

Introduction to SCC & fMRIPrep

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Resources

- Tutorial Slides available:
 - http://rcs.bu.edu/examples/imaging
- Questions, Problems, Need Help?
 - help@scc.bu.edu
 - mhorn@bu.edu



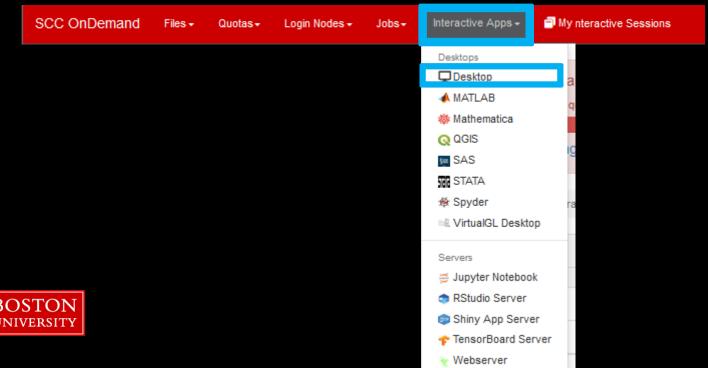
Outline

- SCC
- File Structure
- File Permissions
- Software
- Batch Jobs
- BIDS
- fMRIPrep



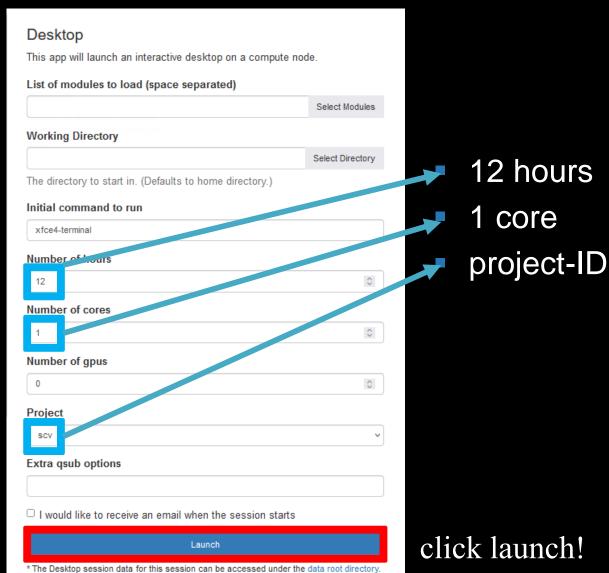
Open an OnDemand session

- Go to: scc-ondemand.bu.edu
- **Interactive Apps**
- Desktop



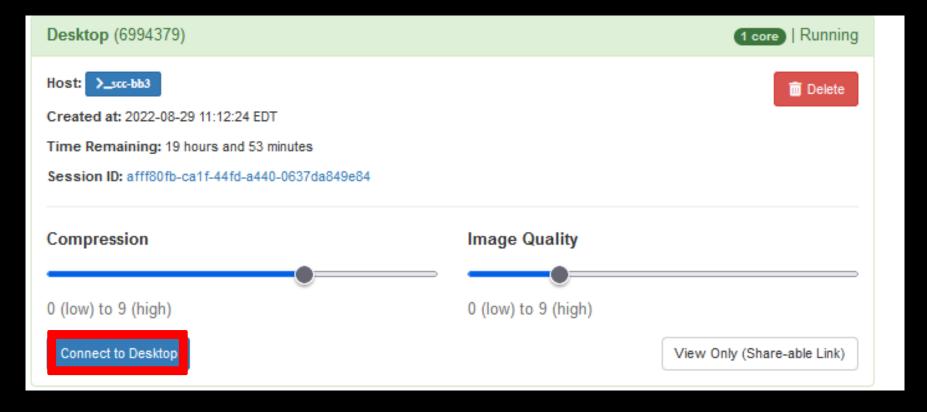


Open an OnDemand session





Open an OnDemand session



click Connect to Desktop!

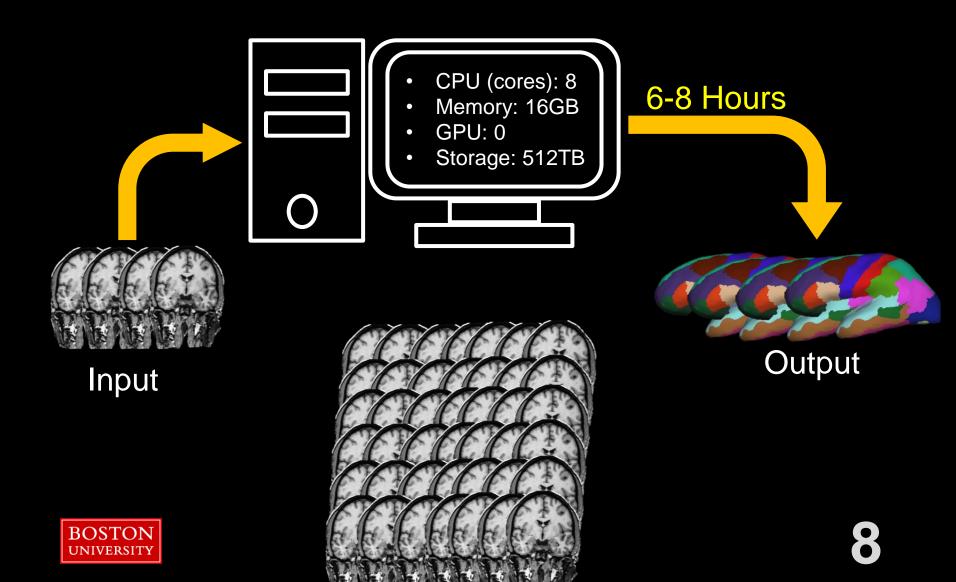


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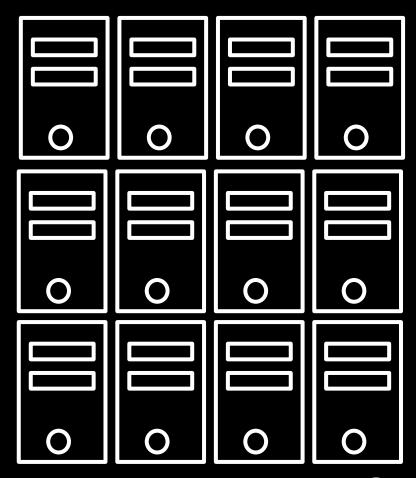


What is the SCC?



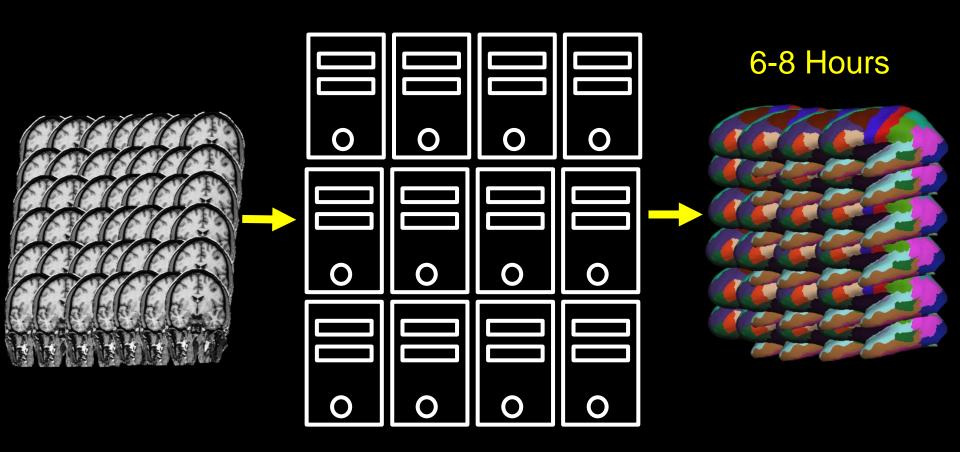
What is the SCC?

- > 900 Computers (nodes)
- > 25,000 CPUs (cores)
- > 100 GPUs
- > 9 PB (storage)

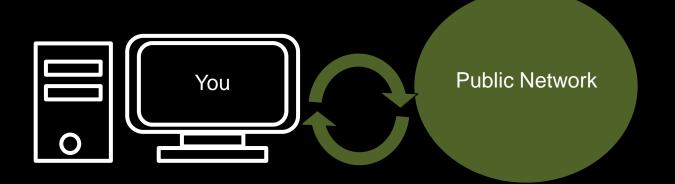




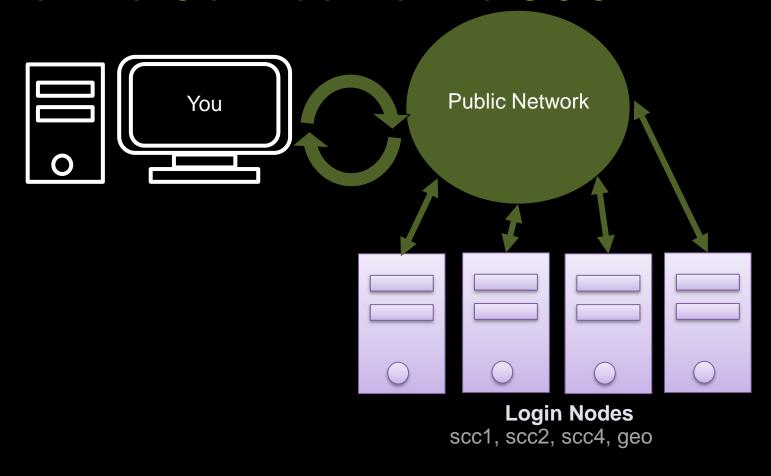
What is the SCC?



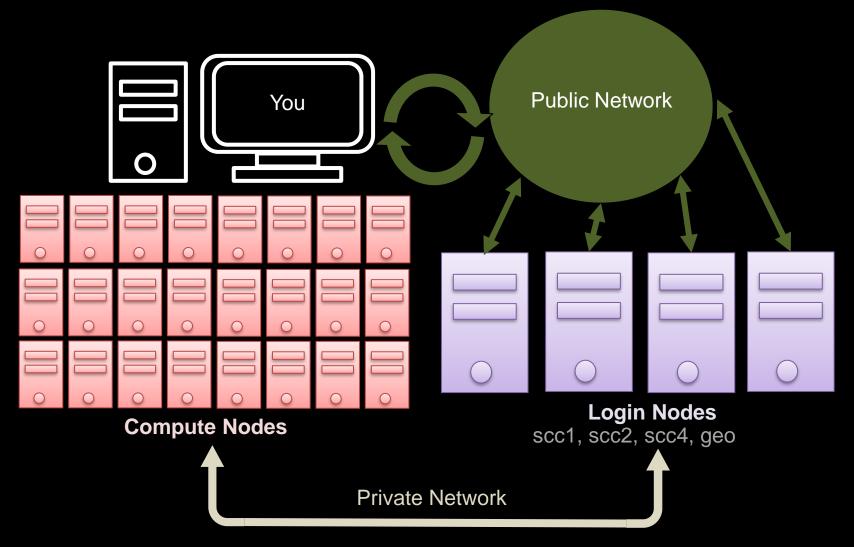




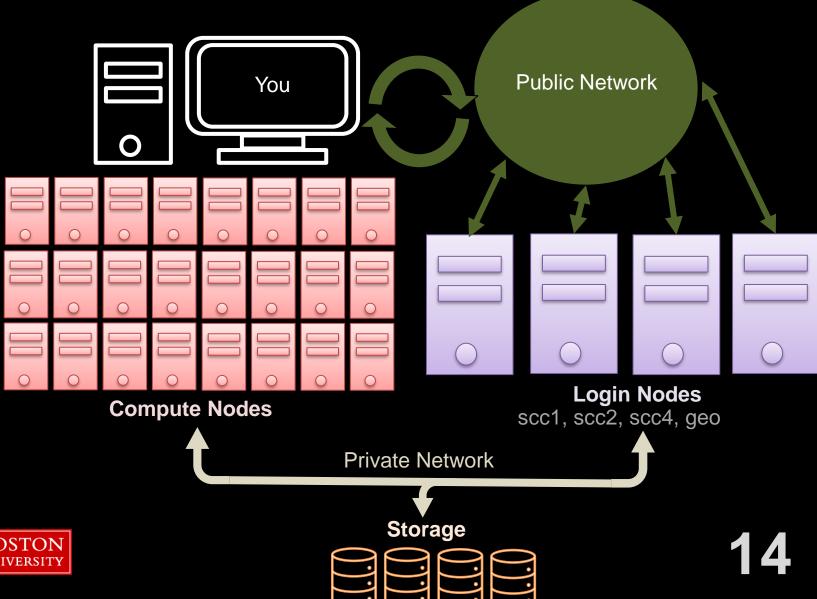




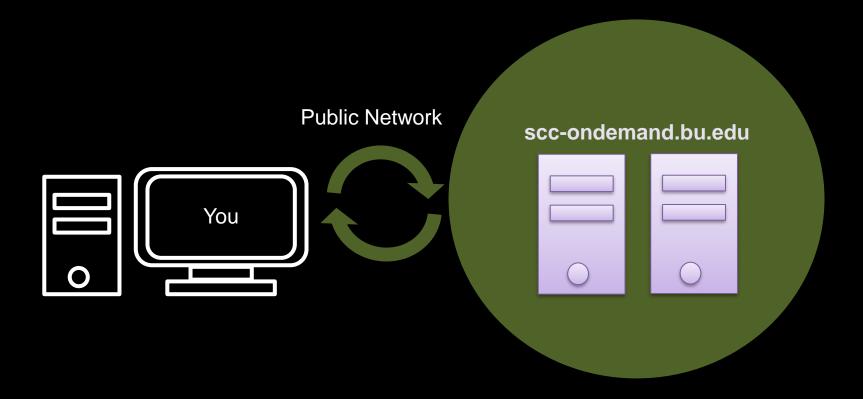














Outline

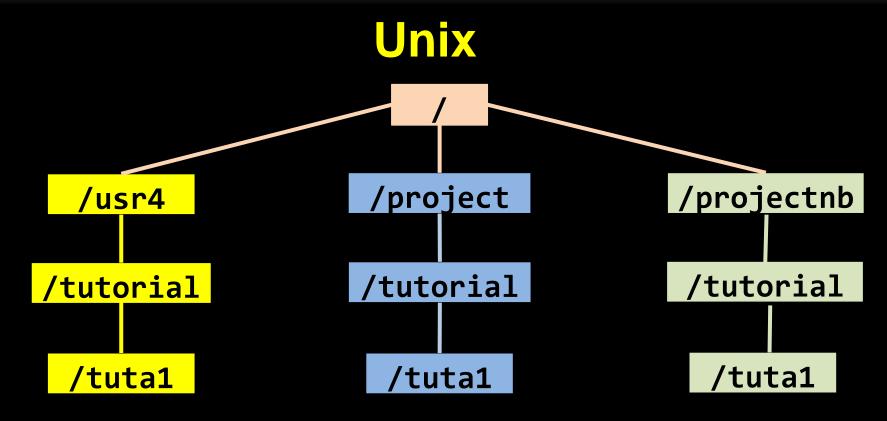
- SCC
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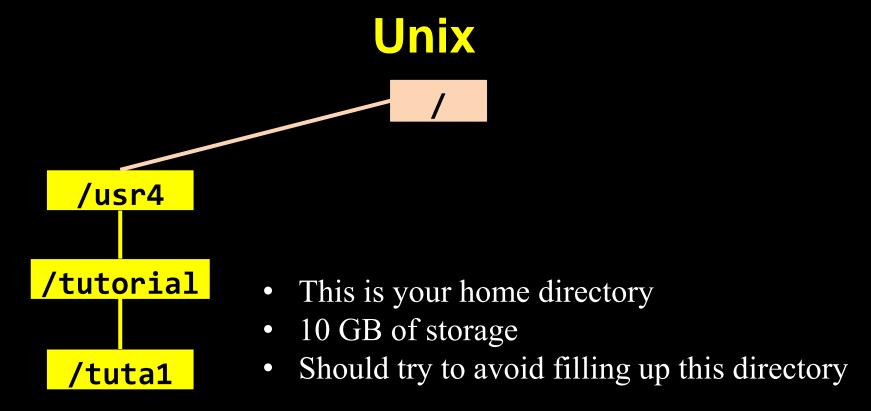
```
[tuta1@scc-v01 ~]$ pwd
/usr4/tutorial/tuta1
```

Shows the directory you are in, with the full path

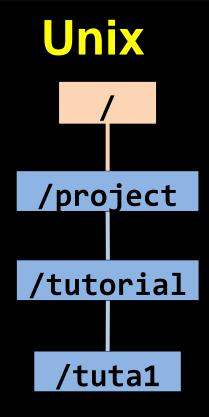












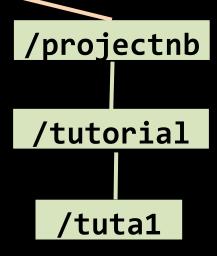
- This is a backed-up project storage
- Backup in a separate location incase of physical damage
- Default 50GB of space, can increase to 200GB







- This is a **NOT** backed-up project storage
- Local snapshots
- Default 50GB of space, can increase to 800GB





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[tuta1@scc-v01 ~]\$ cd /projectnb/ne742

cd = change directory



[tuta1@scc-v01 ~]\$ mkdir mhorn

swap in your username

mkdir = make a directory

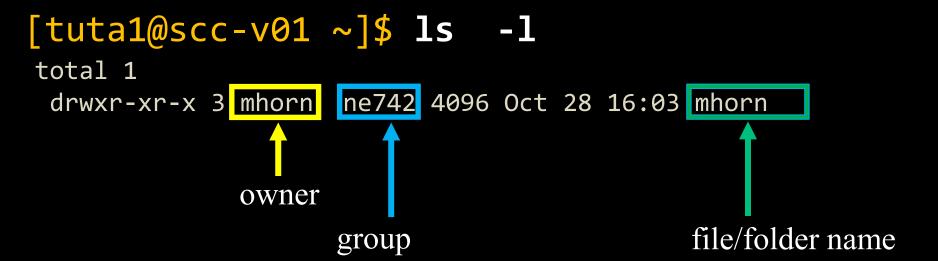


```
[tuta1@scc-v01 \sim]$ ls -1
```

total 1
drwxr-xr-x 3 mhorn ne742 4096 Oct 28 16:03 mhorn

ls = list contents





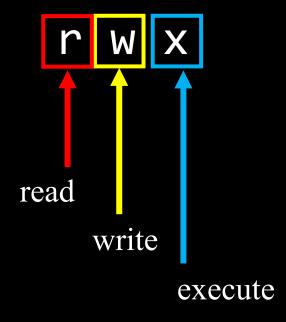
ls = list contents



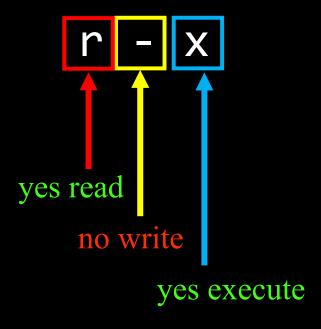
```
[tuta1@scc-v01 ~]$ ls -l
total 1
drwxr-xr-x 3 mhorn ne742 4096 Oct 28 16:03 mhorn
permissions
```

ls = list contents

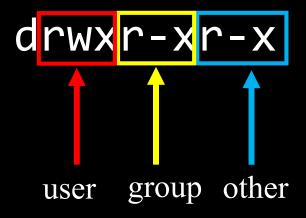






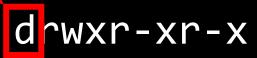








```
[tuta1@scc-v01 ~]$ ls -l
total 1
drwxr-xr-x 3 mhorn ne742 4096 Oct 28 16:03 mhorn
```



Object Type

- d = directory
- -= file



```
[tuta1@scc-v01 ~]$ ls -l
total 1
drwxr-xr-x 3 mhorn ne742 4096 Oct 28 16:03 mhorn
```



User/Owner Permissions:

- mhorn is owner
- currently all enabled



[tuta1@scc-v01
$$\sim$$
]\$ ls -1

total 1

drwxr-xr-x 3 mhorn ne742 4096 Oct 28 16:03 mhorn



Group Permissions:

- ne742 is group
- read and execute are enabled
- write is not enabled



Other/World Permissions:

- read and execute are enabled
- write is not enabled



```
[tuta1@scc-v01 ~]$ ls -l
total 1
drwxr-xr-x 3 mhorn ne742 4096 Oct 28 16:03 mhorn
```



How can we change permissions?

[tuta1@scc-v01 ~]\$ chmod g+w mhorn

To group add write permissions



Unix



Outline

- SCC
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```
[tuta1@scc-v01 tuta1]$ module avail less
```

```
ctrl+z to quit "less"
```



[tuta1@scc-v01 tuta1]\$ module avail dcm2niix

ctrl+z to quit "less"



[tuta1@scc-v01 tuta1]\$ module load dcm2niix

Module system to load software packages



```
[tuta1@scc-v01 tuta1]$ module list
```

```
Currently Loaded Modules:
1) dcm2niix/1.0.20220720
```

Module system to load software packages



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- Up to this point, we've been doing things Interactively
 - request resources
 - connect to the remote session
 - program via interface (terminal &/or GUI's)

- Can process Non-Interactively
 - easier to scale
 - flexible submit and forget
 - more efficient resource management
 - can submit large number (batches) of jobs



Submit from command line

qsub [options] command [arguments]

Submit with a jobscript file:

qsub script.qsub



qsub options

	General Directives	
Directive	Descrip	
-I h_rt=hh:mm:ss	Hard run time limit in hh:mm:ss format. The d	1
-P project_name	Project to which this jobs is to be assigned. Th associated with any Med.Campus project.	-
-N job_name	Specifies the job name. The default is the scrip	
-o outputfile	File name for the stdout output of the job.	
-e errfile	File name for the stderr output of the job.	
-j y	Merge the error and output stream files into a s	2
-m b e a s n	Controls when the batch system sends email t job begins (b), ends (e), is aborted (a), is susp	
-M user_email	Overwrites the default email address used to s	-
-V	All current environment variables should be ex	
-v env=value	Set the runtime environment variable env to va	-
-hold_jid job_list	Setup job dependency list. job list is a con names which must complete before this job ca Usage for more information.	-

	Directives to request SCC resources
Directive	Description
-I h_rt=hh:mm:ss	Hard run time limit in hh:mm:ss format. The default is 12 hours.
-I mem_per_core=#G	Request a node that has at least this amount of memory per core. Recommended choices are: 3G, 4G, 6G, 8G, 12G, 16G, 18G and 28G
-pe omp N	Request multiple slots for Shared Memory applications (OpenMP, pthread). This option can also be used to reserve a larger amount of memory for the application. <i>N</i> can vary. Currently, to request multiple cores on SCC's shared nodes, we recommend to select 1-4, 8, 16, 28, or 36 cores.
-pe mpi_#_tasks_per_node //	Select multiple nodes for an MPI job. Number of tasks can be 4, 8, 12, 16, or 28 and N must be a multiple of this value. See Running Parallel Batch Jobs for more information.
-t N	Submit an Array Job with N tasks. N can be up to 75,000. For more information see Array Jobs
-l cpu_arch=ARCH	Select a processor architecture (broadwell, ivybridge, cascadelake). See Technical Summary for all available choices.
-I cpu_type=TYPE	Select a processor type (X5670, X5675, Gold-6132 etc.) See <u>Technical Summary</u> for all available choices.
-l gpus=G	Requests a node with GPUs. <i>G</i> is the number of GPUs. See GPU Computing for more information.
-l gpu_type=GPUMODEL	To see the current list of available GPU models, run <i>qgpus</i> command. See GPU Computing for more information.
-l gpu_c=CAPABILITY	Specify minimum GPU capability. Current choices for <i>CAPABILITY</i> are 3.5, 5.0, 6.0, 7.0, and 8.6
-I gpu_memory=#G	Request a node with a GPU that has 12G, 16G, 24G, 32G, 48G of memory.
-l avx	Request a node that supports AVX and newer <u>CPU instructions</u> . A small number of modules require support for these instructions.



jobscript file

Script Interpreter

Scheduler Directives

Commands to execute

```
#!/bin/bash -1
# Time Limit
#$ -1 h rt=12:00:00
# Project Name
#$ -P ne742
 Job Name
#$ -N testjob
# Load Modules
module load python/3.10.12
# Run the program
python myscript.py
```

Directives to request SCC resources		
Directive	Description	
-I h_rt=hh:mm:ss	Hard run time limit in hh:mm:ss format. The default is 12 hours.	
-I mem_per_core=#G	Request a node that has at least this amount of memory per core. Recommended choices are: 3G, 4G, 6G, 8G, 12G, 16G, 18G and 28G	
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-l gpu_c=CAPABILITY	Specify minimum GPU capability. Current choices for <i>CAPABILITY</i> are 3.5, 5.0, 6.0, 7.0, and 8.6	
-l gpu_memory=#G	Request a node with a GPU that has 12G, 16G, 24G, 32G, 48G of memory.	
-l avx	Request a node that supports AVX and newer <u>CPU instructions</u> . A small number of modules require support for these instructions.	

- All Purpose Nodes:
 - can run single-processor jobs and parallel jobs (≤ 720h)
- Whole Node queues (8, 16, 28, 36 cores):
 - must request whole node (≤ 240h)
- GPU nodes:
 - must request GPU(s) (≤ 48h)
- MPI queues:
 - must request multiple nodes (≤ 120h)
- VirtualGL nodes
 - for interactive graphics jobs

[tuta1@scc-v01 tuta1]\$ qstat -u tuta1

swap in your username

Check status of job



Check status of running job

7238828 0.14071 ood-deskto tuta1 7243957 0.00000 bet.qsub tuta1

[tuta1@scc-v01 tuta1]\$ qstat

```
-u tuta1
r = running
r = 09/12/2022 09:46:27 b@scc-bc2.scc.bu.edu
qw = 09/12/2022 14:47:20
qw = queued
```



Check report of completed jobs

[tuta1@scc-v01 tuta1]\$ qacct -j 596557

```
qname p100
hostname scc-c11.scc.bu.edu
group ozymandias
owner mhorn
project ne742
jobname myjob
jobnumber 551947
qsub time Wed Sep 6 20:08:56 2017
start time Wed Sep 6 20:09:37 2017
end_time Wed Sep 6 23:32:29 2017
granted pe NONE
slots 1
failed 0
exit status 0
cpu 11232.780
mem 611514.460
io 14.138
iow 0.000
maxvmem 71.494G
arid undefined
```



The job should run soon and produce an output file:

```
[tuta1@scc1 tut]$ cat bet.qsub.o1041461
```

There will also be an error file which should be empty:

```
[tuta1@scc1 tut]$ cat bet.qsub.e1041461
```



Get Example Data

cd /projectnb/ne742/students/\$USER

cp /project/scv/examples/imaging/ne742/Data.zip \$PWD

3.
unzip Data.zip



Submit Single FMRIPrep Job

4.
cd Data/scripts

5. qsub fmriprep.qsub



Submit Multi-FMRIPrep Job

6.
qsub fmriprep_m.qsub subjects.txt



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BIDS

Brain Imaging Data Structure

Why?

- To standardize organization of complex data
- Have software develop BIDS.apps to take in datasets
- Avoid lab-specific pipelines
- Facilitate validating datasets and curation
- Facilitate data sharing



BIDS

Brain Imaging Data Structure

What?

- Three main file types in a BIDS dataset:
 - JSON contain metadata as key:value pairs
 - .TSV contain tables of metadata
 - .NII.GZ raw data files for fMRI and MRI data



BIDS

```
BIDS/
    sourcedata
                <DICOMS go here>
    sub-101
               ses-001
                           anat
                                    sub-001_ses-001_run-01_T1w.json
                                    sub-001_ses-001_run-01_T1w.nii.gz
                          – func
                                   sub-001_ses-001_run-01_BOLD.json
                                    sub-101_ses-001_run-01_BOLD.nii.gz
    derivatives
                fmriprep
README
dataset_description.json
participants.tsv
```



BIDS Compliancy

- For BIDS.apps to successfully recognize and import data, you can verify organization
- Not always necessary
- http://bids-standard.github.io/bids-validator/



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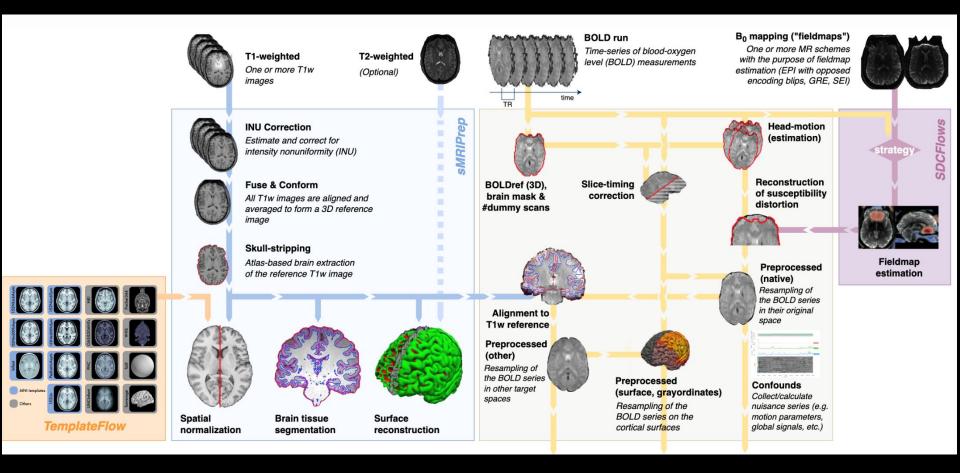
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- Workflow taking principal input images and generates a standardized preprocessed output for analysis
- Uses BIDS standard for input and output
- Customize preprocessing via one command line with arguments
- Generate visual outputs for basic QC analysis

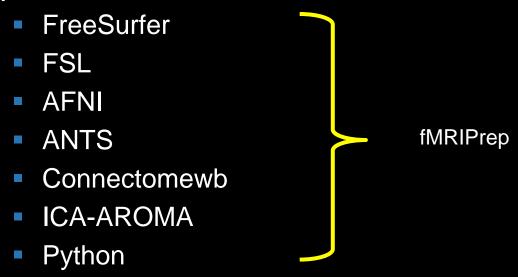


What's fMRIPrep





Adapts multiple software packages into a single standardized process.





fmriprep BIDS BIDS/derivatives participant [options]

command to run fmriprep











Some Useful Options

Performance control

--nprocs : number of cpus for entire process

--omp-nthreads : number of threads per process

--mem : upper bound memory limit for entire process

Workflow configurations

--output-spaces : list spaces i.e. (MNI251Lin6Asym, MNI152Lin2009cAsym)

--bold2t1w-init : DOF when registering BOLD to T1w (6=def, options 6, 9, 12)

--dummy-scans : number of nonsteady-state volumes

--output-spaces : choose your standardized spaces

-w : define where the working directory will save output

Estimating Confounds options

--use-aroma : add ICA_AROMA to your processing steam

--fd-spike-threshold : framewise displacement threshold for flagging frame as an outlier

--dvars-spike-threshold:DVARS threshold for flagging frame as an outlier

FreeSurfer options

--fs-license-file : path to FreeSurfer license key

--fs-subjects-dir : path to existing FreeSurfer directory to import

--fs-no-reconall : disable FreeSurfer preprocessing

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```
fmriprep BIDS BIDS/derivatives participant \
--participant-label $subjid \
-w /scratch/$USER/fmriprep \
--nprocs 8 \
--omp-nthreads 2 \
--mem 64 \
--fs-license-file /share/pkg.8/freesurfer/7.4.1/install/license.txt
```

- command
- required arguments
- option flags
- option arguments



Additional BU RCS Web Resources

- Research Computing Support Pages <u>http://www.bu.edu/tech/support/research/</u>
- Technical Summary of SCC Resources
 http://www.bu.edu/tech/support/research/computing-resources/tech-summary/
- SCC Updates Latest SCC News
 http://www.bu.edu/tech/support/research/whats-happening/updates/
- Code Examples for Popular Software Packages http://scv.bu.edu/examples/



Please open a web browser and:

2. Google "rcs tutorial evaluation"

3. Click 1st link http://scv.bu.edu/survey/tutorial_evaluation.html

4. Fill out the survey!

