py34
has a different version of Python	WINDOWS: activate -/py34
Show the locations of all versions of Python	LINUX, MAC: which -a python
that are currently in the PATH	WINDOWS: where python
NOTE: The first version of Python in the list will be executed.	
Show version information for the current active Python	pythonversion

SPECIFYING VERSION NUMBERS

(~/bio-env), conda removes packages from the current

Ways to specify a package version number for use with conda create or conda install commands, and in meta.yaml files.

Constraint type	Specification	Result
Fuzzy	numpy=1.11	1.11.0, 1.11.1, 1.11.2, 1.11.18 etc.
Exact	numpy==1.11	1.11.0
Greater than or equal to	"numpy>=1.11"	1.11.0 or higher
OR	"numpy=1.11.1 1.11.3 "	1.11.1, 1.11.3
AND	"numpy>=1.8,<2"	1.8, 1.9, not 2.0

NOTE: Quotation marks must be used when your specification contains a space or any of these characters: > < | *

MORE RESOURCES

https://groups.google.com/a/continuum.io/forum/#!forum/anaconda Free Community Support

Online Documentation http://conda.io

Paid Support Options https://www.continuum.io/support Continuum Onsite Training Courses https://www.continuum.io/training

https://www.continuum.io/continuum-consulting Continuum Consulting Services

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Connect with other talented, like-minded data scientists and developers while contributing to the open source movement. Visit https://continuum.io/community

