

Access to Cluster

The following groups are eligible for scientific computing on the HPC cluster:

- Scientists at the TU Berlin
- Students writing their master's/bachelor's thesis at the TU Berlin

The HPC cluster is not intended for personal projects or educational purposes (e.g. exercises, labs). Use of the cluster is therefore linked to a chair or research project at the TU Berlin.

Prerequisites

The prerequisite for access is a [TUB account](#), which is issued to all members of the university at the start of employment via the provisioning procedure.

For cluster access, [a team](#) must be created for each department by the corresponding role administrator, in which all access permissions are maintained. The [group function \("Gruppenfunktion"\) must be activated](#) for said team.

Once the group name (assigned when the group function is added) along with the department's organisational domain has been reported back to <mailto:it-support@tu-berlin.de> with the subject line 'HPC:Team assignment', the group will be approved by us and all team members will then be synchronised with our SLURM database. This synchronisation is done by automated scripts and can therefore take up to an hour after access has been granted.

Students who need the cluster for their thesis are granted access through the chair where the thesis is located.

Login

The login to the cluster is done via SSH (Linux/Mac build-in) using the virtual address `gateway.hpc.tu-berlin.de`. This address can be reached from the TUB subnets. The TUB account is used as the user name.

Within the TUB networks, login is possible as follows:

```
ssh <TUB-Account>@gateway.hpc.tu-berlin.de
```

Outside the TUB networks, the connection is either established via a VPN tunnel [Cisco AnyConnect](#) or the SSH jump host is used:

```
ssh <TUB-Account>@sshgate.tu-berlin.de
<TUB-Account>@sshgate.tu-berlin.de's password:

-bash-4.2$ ssh gateway.hpc.tu-berlin.de
<TUB-Account>@gateway.hpc.tu-berlin.de's password:

-bash-4.2$
```

Windows workstations

Windows Subsystem for Linux (WSL) or an additional software is required for access under Windows. For example, [Putty](#) or [MoabTerm](#) can be used.

Access to the cluster storage space

You can access your own files that are located on the cluster storage space (beegfs) using "scp" or "rsync". This makes it possible, for example, to copy the files directly to your own computer or from there to the cluster. For direct access, i.e. without having to make a copy, [sshfs](#) can be used.

Under Windows, an SFTP client can be used to transfer files. [Here](#) a short explanation.

From:
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Last update: **2025/03/24 11:07**

