

HPC data handling

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Topics

- Data staging
 - Using work
 - Using \$SCRATCH
 - Using \$TMP
- Check whether your job can run in parallel
 - Simple way using top
- Ask for the correct resources
- Make sure environment can be reproduced
- Investigate during the job run

Different locations

- To find out the mounts
 - `df -h`
- `/cluster` is slow
 - `$HOME` is on cluster
- `/work` is fast
 - `/work/users/<USERNAME>`
 - `$SCRATCH` is on work
 - Temp directory during a job is on work
- `$LOCALTMP` is the fastest
 - IT is a local disk on a compute node.
 - Not shared between nodes
 - Very good when there a millions of very small files to be processed.

Check the available directories

- Test job
 - mkdir \$HOME/HPC_NOV2018
 - cp /cluster/teaching/abel_tutorial/NOV2018/location_test.slurm \$HOME/HPC_NOV2018
 - cd \$HOME/HPC_NOV2018
 - chmod +w location_test.slurm
 - nano location_test.slurm
 - Edit the project account
 - sbatch location_test.slurm

When there are hundreds of thousands files

- Test job
 - mkdir /work/users/\$USER/HPC_NOV2018
 - If you are a guest then mkdir \$HOME/HPC_NOV2018
 - cp /cluster/teaching/abel_tutorial/NOV2018/millionfiles.slurm /work/users/\$USER/HPC_NOV2018
 - cd /work/users/\$USER/HPC_NOV2018
 - chmod +w millionfiles.slurm
 - nano millionfiles.slurm
 - Edit the project account
 - sbatch millionfiles.slurm

When there are hundreds of thousands jobs

- Create an archive (no compression)
 - `tar -cvf <ARCHIVE_NM> <FILES>`
 - `tar -cvf file.tar *.txt`
- List content
 - `tar -tvf <ARCHIVE_NM>`
- Append a file
 - `tar --append --file <ARCHIVE_NM> <NEW_FILES>`
 - `tar --append --file files.tar.gz 1.txt`
- Extract all
 - `tar -xvzf <ARCHIVE_NM>`
- Extract one
 - `tar -xvf <ARCHIVE_NM> <FILES>`

Big file and Isof

- Test job
 - `mkdir /work/users/$USER/HPC_NOV2018`
 - `cp /cluster/teaching/abel_tutorial/NOV2018/file_IO* /work/users/$USER/HPC_NOV2018`
 - `cd /work/users/$USER/HPC_NOV2018`
 - `chmod +w file_IO.slurm`
 - `nano file_IO.slurm`
 - Edit the project account

Big file and Isov

- Test job
 - sbatch file_IO.slurm
 - squeue -u \$USER
 - When the job starts get the compute node address
 - Login to the compute node
 - ssh cx.xx
 - top -u \$USER
 - Find PI and investigate with Isov

sacct

- Investigate resource usage after the job is finished
- `sacct -j <JOB_ID>`
- `sacct -j <JOB_ID> -o all`

Qlogin

- `qlogin --account=<ACCOUNT> --ntasks=4
--mem-per-cpu=2G --time=1:00:00`
- `qlogin --account=<ACCOUNT> --partition=accel
--ntasks=8 --mem-per-cpu=6G --gres=gpu:2
--time=02:00:00`