Stony Brook University

Alejandro Gil Gomez <alejandro.gilgomez@stonybrook.edu>

Re: SeaWulf access [#AR850034]

Support <support@iacs.supportsystem.com>   Mon, Dec 9, 2019 at 12:47 PM
To: Alejandro Gil Gomez <alejandro.gilgomez@stonybrook.edu>
Cc: Joshua Rest <joshua.rest@stonybrook.edu>

Dear Alejandro Gil Gomez,

Hi Alejandro,

Your account on the SeaWulf cluster is ready for use.

SeaWulf uses the campus NetID and NetID password for authentication.
If you need to reset your NetID password, please visit this site.

You may access the SeaWulf cluster login node using the command-line from any modern workstation via secure shell (SSH). In Linux or MacOSX, simply `ssh` to the SeaWulf login node by issuing the command:

`ssh -X` agilgomez@login.seawulf.stonybrook.edu

Your home folder is located in: `/gpfs/home/agilgomez` [Quota 20 GB]
Your scratch space is located in: `/gpfs/scratch/agilgomez` [Quota 20 TB]

NOTE: You can run jobs from your `home` directory, however due to the 20 GB quota, it is recommended that you run jobs from the `scratch` space, and save the results in your `home` directory. Files older than 1 month will be purged from the `scratch` directory, which is **not** backed up. By default only you and the System Administrators have access to the contents of your `home` and `scratch` directories.

For more information on the SeaWulf storage please see here.

In order to submit jobs on the cluster, first load the required modules via:

`module load shared slurm`

To see a list of modules available for use, please type:

`module avail`

For more information on modules and software packages available on the cluster, please visit our HPC FAQ site.

Lastly, please report any issues and suggestions via our support portal.

Regards,
Joey

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