

Python and Conda Environments in HPC: *From Basics to Best Practices*

Hui (Julia) Zhao

NJIT High Performance Computing

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 | 3.9.6 | foss/2021b | <code>module load foss/2021b Python/3.9.6</code> |
| Python | 3.11.5 | foss/2023b | <code>module load foss/2023b Python/3.11.5</code> |
| Python | 3.10.8 | foss/2022b | <code>module load foss/2022b Python/3.10.8</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 <module-name>: Always use `python -m pip` instead of just `pip`

Ensures pip runs using the **correct Python interpreter**

Avoids conflicts with multiple Python installations

--user Flag: Installs packages **to user account only**

Ensures installation **without admin/root privileges**

Useful on shared HPC systems

--no-cache-dir Option: Prevents pip from storing package caches in the home directory

Saves disk space, especially in HPC environments

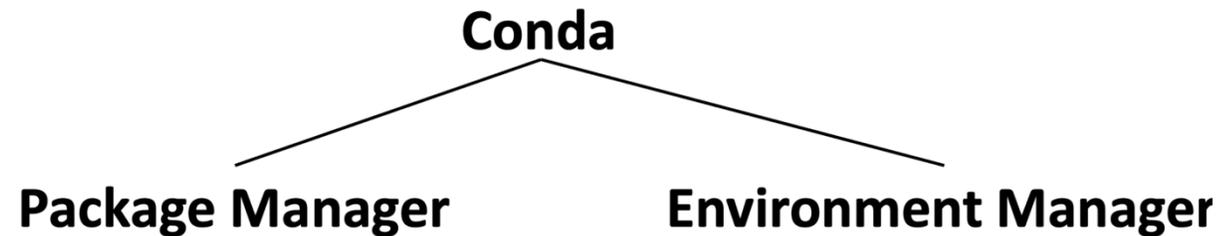
Method 3: Conda

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?

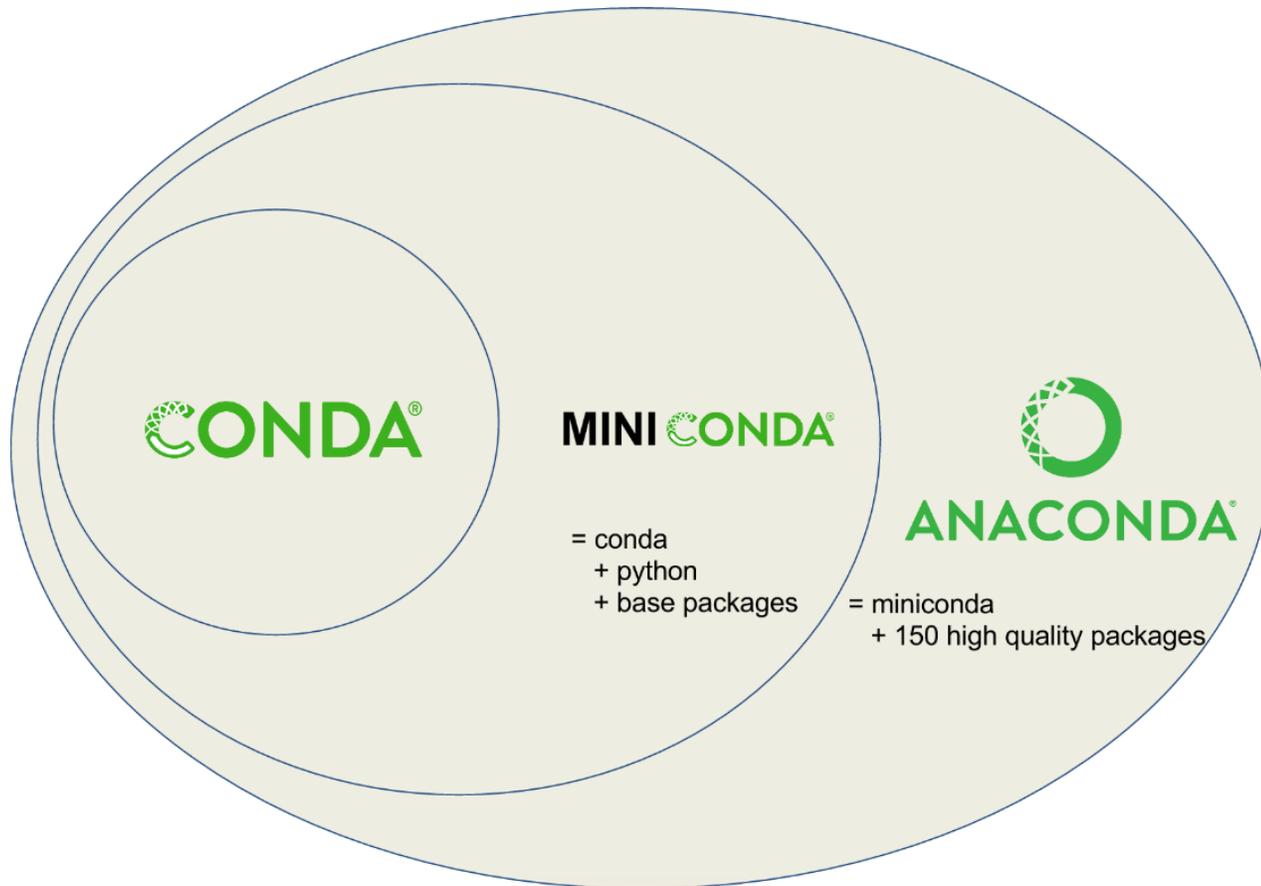
Key Benefits

- Simplifies installation, dependency resolution, and reproducibility
- Ideal for scientific computing, data science, and HPC
- Manages dependencies automatically
- Creates isolated environments to prevent package conflicts
- Supports multiple programming languages, including R, C, and Java

Additional Advantages

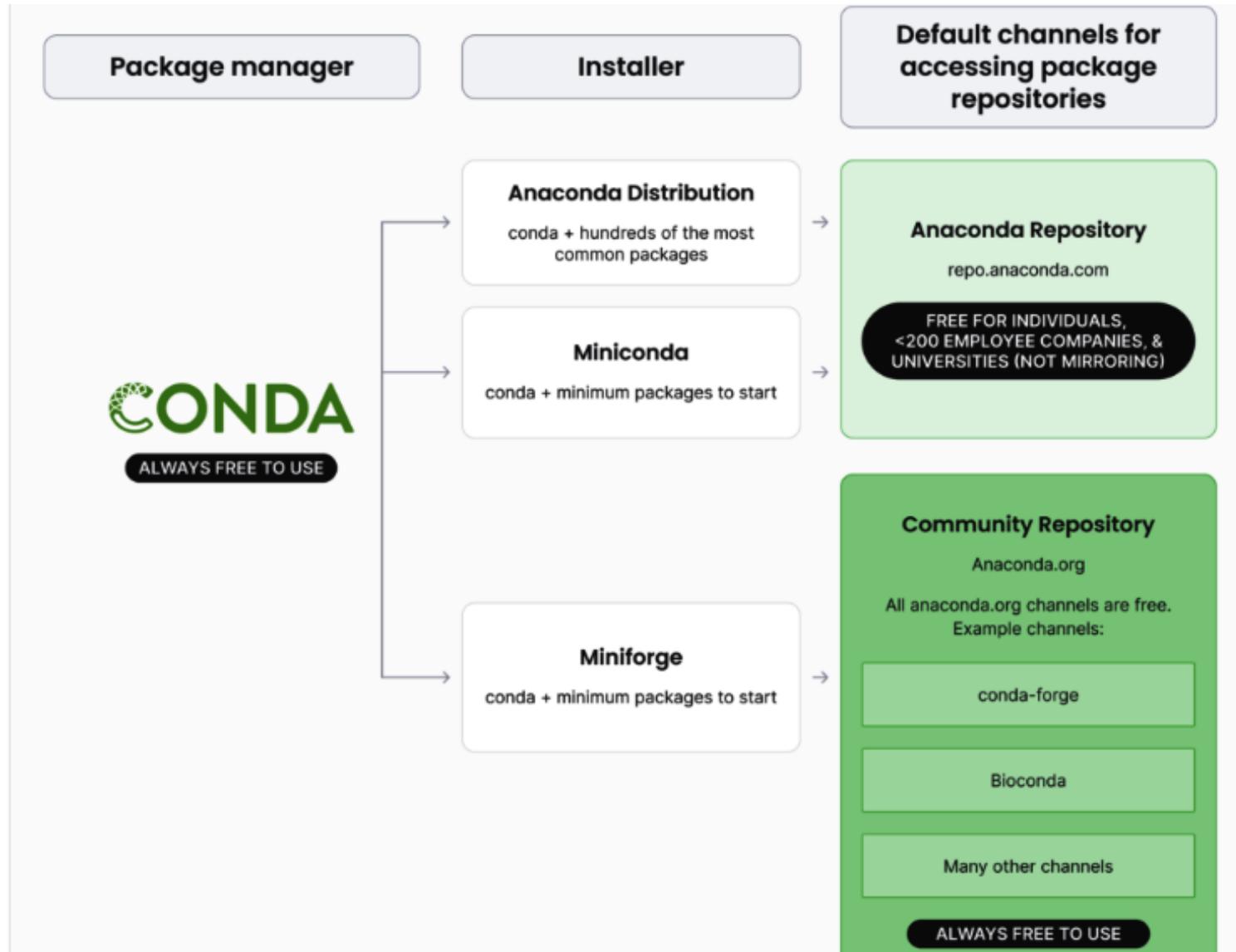
- Ensures smooth package management across different platforms
- Prevents version conflicts with isolated environments
- Optimized for performance and scalability in scientific applications

Anaconda vs Miniconda vs Conda



- Conda: Open-source package manager
- 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
- 2) wulver
- 3) slurm/wulver
- 4) null
- 5) Miniforge3/24.1.2-0

What is Miniforge

\$ module whatis Miniforge3

Miniforge3/24.1.2-0 : Description: Miniforge is a free minimal installer for conda and Mamba specific to conda-forge.

Miniforge3/24.1.2-0 : Homepage: <https://github.com/conda-forge/miniforge>

Miniforge3/24.1.2-0 : 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 : /mmfs1/apps/easybuild/software/Miniforge3/24.1.2-0/.condarc
  conda version    : 24.1.2
  conda-build version : not installed
  python version   : 3.10.14.final.0
  solver          : libmamba (default)
virtual packages  : __archspec=1=skylake_avx512
                  __conda=24.1.2=0
                  __glibc=2.28=0
                  __linux=4.18.0=0
                  __unix=0=0
base environment  : /mmfs1/apps/easybuild/software/Miniforge3/24.1.2-0 (read only)
conda av data dir : /mmfs1/apps/easybuild/software/Miniforge3/24.1.2-0/etc/conda
conda av metadata url : None
  channel URLs    : https://conda.anaconda.org/conda-forge/linux-64
                  https://conda.anaconda.org/conda-forge/noarch
  package cache   : /mmfs1/apps/easybuild/software/Miniforge3/24.1.2-0/pkgs
                  /home/g07396/.conda/pkgs
  envs directories : /home/g07396/.conda/envs
                  /mmfs1/apps/easybuild/software/Miniforge3/24.1.2-0/envs
  platform        : linux-64
  user-agent      : conda/24.1.2 requests/2.31.0 CPython/3.10.14 Linux/4.18.0-372.26.1.el8_6.x86_64 rhel/8.6
glibc/2.28 solver/libmamba conda-libmamba-solver/24.1.0 libmambapy/1.5.7
  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 conda-forge
```

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

```
$conda config --append channels conda-forge
```

This would add the conda-forge to the end of the channel list, giving it the lowest priority.

Conda on HPC

- Introduction to Conda
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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.

Ease of Use

- User-friendly commands for project management.

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].yaml</code> |
| Import environment from YAML: | <code>conda env create -f [environment_name].yaml</code> |
| Import environment to different location: | <code>conda env create -f [environment_name].yaml -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

```
[n0088:~ hz3$ conda info --envs
# conda environments:
#
base                /apps/easybuild/software/Anaconda3/2023.09-0
my_env              /home/hz3/.conda/envs/my_env
tensorflow           /home/hz3/.conda/envs/tensorflow
tf                  /home/hz3/.conda/envs/tf
tf-gpu              /home/hz3/.conda/envs/tf-gpu
                    /project/hpcadmins/hz3/conda_env/my_env
```

```
[n0088:~ hz3$ conda env list
# conda environments:
#
base                /apps/easybuild/software/Anaconda3/2023.09-0
my_env              /home/hz3/.conda/envs/my_env
tensorflow           /home/hz3/.conda/envs/tensorflow
tf                  /home/hz3/.conda/envs/tf
tf-gpu              /home/hz3/.conda/envs/tf-gpu
                    /project/hpcadmins/hz3/conda_env/my_env
```

Conda on HPC

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- **Conda packages**
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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:

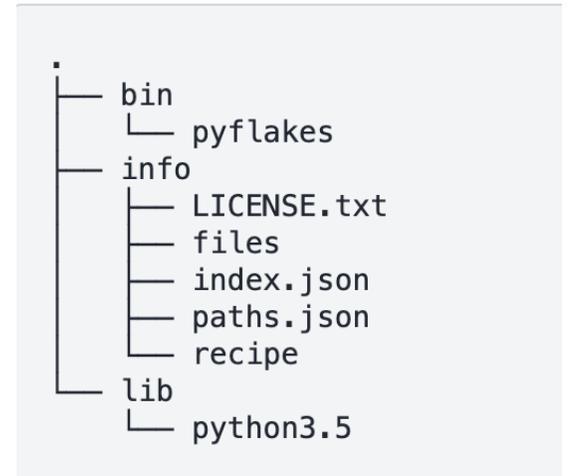
\$ **conda list -n env_name or conda list -p /path/to/environment**

Search for a Package:

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

Check for Specific Package Installation:

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



List packages in all environments

```
[n0088:~ hz3$ conda list
# packages in environment at /apps/easybuild/software/Anaconda3/2023.09-0:
#
# Name                               Version                               Build                               Channel
_anaconda_depends                    2023.09                               py311_mkl_1
_libgcc_mutex                         0.1                                    main
_openmp_mutex                         5.1                                    1_gnu
abseil-cpp                            20211102.0                             hd4dd3e8_0
aiobotocore                           2.5.0                                  py311h06a4308_0
aiofiles                              22.1.0                                 py311h06a4308_0
aiohttp                               3.8.5                                  py311h5eee18b_0
aioitertools                          0.7.1                                  pyhd3eb1b0_0
aiosignal                             1.2.0                                  pyhd3eb1b0_0
aiosqlite                             0.18.0                                 py311h06a4308_0
alabaster                             0.7.12                                 pyhd3eb1b0_0
anaconda-anon-usage                   0.4.2                                  py311hfc0e8ea_0
anaconda-catalogs                    0.2.0                                  py311h06a4308_0
anaconda-client                       1.12.1                                 py311h06a4308_0
anaconda-cloud-auth                  0.1.3                                  py311h06a4308_0
anaconda-navigator                   2.5.0                                  py311h06a4308_0
anaconda-project                     0.11.1                                 py311h06a4308_0
```

List packages in an environment

```
[n0088:~ hz3$ conda list -n my_env
# packages in environment at /home/hz3/.conda/envs/my_env:
#
# Name                Version                Build                Channel
_libgcc_mutex        0.1                    conda_forge         conda-forge
_openmp_mutex        4.5                    2_gnu               conda-forge
alm                   2.0.0_dev.2           py312h63811a6_8     conda-forge
blas                  1.0                    mkl                  conda-forge
bzip2                 1.0.8                  h7b6447c_0          conda-forge
ca-certificates      2024.2.2               hbcca054_0          conda-forge
expat                 2.5.0                  h6a678d5_0          conda-forge
icu                   73.2                   h59595ed_0          conda-forge
intel-openmp         2023.1.0               hdb19cb5_46306     conda-forge
ld_impl_linux-64    2.38                   h1181459_1          conda-forge
libblas               3.9.0                  1_h86c2bf4_netlib   conda-forge
libboost              1.82.0                 h6fcfa73_6          conda-forge
libboost-python     1.82.0                 py312hfb10629_6     conda-forge
libexpat              2.5.0                  hcb278e6_1          conda-forge
libffi                3.4.4                  h6a678d5_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`

Mamba

Mamba is a reimplementaion 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

- Introduction to Conda
- Conda environment
- 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 yml 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/Miniforge3/24.1.2-0
 - * /project/\$GROUP/\$USER/conda_env/my_env

Updating a Conda environment

When to update your conda environment?

- One of your core dependencies just released a new version
- You need an additional package for data analysis (add a new dependency).
- You have found a better visualization package and no longer need to old visualization package

Update the contents of your environment.yml file and run the following command:

```
$ conda env update --file environment.yml --prune
```

- **--prune** option tells Conda to remove any dependencies that are no longer required from the environment

Best practices

- **Use interactive sessions on a compute node**

- Use an interactive session on a compute node to install software with Conda
- **`$ srun -p general -n 1 --qos=standard --account=$PI_ucid --mem-per-cpu=2G --time=59:00 --pty bash`** *#modify srun options as desired*
- **`$ 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. Using your home directory can fill its limited space.
- Managing Conda Cache and changing the default caching behavior

- **Avoid installing packages into your base Conda environment**

Configuring Conda Package Cache

Default Location: **\$HOME/.conda/pkg**s

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

Verify Change: **conda info**

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 --user multiregex
```

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

- ⚠ 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

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

# Example - install tensorflow-gpu

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

```
$conda activate tensorflow
```

```
$conda install -c anaconda tensorflow-gpu numpy=1.21.6
```

Simple TensorFlow test program to make sure the virtual env can access a GPU.

 [tf.gpu.test.py](#) >

 [Slurm script to submit the job](#) >

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

# 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>

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