

PowerIDE User Guide

PowerIDE provides interactive access to the HPC cluster through a web browser. Run Jupyter notebooks, VS Code, and RStudio directly on compute nodes without needing SSH access.

Access PowerIDE at: <https://poweride.tau.ac.il/jupyter>

Getting Started

Log in with your TAU university credentials (same as email and other university services).

After logging in, click **Start My Server**. You'll see a Server Options form to configure your compute resources.

When you start your server, PowerIDE submits a Slurm job to the PowerSlurm cluster. Your session runs on a compute node — not on the PowerIDE server itself. This means:

- You get dedicated resources (CPUs, memory, GPUs) on a compute node
- Your job runs through the same Slurm scheduler as other HPC jobs
- Your session will queue if the cluster is busy

Configuring Resources

Partition

Select which partition to run on. The dropdown shows only partitions you have access to. Common options:

- `power-general-shared-pool` — general purpose computing
- `gpu-general-pool` — GPU-enabled nodes

Check with your PI or HPC admin if unsure which partition to use.

QOS

Controls priority and resource limits. **Default (owner)** is usually the right choice. Only valid QOS options for your selected partition are shown.

GPUs

Appears only when a GPU partition is selected. Specify how many GPUs you need (0 if none).

Time (D-HH:MM:SS)

How long your session should run. Default: . Your session is terminated when time runs out — save your work regularly.

- — 2.5 hours
- — 1 day and 12 hours

CPUs per task

Default: 1. Increase for multi-threaded code.

Memory

Default: 1G. Examples: , , . Start small and increase if needed — over-requesting delays job start.

Working Directory

Default: your home directory. Change to your project directory to save navigation time after launch.

Stdout / Stderr Directory

Where job logs are written. Default (home directory) is fine for most users.

Starting Your Session

Click the orange **Start** button. PowerIDE submits a Slurm job and shows a progress page. Once a compute node is available (usually 10–60 seconds), you're automatically redirected to JupyterLab.

If the cluster is busy, you can close the browser and come back — your session will start when resources are available.

Using JupyterLab

- **Left sidebar** — file browser, running kernels, extensions
- **Main area** — notebooks, text files, terminals
- **+** button — opens the launcher for new tools

Common tasks:

- **New notebook** — click **+** → choose a Python kernel
- **Terminal** — click **+** → click Terminal (bash shell on the compute node)
- **Upload files** — drag and drop into the file browser
- **Download files** — right-click file → Download

Using VS Code

PowerIDE includes VS Code running in your browser:

1. Click **+** to open the launcher
2. Click the **VS Code** icon
3. VS Code opens in a new tab with access to all your files and the same resources as JupyterLab

Using RStudio

RStudio runs as a separate service on a dedicated compute node. It has its own launch form with R-specific options:

1. Click **+** to open the launcher
2. Click the **RStudio** icon
3. Fill in the resource form and select your R environment
4. Click **Start** — RStudio opens in a new tab once the job is running

R Environment

Select the R environment to load. Each environment is a named module (e.g. `webR-genomics-2024`) with R and a pre-installed set of packages. Contact HPC support if you need a package that isn't available.

R Library Path (optional)

If you have a personal R package library installed in a directory on the cluster, enter its full path here (e.g. `/home/user/R/library`). R will search this directory first, before the environment's default library.

Stopping RStudio

Use the **Stop** button in the PowerIDE topbar to terminate your RStudio job. **Do not use File → Quit Session** — that ends the R session but leaves the Slurm job running, continuing to consume resources.

Python Environments

PowerIDE provides one default kernel: **Python 3.12 (Base)**.

You can register your own conda/mamba environments as kernels:

```
module load mamba/mamba-2.1.1
mamba create -n my-project python=3.11 pandas matplotlib
mamba activate my-project
mamba install ipykernel

# Register as kernel (only visible to you)
python -m ipykernel install --user --name my-project --display-name "My Project (Python 3.11)"
```

Refresh your browser — the new kernel appears in the launcher. To remove a kernel:

```
jupyter kernelspec uninstall kernel-name
```

Stopping Your Server

Always stop your server when done to free resources for others.

- **From JupyterLab** — File → Hub Control Panel → Stop My Server
- **From PowerIDE home** — navigate to <https://poweride.tau.ac.il/jupyter/hub/home> → Stop My Server
- **VS Code / RStudio** — use the **Stop** button in the PowerIDE topbar

Best Practices

- Request only what you need — over-requesting delays your job and others
- Set a realistic time limit; restart if you need more
- Only request GPUs if your code actually uses them
- Store large datasets in scratch space, not your home directory
- Use Git for code — not for large data files

Created 2026-06-14 08:31:40 UTC by levk

Updated 2026-06-18 13:02:09 UTC by levk