

Python and Conda Environments in HPC:

From Basics to Best Practices

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Outline

- Why High Performance Computing
- How to access Python on Wulver at HPC
- Introduction to Conda environments
- Install, uninstall and upgrade packages
- Best Practices for managing conda environments
- Common Python libraries for scientific computing

Why High Performance Computing?

- Handling Complex Problems
- Big Data Analysis
- Speeding up Research
- Parallel Computing
- Resource Sharing and Collaboration

Why Use Python for HPC?

- **Clear Syntax** – Simple, readable, and easy to learn
- **Extensive Libraries** – Optimized packages for scientific computing
- **Multi-language Integration** – Works seamlessly with C, C++, and Fortran
- **Parallel Computing Capabilities** – Supports multi-threading & distributed computing
- **Strong Community Support** – Actively maintained & widely adopted

Python on Wulver

Software	Version	Dependent Toolchain	Module Load Command
Python	3.13.1	foss/2025a	<code>module load foss/2025a Python/3.13.1</code>
Python	3.12.3	foss/2024a	<code>module load foss/2024a Python/3.12.3</code>

Installing Python packages

Method 1: Installing Python Packages from Source

1 Clone the repository

```
$ git clone https://github.com/pandas-dev/pandas.git
```

2 Navigate into the package directory

```
$ cd pandas
```

3 Install the package to a custom location

```
$ python setup.py install --prefix=/project/$GROUP/$USER/python_pkg/
```

Possible Installation Error

Error Message:

Traceback (most recent call last):

File "/usr/lib64/python3.6/site-packages/numpy/core/__init__.py", line 16, in
<module>

from . import multiarray

ImportError: libopenblas.so.0: cannot open shared object file: No such file or directory

Reason: The required shared library (`libopenblas.so.0`) is missing or not found.

Installing Python packages - PiP

Method 2: pip

pip – stands for “**Preferred Installer Program**”

A package manager for Python packages **only**

Installs packages from the **Python Package Index (PyPI)**

```
$ python -m pip install --user <python-module-name> --no-cache-dir
```

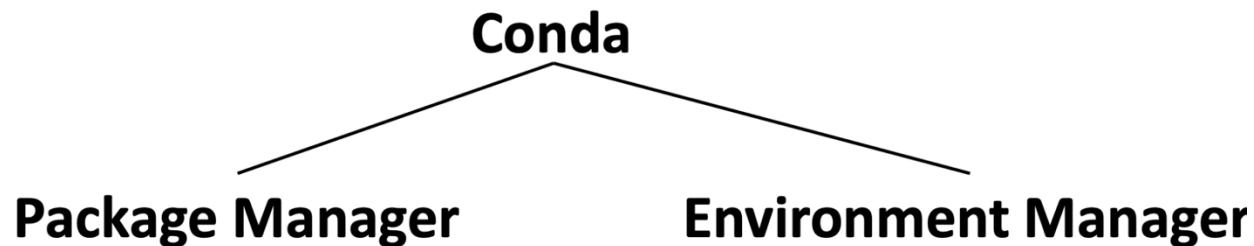
- -m pip – Run the pip module *using this exact Python*
- --user – Install to your user account only
- --no-cache-dir – Disable pip’s cache (helpful for small quotas)

Conda on HPC

- **Introduction to Conda**
- Conda channels
- Conda environment
- Conda packages
- Sharing environments

Introduction to Conda

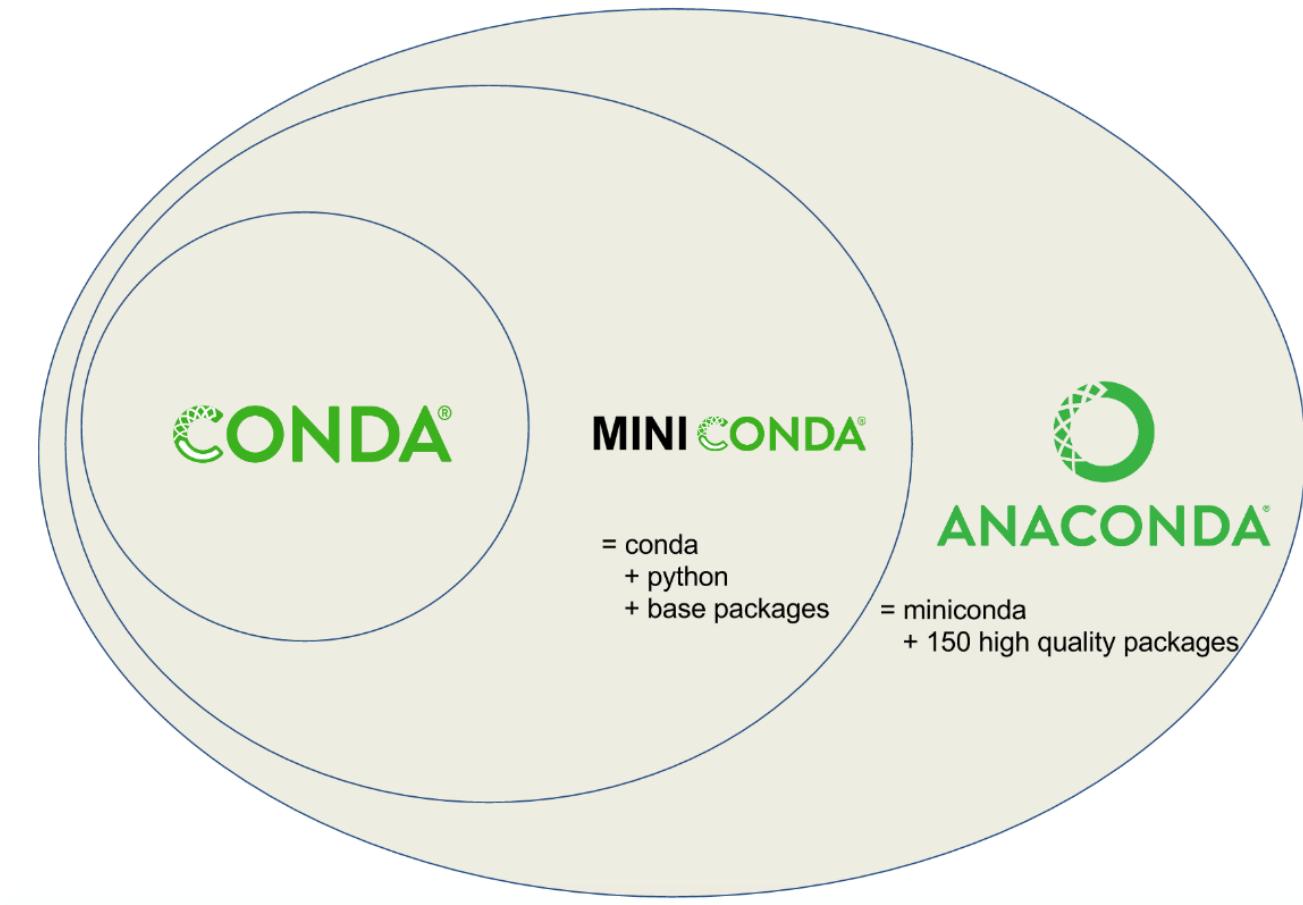
- **Conda is an open-source package and environment manager**
Supports **Python and non-Python** packages
Works across **Windows, macOS, and Linux**
- **Conda is a powerful package & environment manager**



Why use Conda?

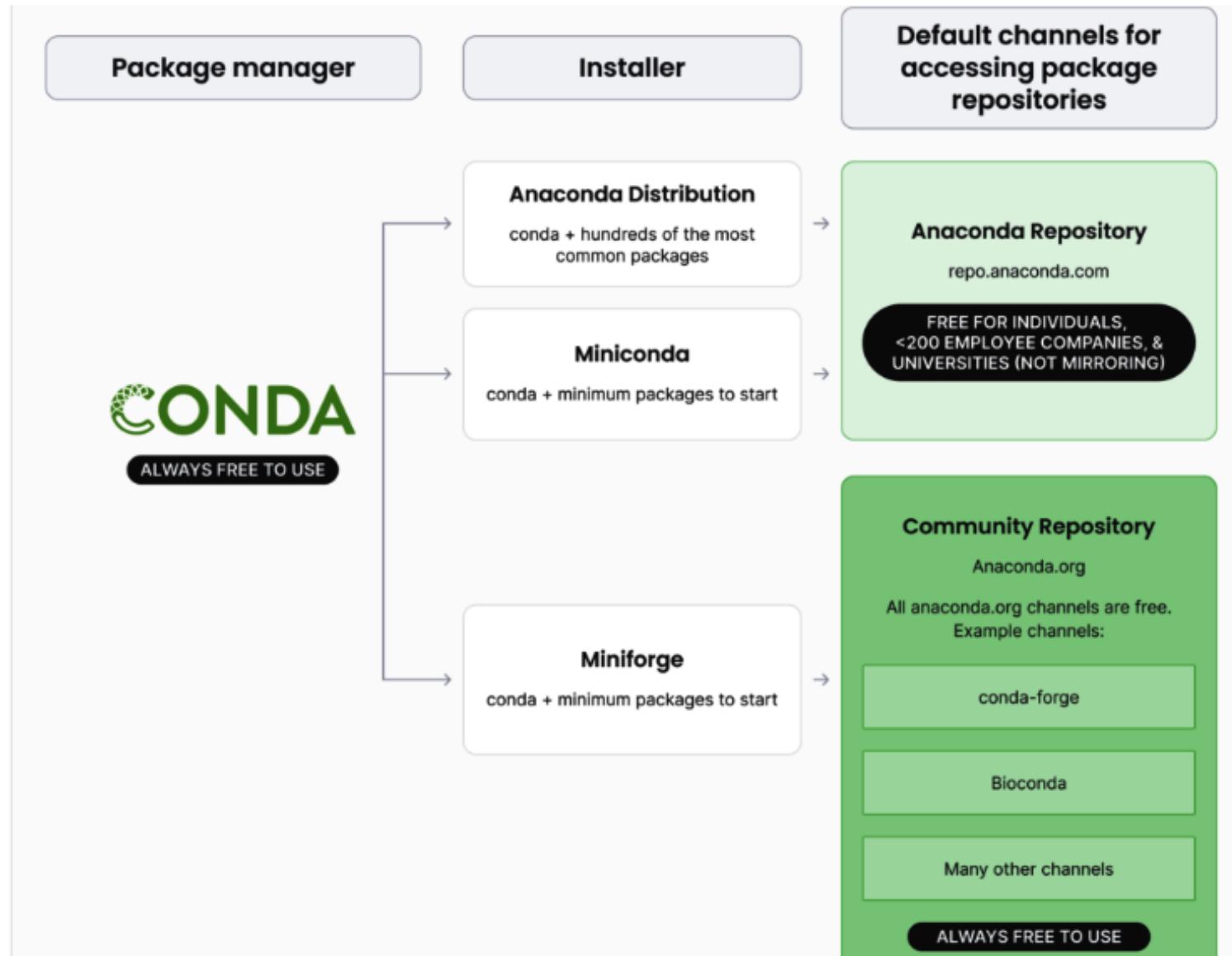
- Simplifies package management and installation
- Automatically handles dependencies
- Creates isolated environments to avoid conflicts
- Supports Python, R, C, and more
- Reproducible and portable across platforms
- Ideal for HPC and scientific computing

Anaconda vs Miniconda vs Conda



- Conda: Open-source tool
- Anaconda: A software distribution: open-source (personal) and Commercial
- Miniconda: minimal installer for conda

Anaconda Portfolio



Load Miniforge Module on Wulver

Load Miniforge3 Module

```
$ module load Miniforge3
```

Check Loaded Modules

```
$ module list
```

Currently Loaded Modules:

- 1) easybuild 3) slurm/wulver 5) Miniforge3/24.11.3-0
- 2) wulver 4) null

What is Miniforge

\$ module whatis Miniforge3

Miniforge3/24.11.3-0 : Description: Miniforge is a free minimal installer for conda and Mamba specific to conda-forge.

Miniforge3/24.11.3-0 : Homepage: <https://github.com/conda-forge/miniforge>

Miniforge3/24.11.3-0 : URL: <https://github.com/conda-forge/miniforge>

Conda info

```
g07396@login02:~$ conda info

active environment : None
shell level       : 0
user config file  : /home/g07396/.condarc
populated config files : /apps/easybuild/el9_5.x86_64/software/Miniforge3/24.11.3-0/.condarc
                        /home/g07396/.condarc
conda version     : 24.11.3
conda-build version: not installed
python version    : 3.12.8.final.0
solver           : libmamba (default)
virtual packages  : __archspec=1=icelake
                    __conda=24.11.3=0
                    __glibc=2.34=0
                    __linux=5.14.0=0
                    __unix=0=0
base environment   : /apps/easybuild/el9_5.x86_64/software/Miniforge3/24.11.3-0 (read only)
conda av data dir  : /apps/easybuild/el9_5.x86_64/software/Miniforge3/24.11.3-0/etc/conda
conda av metadata url: None
channel URLs      : https://conda.anaconda.org/conda-forge/linux-64
                    https://conda.anaconda.org/conda-forge/noarch
                    https://repo.anaconda.com/pkgs/main/linux-64
                    https://repo.anaconda.com/pkgs/main/noarch
                    https://repo.anaconda.com/pkgs/r/linux-64
                    https://repo.anaconda.com/pkgs/r/noarch
package cache     : /project/walsh/g07396/conda_env/pkgs_dirs
envs directories  : /home/g07396/.conda/envs
                    /apps/easybuild/el9_5.x86_64/software/Miniforge3/24.11.3-0/envs
platform          : linux-64
user-agent        : conda/24.11.3 requests/2.32.3 CPython/3.12.8 Linux/5.14.0-503.11.1.el9_5.x86_64 rhel/9.6 glibc/2.34 solver/libmamba conda-libmamba-solver/24.9.0 libmambapy/1.5.12
UID:GID          : 580857:580857
netrc file        : None
offline mode     : False
```

Conda on HPC

- Introduction to Conda
- **Conda channels**
- Conda environment
- Conda packages
- Sharing environments

Configuring Conda channels

A conda channel is a repository of conda packages

```
$ conda config --help
```

```
$ conda config --show
```

```
$ conda config --show channels
```

channels:

- conda-forge
- defaults

```
$conda config --describe channels
```

```
$conda config --add channels bioconda
```

This would add the conda-forge channel to the top of the channel list.

```
$conda config --append channels bioconda
```

This would add the bioconda to the end of the channel list, giving it the lowest priority.

Conda on HPC

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Why create a Conda environment?

A **conda environment** is a directory that contains a specific collection of conda packages.

Isolation from other projects

Control Over Packages

- Manage versions and dependencies.

Reproducibility

- Consistent setups across systems.

Dependency Management

- Handles Python and non-Python dependencies.

Python Versatility

- Manage and switch Python versions easily.

Cross-Platform

- Works on Linux, Windows, and macOS.

Commonly used Conda commands

Task	Command
Activate environment:	<code>conda activate [environment_name]</code>
Deactivate environment:	<code>conda deactivate [environment_name]</code>
Show the list of environments:	<code>conda env list</code>
Delete environment:	<code>conda remove [environment_name]</code>
Export environment:	<code>conda env export > [environment_name].yml</code>
Import environment from YAML:	<code>conda env create -f [environment_name].yml</code>
Import environment to different location:	<code>conda env create -f [environment_name].yml -p [PATH]</code>

[Conda cheat sheet](#) - Link to Conda Doc for more helpful commands



Creating Conda Environment

Creating a new conda environment

```
$ conda create --name my_env
```

Creating a new conda environment with a specific python version

```
$ conda create --name my_env python=3.9
```

Creating a new conda environment with a specific python version and scipy package

```
$ conda create --name my_env python=3.9 scipy=0.15.0
```

Creating a new conda environment in difference location with **--prefix** or **-p**

```
$ conda create --prefix /project/$GROUP/$USER/conda_env/AAA
```

Enter, Exit and Remove conda environment

Entering a Conda environment

```
$ conda activate my_env
```

```
$ conda activate /project/$GROUP/$USER/conda_env/AAA
```

Exiting a Conda environment we are currently in

```
$ conda deactivate
```

Removing a Conda environment

```
$ conda env remove -n my_env
```

Renaming a Conda environment

```
$conda rename -n old_env_name new_env_name
```

List conda environments

A user may list all shared virtual environments and your own private virtual environments

```
g07396@login02:~$ conda info --envs

# conda environments:
#
base          /apps/easybuild/el9_5.x86_64/software/Miniforge3/24.11.3-0
matlab-proxy   /apps/easybuild/el9_5.x86_64/software/Miniforge3/24.11.3-0/envs/matlab-proxy
r_env         /home/g07396/.conda/envs/r_env
tensorflow     /home/g07396/.conda/envs/tensorflow
torch-cuda    /home/g07396/.conda/envs/torch-cuda
              /project/walsh/g07396/conda_env/torch-cuda_12.1
              /project/walsh/g07396/conda_env/torch-cuda_12.2
```

```
g07396@login02:~$ conda env list
```

```
# conda environments:
#
base          /apps/easybuild/el9_5.x86_64/software/Miniforge3/24.11.3-0
matlab-proxy   /apps/easybuild/el9_5.x86_64/software/Miniforge3/24.11.3-0/envs/matlab-proxy
r_env         /home/g07396/.conda/envs/r_env
tensorflow     /home/g07396/.conda/envs/tensorflow
torch-cuda    /home/g07396/.conda/envs/torch-cuda
              /project/walsh/g07396/conda_env/torch-cuda_12.1
              /project/walsh/g07396/conda_env/torch-cuda_12.2
```

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Conda packages

A conda package is a compressed tarball file

List All Installed Packages:

- **\$ conda list**
- Displays all packages installed in the active Conda environment.

List Packages in a Specific Environment:

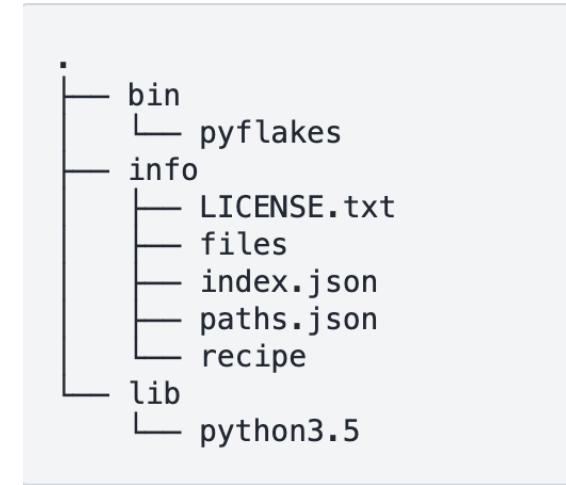
\$ conda list -n env_name or conda list -p /path/to/environment

Search for a Package:

- **\$ conda search package_name**
- Searches for a package across all channels in Conda.

Check for Specific Package Installation:

- **\$ conda list package_name -n myenv**
- **\$ conda list -n myenv | grep package_name**
- Filters the list of installed packages to show only the entries related to package_name.



List packages in all environments

```
g07396@n0094:~$ conda list
# packages in environment at /apps/easybuild/el9_5.x86_64/software/Miniforge3/24.11.3-0:
#
# Name           Version        Build  Channel
_libgcc_mutex   0.1            conda_forge
_openmp_mutex   4.5            2_gnu   conda-forge
archspec        0.2.5          pyhd8ed1ab_0
boltons         24.0.0          pyhd8ed1ab_1
brotli-python   1.1.0          py312h2ec8cdc_2
bzip2           1.0.8          h4bc722e_7
c-ares          1.34.4          hb9d3cd8_0
ca-certificates 2024.12.14    hbcca054_0
certifi         2024.12.14    pyhd8ed1ab_0
cffi            1.17.1          py312h06ac9bb_0
charset-normalizer 3.4.1          pyhd8ed1ab_0
colorama        0.4.6          pyhd8ed1ab_1
conda           24.11.3          py312h7900ff3_0
conda-libmamba-solver 24.9.0    pyhd8ed1ab_0
conda-package-handling 2.4.0     pyh7900ff3_2
conda-package-streaming 0.11.0    pyhd8ed1ab_0
distro          1.9.0          pyhd8ed1ab_1
fmt             11.0.2          h434a139_0
```

List packages in an environment

```
g07396@n0094:~$ conda list -n tensorflow
# packages in environment at /home/g07396/.conda/envs/tensorflow:
#
# Name           Version        Build  Channel
_libgcc_mutex   0.1            conda_forge  conda-forge
_openmp_mutex   4.5            2_gnu      conda-forge
bzip2          1.0.8          h4bc722e_7  conda-forge
ca-certificates 2025.1.31    hbd8a1cb_1  conda-forge
certifi         2025.1.31    pyhd8ed1ab_0  conda-forge
ld_impl_linux-64 2.43          h712a8e2_4  conda-forge
libffi          3.4.6          h2dba641_0  conda-forge
libgcc          14.2.0         h767d61c_2  conda-forge
libgcc-ng       14.2.0         h69a702a_2  conda-forge
libgomp         14.2.0         h767d61c_2  conda-forge
liblzma         5.6.4          hb9d3cd8_0  conda-forge
libnsl          2.0.1          hd590300_0  conda-forge
```

List the installed packages for the present environment

(myenv) \$ conda list

Installing Conda packages

- Entering a Conda environment
 - \$ **conda activate my_env**
 - (my_env) \$: **conda install scipy=1.6 --channel conda-forge**
- Create an environment called my_biowork-env and install blast from the bioconda channel:
 - \$ **conda create --name my_biowork-env blast --channel bioconda**
- The name flag can be used to specify the environment in which we install the package
 - \$ **conda install -n my_env scipy**

```
$conda install conda-forge::tensorflow --prefix /project/$GROUP/$USER/my_env
```

Example - Install PyTorch with GPU

```
$conda create --name torch-cuda python=3.10  
$conda activate torch-cuda  
$conda install -c "nvidia/label/cuda-12.2.0" cuda-toolkit  
$conda install -c pytorch -c nvidia pytorch torchvision torchaudio pytorch-cuda -y
```

A simple PyTorch test program is given below to check whether PyTorch has been installed properly. Program is called

 **torch_tensor.py**



User can use the following job script to run the script.

 **torch-cuda.submit.sh**



<https://hpc.njit.edu/Software/programming/python/conda/#install-tensorflow-with-gpu>

Mamba

Mamba is a reimplementation of the conda package manager in C++ for maximum efficiency

- Parallel downloading of repository data and packages files using multi-threading
- Libsolv for much faster dependency solving
- a *drop-in* replacement for conda
- Same commands as conda
- Robust and fast but not 100% drop-in replacement yet (especially for conda-env commands)

<https://mamba.readthedocs.io/en/latest/>

Mamba on Wulver

```
module load Miniforge3

# create new environment
mamba create --name env_name python numpy pandas
# install a new package into an existing environment
conda activate env_name
mamba install scipy
```

[Mamba on wulver](#)



Conda on HPC

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- Conda channels
- Conda packages
- **Sharing environments**

Exporting Conda environment

Export a conda environment to a new directory or a different machine

1. activate the environment first that you intend to export.
2. export it to a YAML file:

```
$ conda env export > my_environment.yml
```

```
name: my_env
channels:
- defaults
dependencies:
- _libgcc_mutex=0.1=main
- _openmp_mutex=5.1=1_gnu
- blas=1.0=mkl

<output snipped>

#the last line is the path of the env
prefix: /home/a/abc3/.conda/envs/my_env.
```

Create an Conda environment from YAML file

- First load Miniforge
- Create the environment from the YAML file:

```
conda env create -f my_environment.yml
Collecting package metadata (repodata.json): done
Solving environment: done

<output snipped>

Downloading and Extracting Packages
Preparing transaction: done
Verifying transaction: done
Executing transaction: done
#
# To activate this environment, use
#
# $ conda activate my_env
#
# To deactivate an active environment, use
#
# $ conda deactivate
```

Importing Conda environment to a new location

Use the **--prefix** or **-p** option to **specify the environment location**

- \$ **conda env create -f my_environment.yml -p /project/\$GROUP/\$USER/conda_env/my_env**
- This will create the environment in the specified directory instead of the default conda environment directory.

Provide the full path of the environment to activate it.

- \$ **conda activate /project/\$GROUP/\$USER/conda_env/my_env**
- \$ **conda env list**

conda environments:

```
base      /mmfs1/apps/easybuild/software/Miniforge3/24.11.3-0
          * /project/$GROUP/$USER/conda_env/my_env
```

Updating a Conda environment

When to Update:

- A core dependency has a new version

- You need to add a new package (e.g., for analysis or visualization)

- You want to remove outdated packages or switch to better ones

Example: Update a Single Package

```
$ conda update -n myenv scipy
```

Update from environment.yml

```
$ conda env update --name=foo --file=environment.yml --prune
```

--prune removes packages no longer required and keeps your environment clean

Tip:

Always save and version-control your environment.yml file to track changes.

Conda Revision and history

Task	Command	Description
List revisions	conda list --revisions	Show numbered history of changes
Roll back	conda install --revision <n>	Restore environment to revision <n>
Export current state	conda env export > env.yaml	Backup environment definition

Best practices

Use interactive sessions on a compute node

- Use an interactive session on a compute node
- `$ srun -p general -n 1 --qos=standard --account=$PI_ucid --mem-per-cpu=2G --time=59:00 --pty bash`
- `$ interactive -a $PI_UCID -q standard -j cpu`

Use /project directory with large quotas

- Use /project/\$PI/\$USER directory other than the home directory for conda environments and packages.
- Managing Conda Cache and changing the default caching behavior

Avoid installing packages into your base Conda environment

Conda Cache

Default Location: **\$HOME/.conda/pkgs**

Check Current Cache Directory: **conda info**

Change Cache Location:

- Edit **.condarc**
pkgs_dirs:
 - /path/to/desired/cache/directory
- Use Conda Command:
conda config --add pkgs_dirs /project/\$GROUP/\$USER/conda_env/pkgs_dirs
- Set Environment Variable:
export CONDA_PKGS_DIRS=/path/to/desired/cache/directory

Clean Cache and Unused Packages

- **conda clean --tarballs**
- **conda clean --all** (remove index cache, lock files, unused cache pkgs, tarballs, logs)

More Options: [**official .condarc user guide**](#)



PiP vs Conda



- Favor Conda over Pip whenever possible**
- Use Conda first, then Pip only if necessary**

Why Choose Conda?

- Pre-compiled packages** – No need to build from source
- Automatic dependency resolution** – Handles package conflicts
- Better for scientific computing** – Optimized for numerical libraries

When to Use Pip?

- If the package **is not available** in Conda
If you need **the latest version** of a package \$ pip install latest-package

Pip drawbacks

- Dependencies may need manual resolution
Possible compatibility issues with Conda-installed packages

Pip installs in a Conda environment

Recommend

- Use Conda environments for isolation
- Always install Conda packages first, then use `pip`
- ⚠ Avoid installing Conda packages after using `pip`

Create and activate a Conda environment

```
$ conda create --name my_env pandas  
$ conda activate my_env  
Install additional packages with pip  
(my_env)$ python -m pip install multiregex
```

Recreate the environment if you need to modify packages after using `pip`

- once pip has been used conda will be unaware of the changes
- to install additional conda packages it is best to recreate the environment

Store conda and pip requirements in text files

⚠ Use `--no-cache-dir` to prevent pip from filling your home directory
(my_env)\$ `python -m pip install --no-cache-dir package_name`

Refer to [Conda guide for using pip in a Conda environment](#)

Common Python libraries for scientific computing

Library	Key features	Common Use Case
Numpy	Multidimensional arrays, Broadcasting, Vectorization	Mathematical operations, Basic statistics
SciPy	Numerical integration, Optimization, Linear algebra	Solving differential equations, Signal processing
Matplotlib	2D and 3D plotting, Customizable plots	Visualizing data, Scientific charts
Pandas	DataFrame and Series, Data manipulation, Cleaning	Data analysis, Time series analysis
Scikit-learn	Machine learning algorithms, Data preprocessing tools	Classification, Regression, Clustering
TensorFlow	Computational graph, Automatic differentiation	Building deep learning models, Neural networks
PyTorch	Dynamic computational graph, TorchScript for deployment	Machine learning, Computer vision

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Questions?