

The University of Queensland

Research Computing Centre

Batch Computing using SLURM

Document Status

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Background

The Wiener GPU based cluster has adopted SLURM as the batch system because a number of features it has for supporting GPU computing.

This guide is still under construction.

In the meantime, you could refer to the

- SLURM documentation available at the [SchedMD website](#)
- SLURM Quick Start Tutorial at the [Consortium des Équipements de Calcul Intensif](#)
- SLURM Tutorial at [NYU](#)
- our very own Wiener User Guide (linked on the right ... just over there)

Command Subset Comparison

To do this	SLURM	PBS Pro
Check queue status	<code>squeue</code>	<code>qstat</code>
Submit a batch script	<code>sbatch</code>	<code>qsub</code>
Delete a job	<code>scancel</code>	<code>qdel</code>
Node information	<code>sinfo</code>	<code>pbsnodes</code>
Start an interactive session	<code>srun ... --pty bash</code>	<code>qsub -I</code>
Denote a setting in a job script	<code>#SBATCH</code>	<code>#PBS</code>