

Commands for working on the cluster:

Linux commands:

ls – list the folders in the current working directory

cd \$PATH – change directory

pwd – prints the working directory

cp -r \$SOURCE \$TARGET – for copying any file or folder from one path to another locally or within the server

mv \$SOURCE \$TARGET – moving a file/folder from one path to another locally or within the server

Communicating with the server

ssh – for logging in to the server

e.g.: **ssh yadav@cluster-g.math.tu-berlin.de**

scp – for securely copying the files/folders from local machine to remote server and vice-versa (**scp -r \$SOURCE \$TARGET**)

e.g.: **scp -r /home/vikas/PINN/Cases/Kornilov_Adiabatic/example.py yadav@cluster-g.math.tu-berlin.de:/homes/smta/yadav/Vikas/**

After logging in to the server, you can activate your conda environment by the following command:

conda activate /work/yadav/temp-env/

In case you do want to create your own environment, you can also do that by the following command:

conda create -p /work/yadav/\$ENV_NAME python=3.10.12

You can then activate your new conda environment.

Once you are in your preferred environment, you can install the packages using the following command:

conda install \$PACKAGE_NAME

e.g: **conda install pandas** or **pip install pandas**

In case you want to deactivate the environment, you can do that by,

conda deactivate

I think you will not have to install

Job submission commands

Following are the most common commands that you would have to use in Slurm:

sbatch #JOB_FILE - For submitting the job, e.g., **sbatch job.sh** (Description of job file already in the comments of the job file)

squeue -me - For checking the job status (PD – pending, R – running)

scancel #JOB_ID - For canceling the job