

## Useful links

- MARCC Homepage
- The Queuing System
- Storage Guidelines
- Slurm Array jobs

# The setup

```
1 git clone https://github.com/neurodata/marcc_examples.git
```

```
2 ml python/3.8
```

```
3 python -m venv ~/env_examples
```

```
4 pip install -r marcc_examples/requirements.txt
```



<i>skrf.py</i>	The main python script that takes parameters such as dataset ID, run ID, n_jobs, output directory.
<i>workerScript.scr</i>	The slurm script used for each individual job.
<i>cleanUp.scr</i>	The script that runs the cleanup job/s.
<i>MASTER.scr</i>	The master script that sets up the job array and sets cleanUp as a dependant job to be run after all exit cleanly.