

Anatomical analysis with Freesurfer

31082024

WhatsApp group

<https://chat.whatsapp.com/L5qwsHRiBkZ8Z0PfiOhpSb>

NeuroImaging Workshop IIITH
BCCL

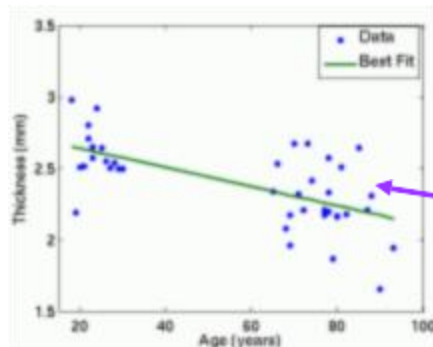
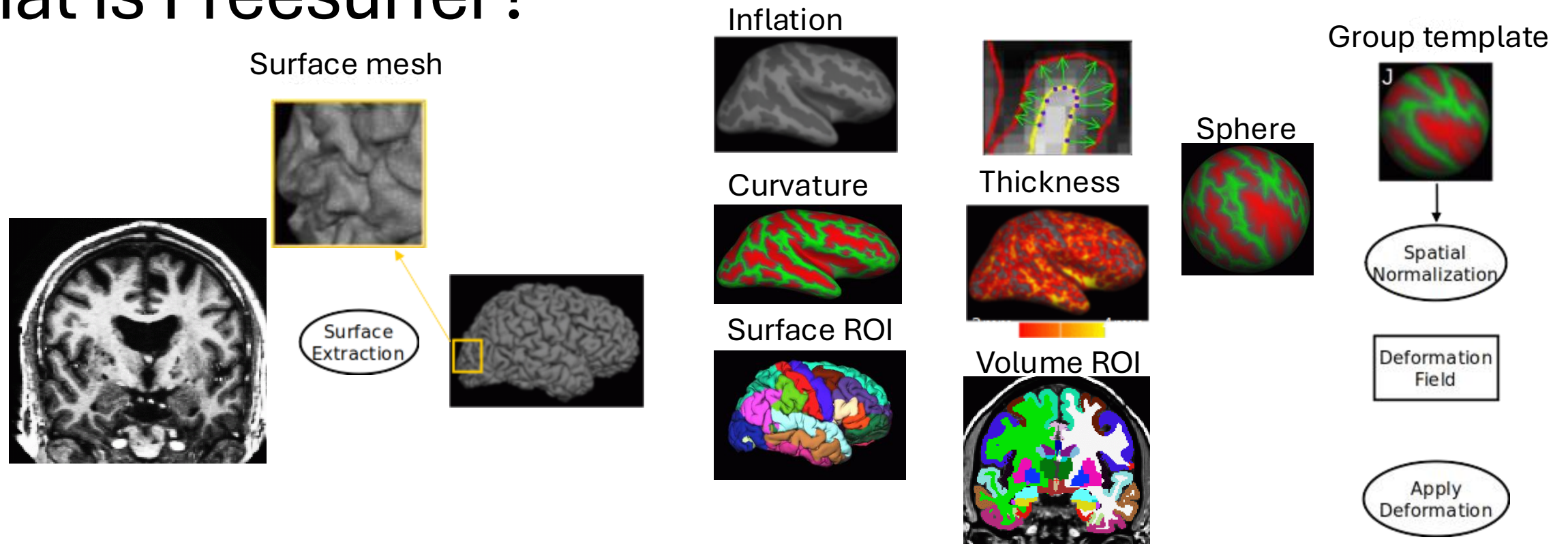
WhatsApp group



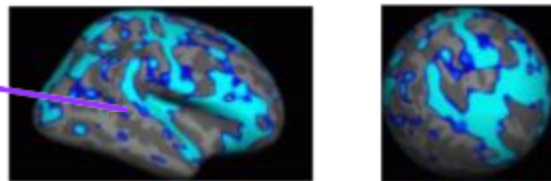
Outline

- What is Freesurfer?
- Recap Freesurfer terms
- Recon-all
 - Components of each step in recon-all and its output
- Exploring ROI-based summary of cortical thickness or gray matter volumes
- Statistical comparisons between CN vs AD

What is Freesurfer?

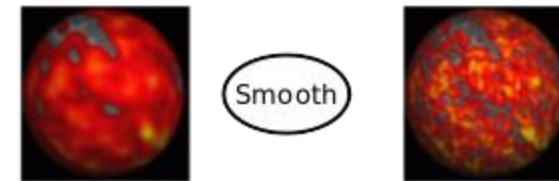


Statistical analysis



$P < 0.01$

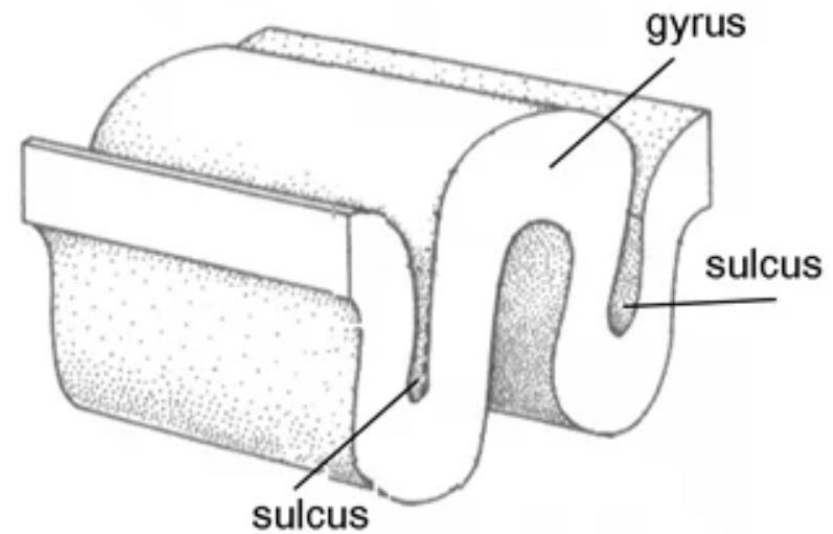
Group analysis



Thickness
group space

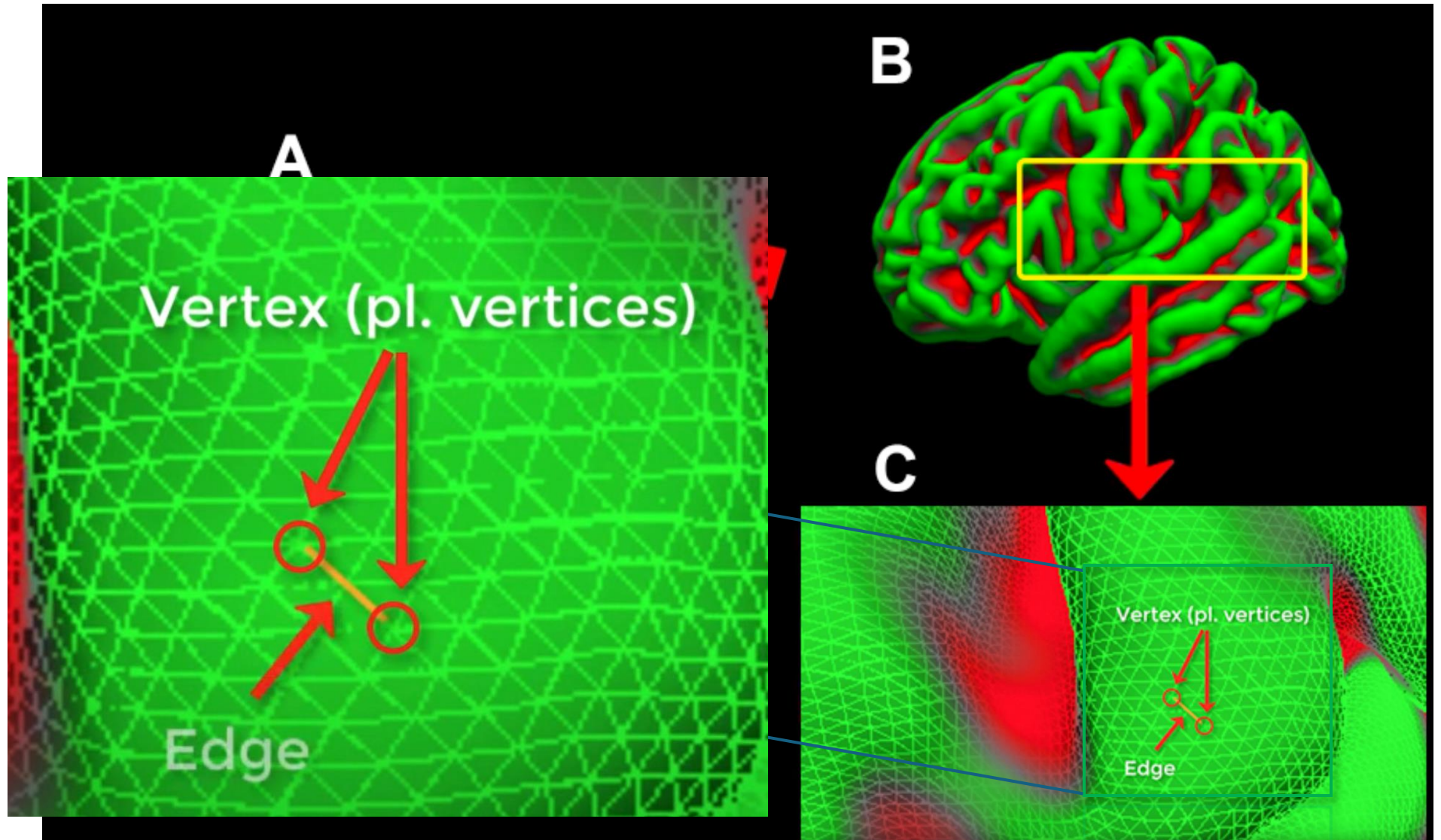
Cortex

- Gyri and Sulci of the Brain



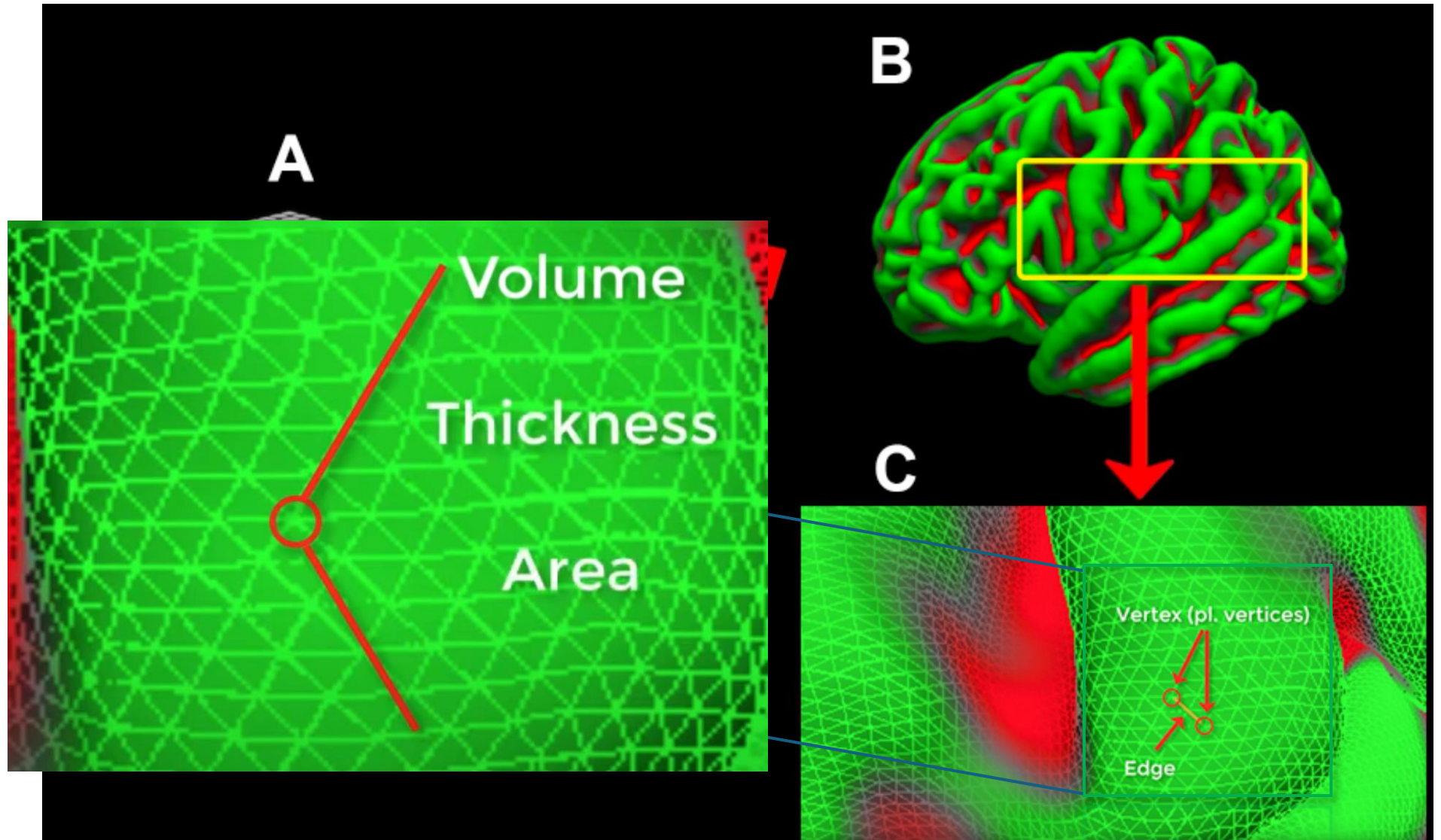
Reconstructs cortical surface from a 3D volume

recon-all



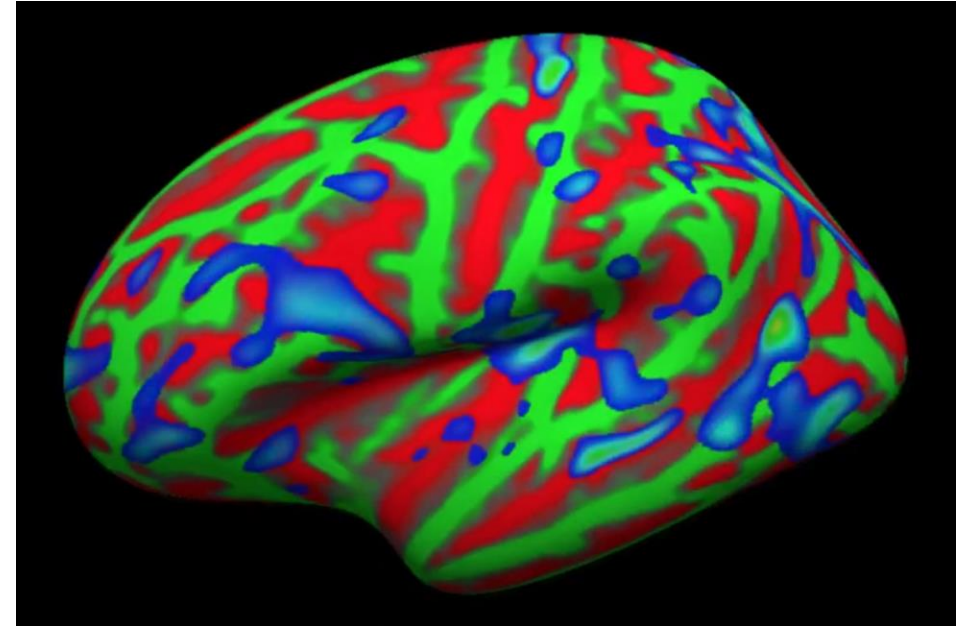
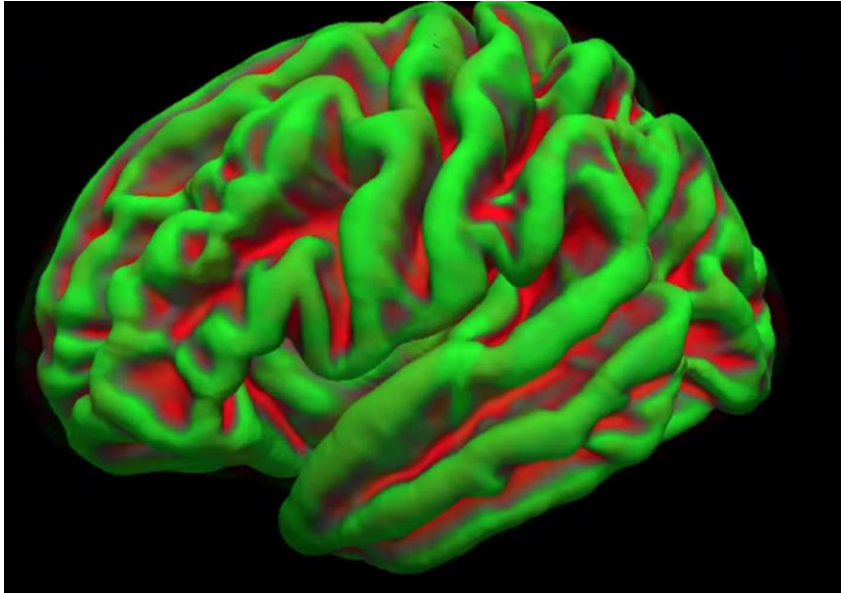
Reconstructs cortical surface from a 3D volume

recon-all



Inflated brain for better visualization

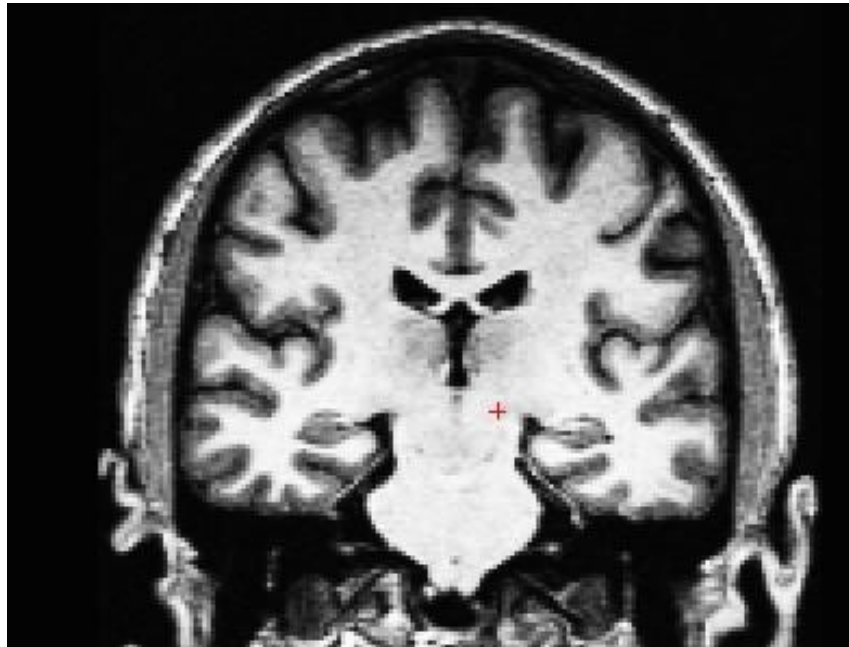
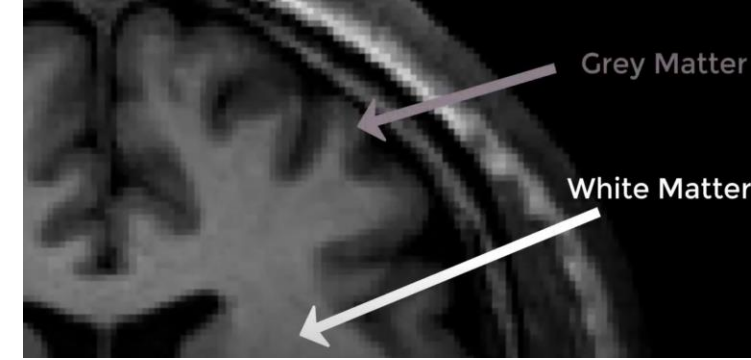
freeview



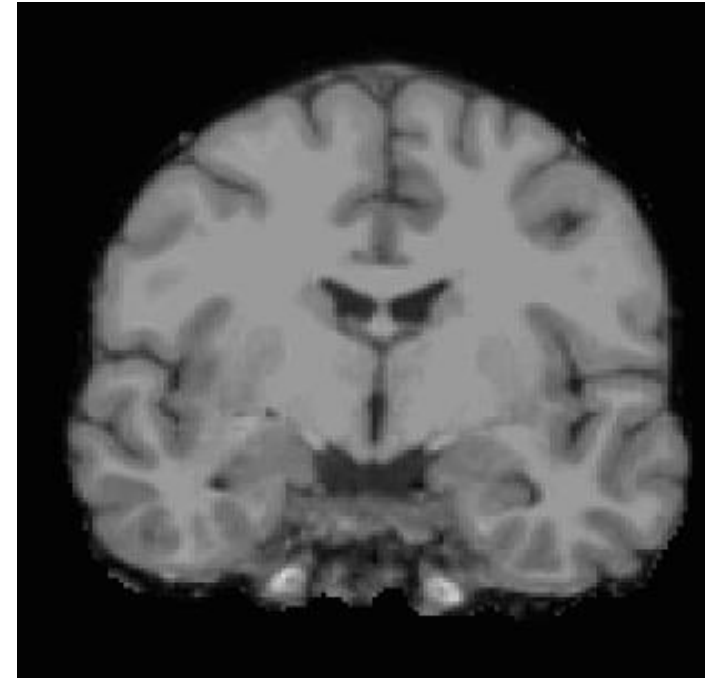
Gyrus – Green
Sulcus - Red

Grey Matter
differences – Blue

The output files of recon-all



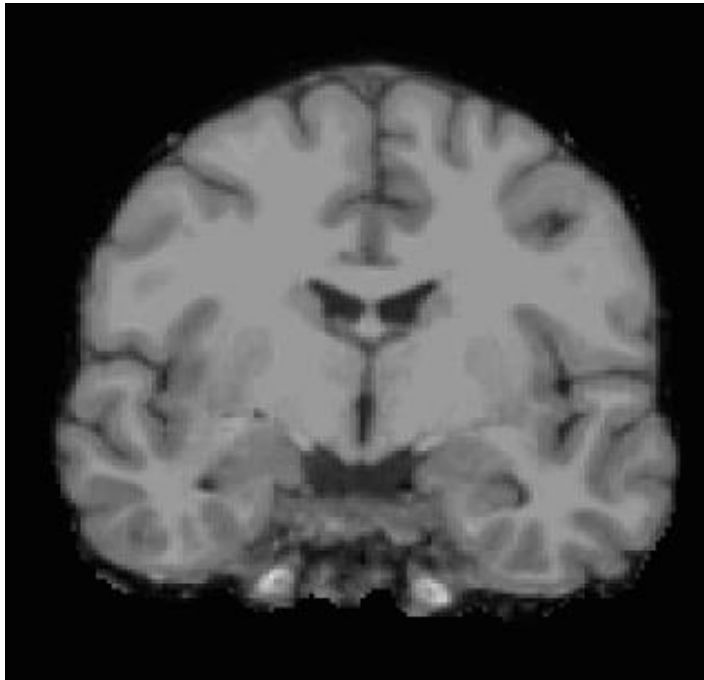
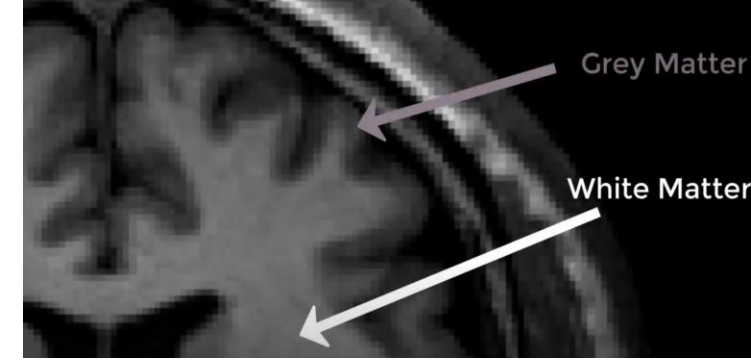
T1 weighted Input



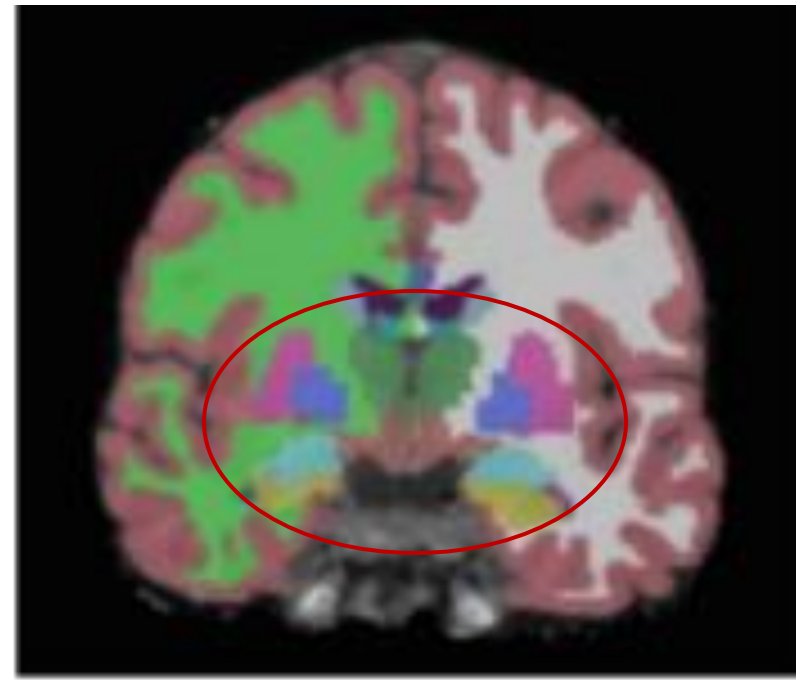
Skull stripping

brainmask.mgz
Unique to
Freesurfer

The Output of Recon-all



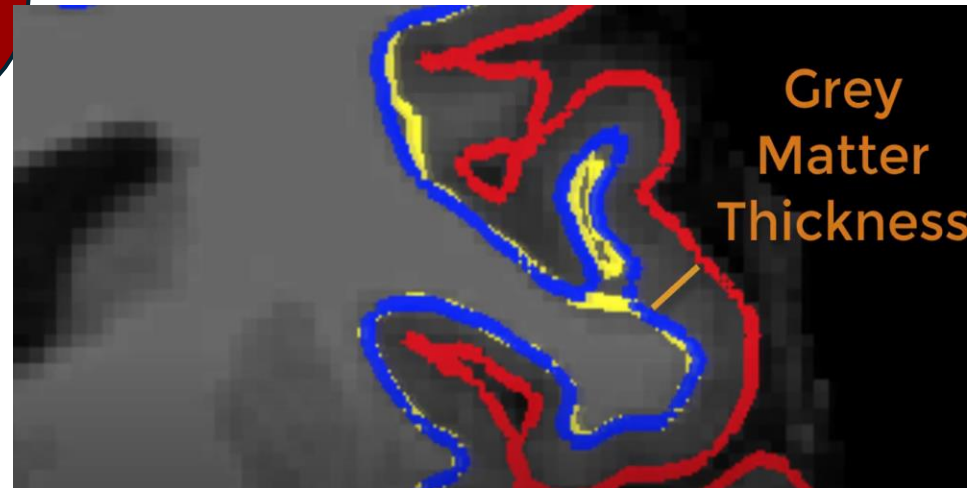
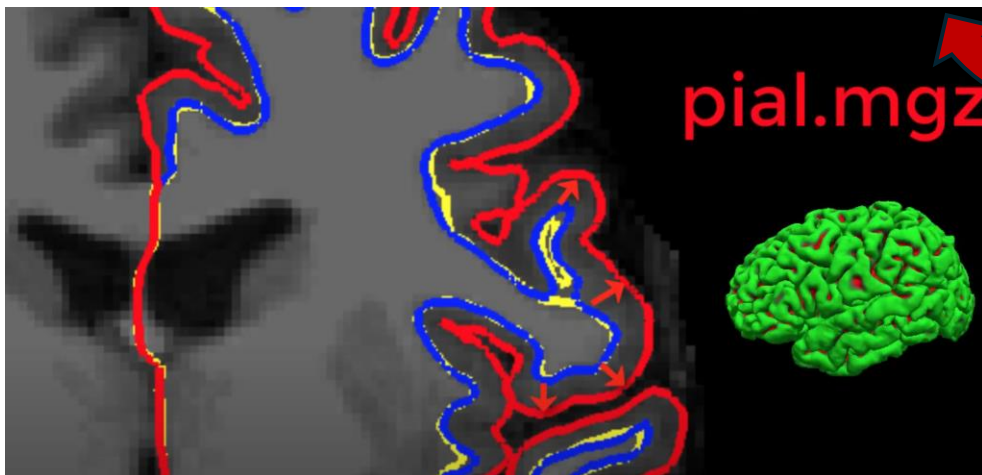
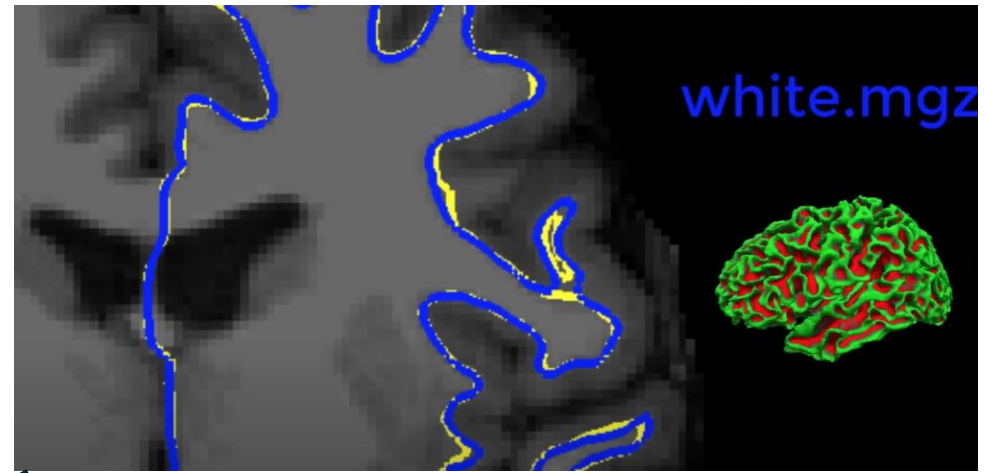
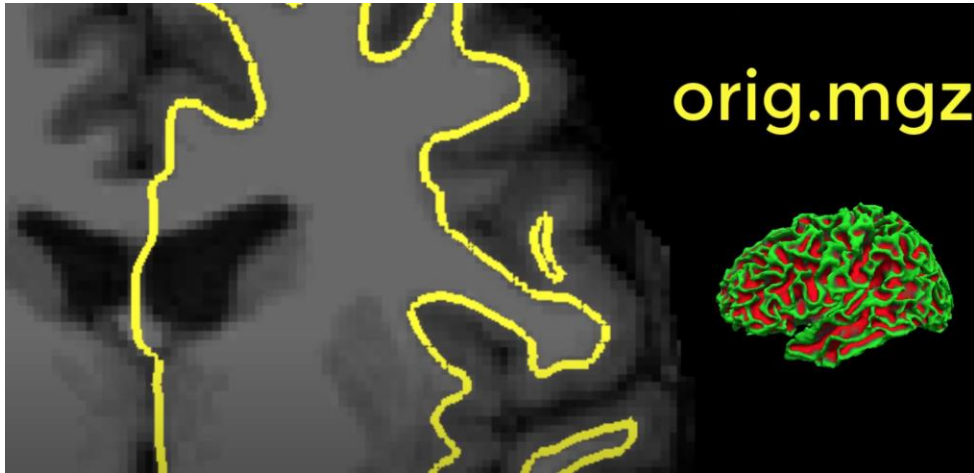
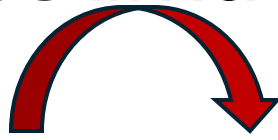
Skull stripping



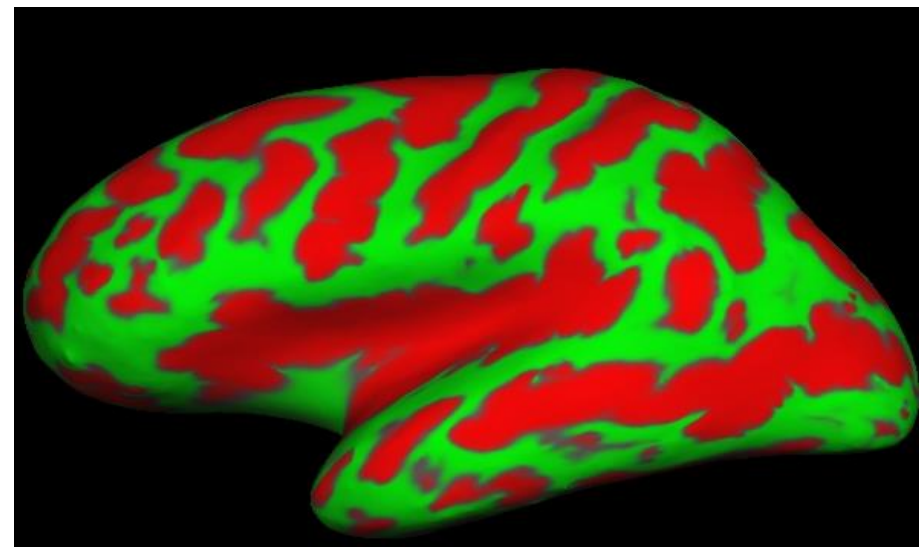
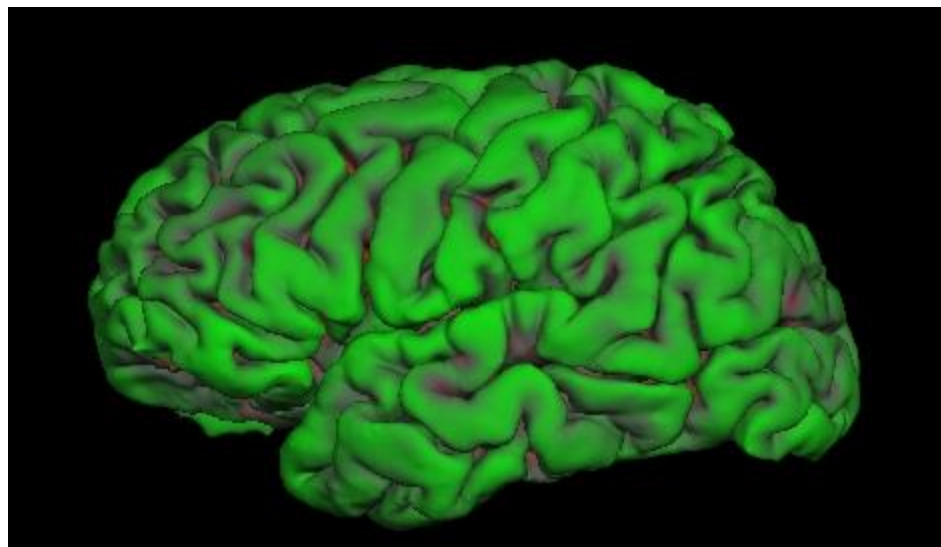
Volume labelling

aseg.mgz

Surface and thickness estimates

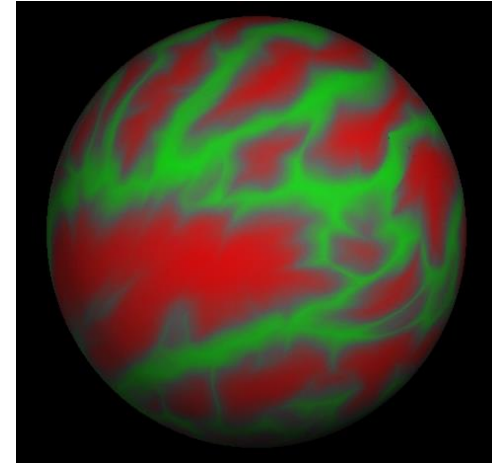
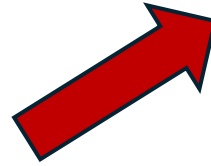
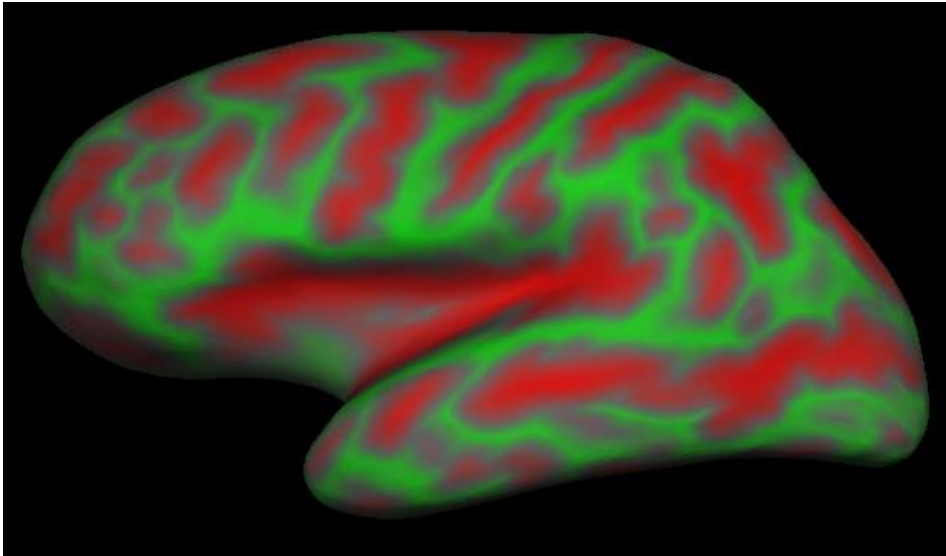


The output of recon-all

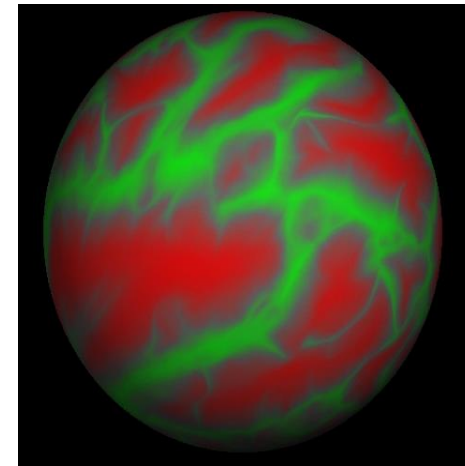
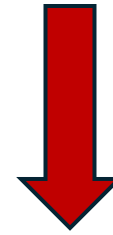


Inflated (lh and rh)

Spherical registration



Lh.sphere
Rh.sphere

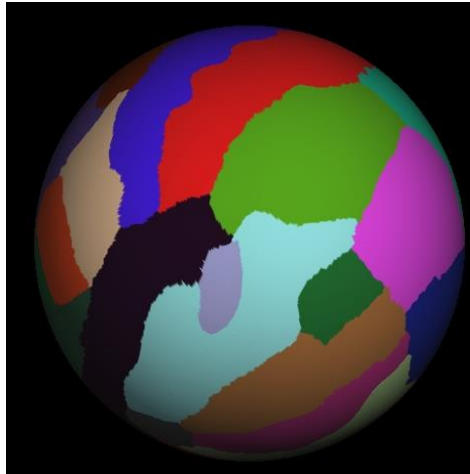


Lh.sphere.reg
Rh.sphere.reg

\$SUBJECTS_DIR/fsaverage; 40 subjects

Cortical Surface Segmentation

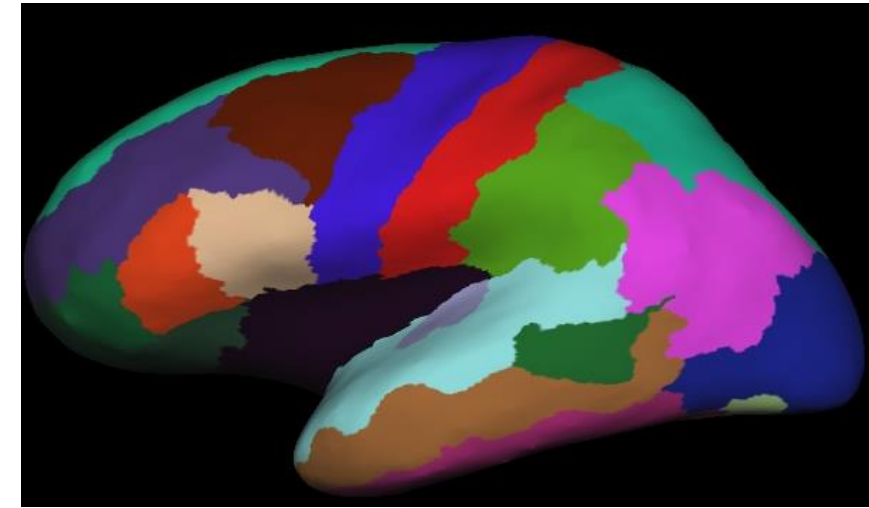
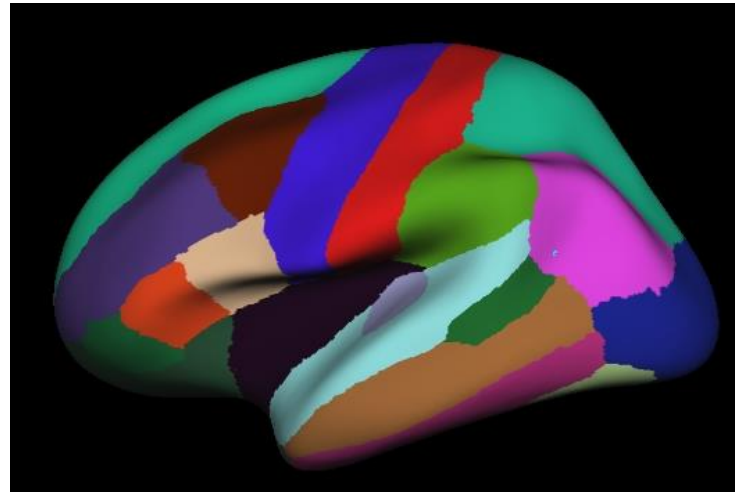
`$SUBJECTS_DIR/label/
{lh,rh}.aparc.annot`



Spherical Atlas based on Manual
Labeling

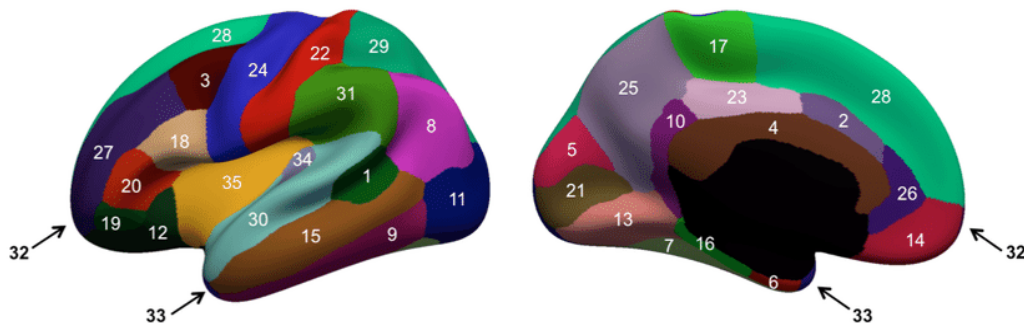


Map to Individual
Thru Spherical Reg

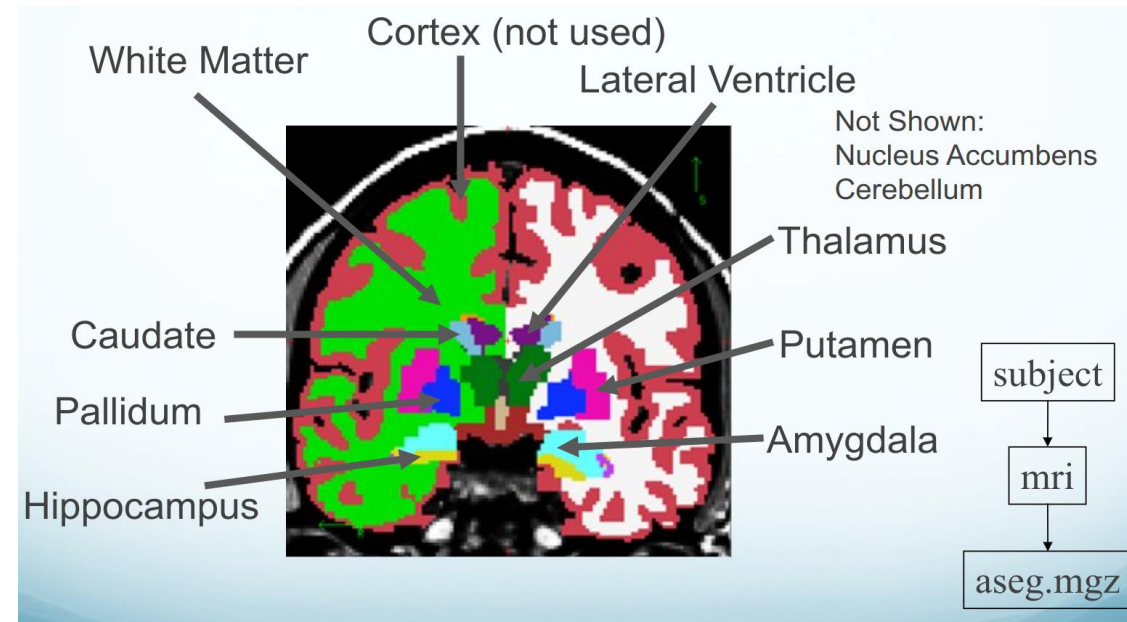


Fine-tune based on
individual anatomy

Surface parcellations and Segmentations

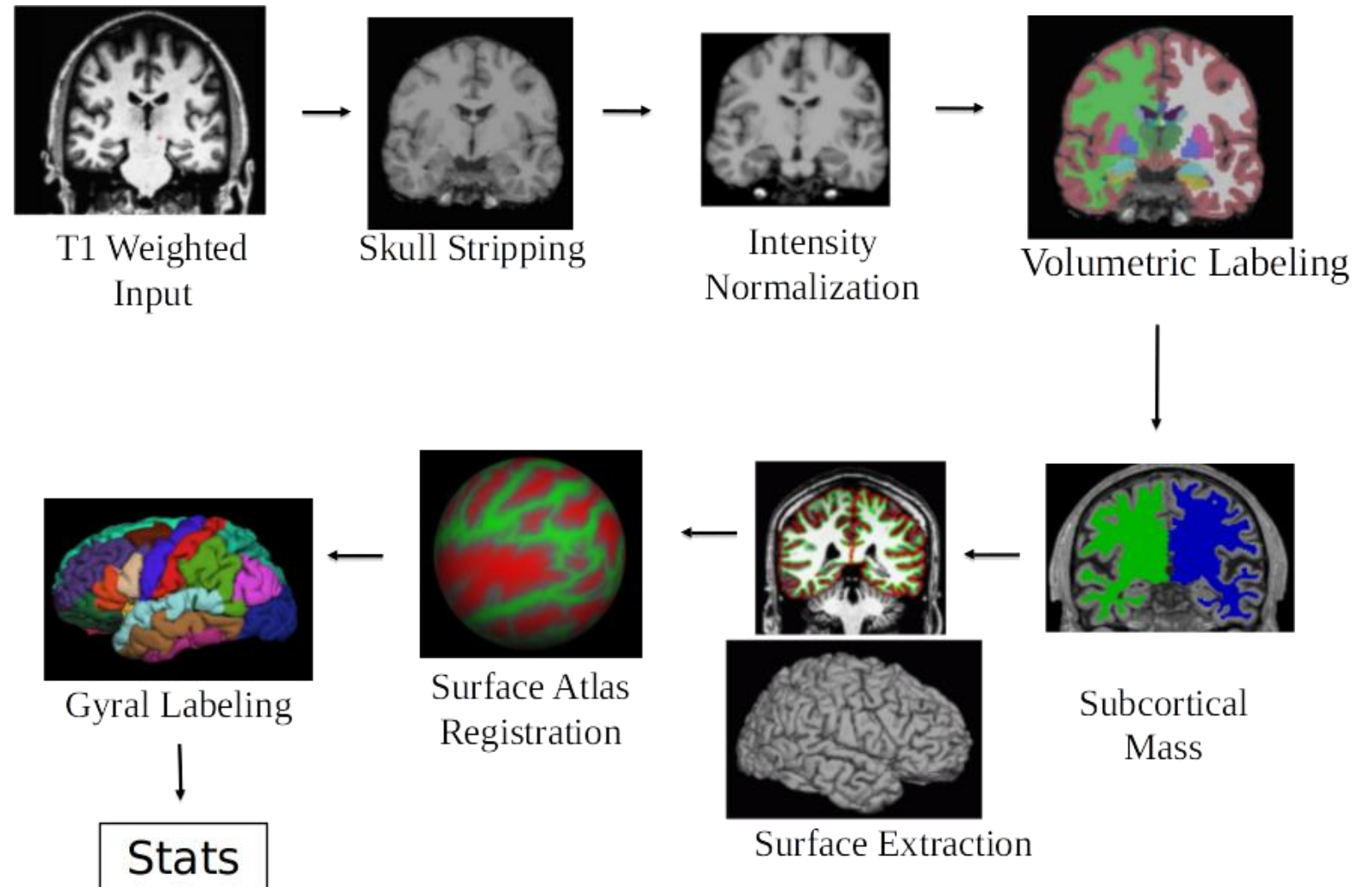


1. Bank of the Superior Temporal Sulcus	2. Caudal Anterior-cingulate Cortex
3. Caudal Middle Frontal Gyrus	4. Unmeasured Corpus Callosum
5. Cuneus Cortex	6. Entorhinal Cortex
7. Fusiform Gyrus	8. Inferior Parietal Cortex
9. Inferior Temporal Gyrus	10. Isthmus-cingulate Cortex
11. Lateral occipital cortex	12. Lateral orbital frontal cortex
13. Lingual gyrus	14. Medial orbital frontal cortex
15. Middle temporal gyrus	16. Parahippocampal gyrus
17. Paracentral lobule	18. Pars opercularis
19. Pars orbitalis	20. Pars triangularis
21. Pericalcarine cortex	22. Postcentral gyrus
23. Posterior-cingulate cortex	24. Precentral gyrus
25. Precuneus cortex	26. Rostral anterior cingulate cortex
27. Rostral middle frontal gyrus	28. Superior frontal gyrus
29. Superior parietal cortex	30. Superior temporal gyrus
31. Supramarginal gyrus	32. Frontal pole
33. Temporal pole	34. Transverse temporal cortex
35. Insula cortex	



Desikan RS, ..., Killiany RJ. An automated labeling system for subdividing the human cerebral cortex on MRI scans into gyral based regions of interest. Neuroimage. 2006

Individual Processing Stream Overview



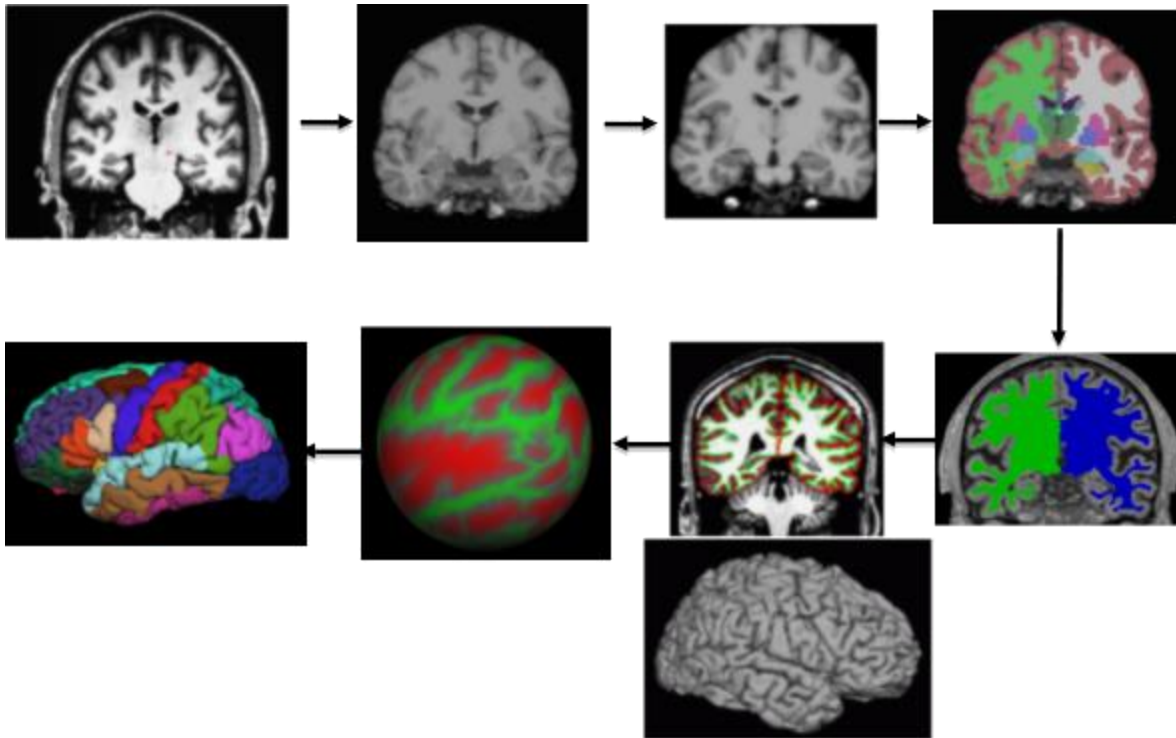
Freesurfer anatomical processing command

Set path to freesurfer installation

Then try "recon-all"

Fully Automated Reconstruction

```
recon-all -i file.nii -subject bert -all
```



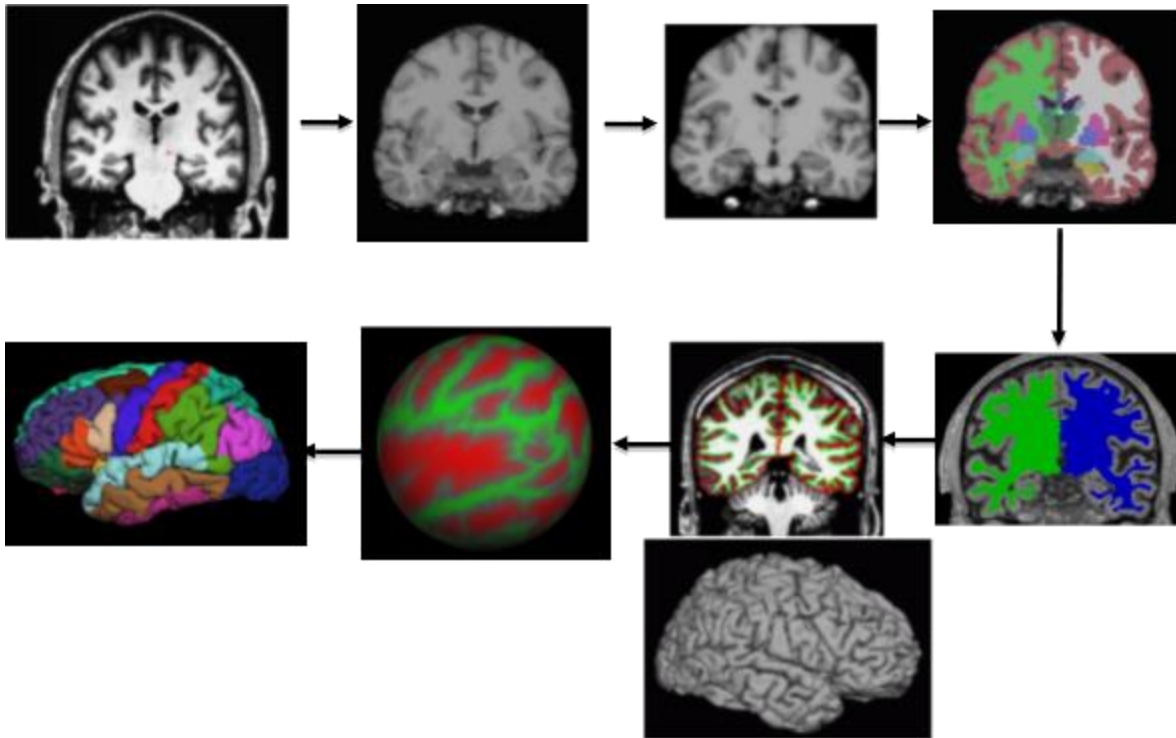
"file.nii" is a single NiFTI file pointing to the T1 MRI image

Freesurfer output variable

"\$SUBJECTS_DIR"

Fully Automated Reconstruction

```
recon-all -i file.nii -subject bert -all
```



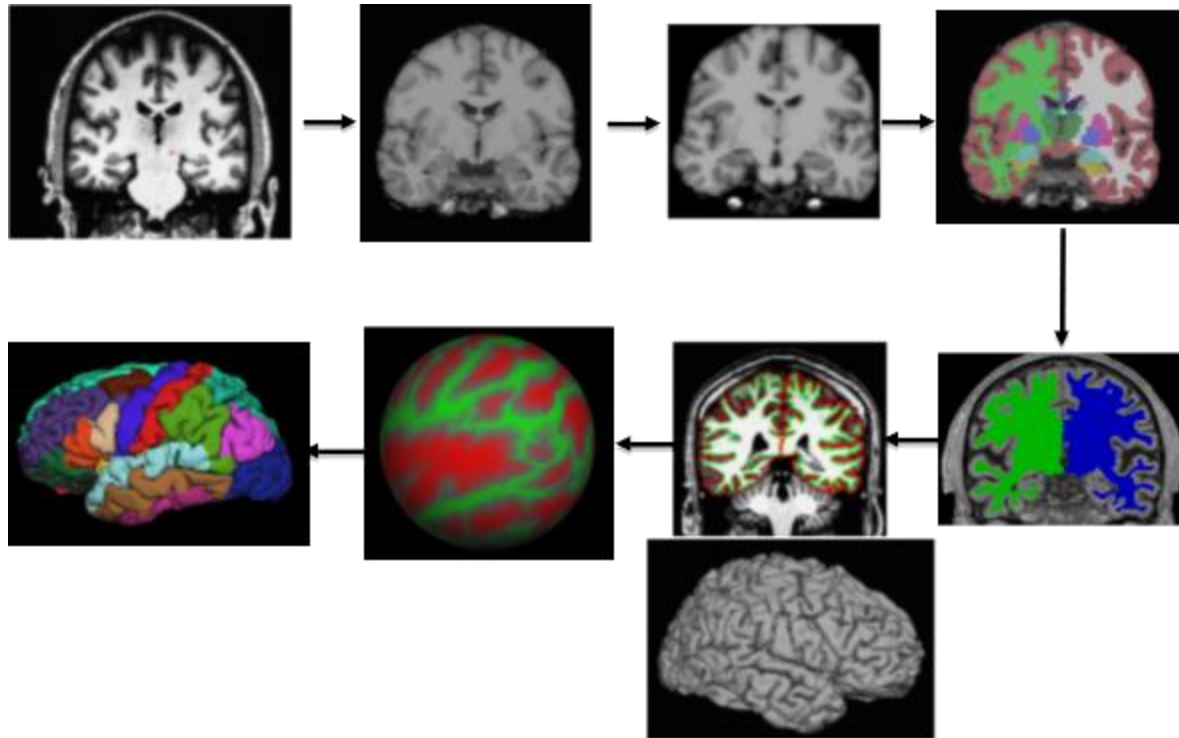
"bert" is the "name" of the subject

Creates a folder in
\$SUBJECTS_DIR

Fully Automated Reconstruction

```
recon-all -i file.nii -subject bert
```

-all



"-all" means do everything

5-10 hours on single CPU

Fully-Automated Directive

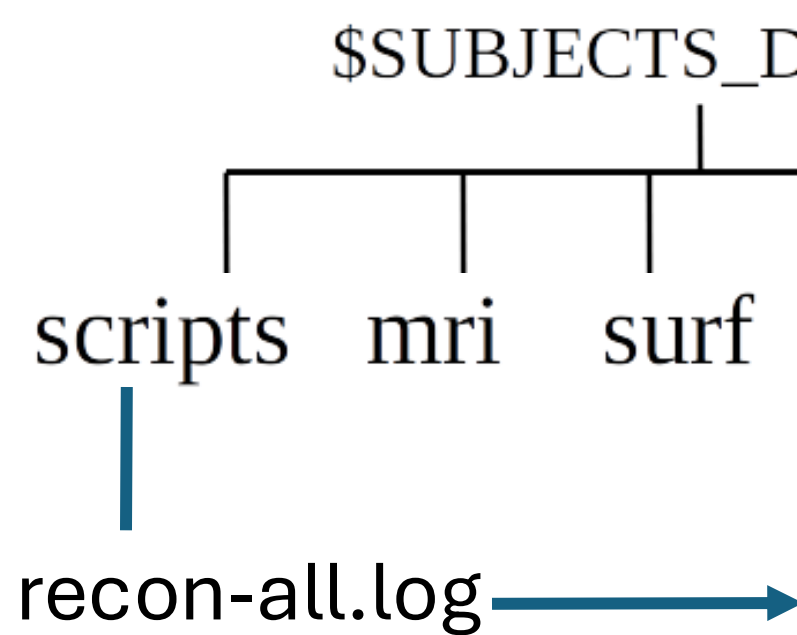
-autorecon1	process steps 1-5 (see below)
-autorecon2	process steps 6-23
	after autorecon2, check final surfaces:
	a. if wm edit was required, then run -autorecon2-wm
	b. if control points added, then run -autorecon2-cp
	c. if edits made to correct pial, then run -autorecon2-pial
	d. proceed to run -autorecon3
-autorecon2-cp	process stages 12-23 (uses -f w/ mri_normalize, -keep w/ mri_seg)
-autorecon2-wm	process stages 15-23
-autorecon2-pial	process stages 21-23
-autorecon3	process stages 24-31
-hemi ?h	just do lh or rh (default is to do both)

Autorecon Processing Stages (see -autorecon# flags above):

1. Motion Correction and Conform
2. NU (Non-Uniform intensity normalization)
3. Talairach transform computation
4. Intensity Normalization 1
5. Skull Strip
6. EM Register (linear volumetric registration)
7. CA Intensity Normalization
8. CA Non-linear Volumetric Registration
9. Remove Neck
10. LTA with Skull
11. CA Label (Volumetric Labeling, ie Aseg) and Statistics
12. Intensity Normalization 2 (start here for control points)
13. White matter segmentation
14. Edit WM With ASeg
15. Fill (start here for wm edits)
16. Tessellation (begins per-hemisphere operations)
17. Smooth1
18. Inflate1
19. QSphere
20. Automatic Topology Fixer
21. Final Surfs (start here for brain edits for pial surf)
22. Smooth2
23. Inflate2
24. Spherical Mapping
25. Spherical Registration
26. Spherical Registration, Contralateral hemisphere
27. Map average curvature to subject
28. Cortical Parcellation - Desikan_Killiany and Christophe (Labeling)
29. Cortical Parcellation Statistics
30. Cortical Ribbon Mask
31. Cortical Parcellation mapping to Aseg

```
recon-all -i file.nii -subject bert -all
```

Upon successful completion of recon-all



```
Thu Aug 5 14:11:38 IST 2021
/media/cogsci/SeagateBackupPlusDrive/madhukar/ADNI/processed/011_S_4075_tp0
/home/cogsci/AJP/freesurfer/bin/recon-all
-recon-all -s 011_S_4075_tp0 -i 011_S_4075_tp0_T1w.nii.gz -cw256
subjid 011_S_4075_tp0
setenv SUBJECTS_DIR /media/cogsci/SeagateBackupPlusDrive/madhukar/ADNI/processed
FREESURFER_HOME /home/cogsci/AJP/freesurfer
Actual FREESURFER_HOME /home/cogsci/AJP/freesurfer
build-stamp.txt: freesurfer-Linux-centos6_x86_64-stable-pub-v6.0.0-2beb96c
Linux cogsci-server 4.15.0-142-generic #146~16.04.1-Ubuntu SMP Tue Apr 13 09:27:15 UTC 2021 x86_64 x86_64 x86_64 GNU/Linux
cputime unlimited
filesize unlimited
datasize unlimited
stacksize 8192 kbytes
coredumpsize 0 kbytes
memoryuse unlimited
vmemoryuse unlimited
descriptors 1024
memorylocked 64 kbytes
maxproc 192534
maxlocks unlimited
maxsignal 192534
maxmessage 819200
maxnice 0
maxrtprio 0
maxrttime unlimited

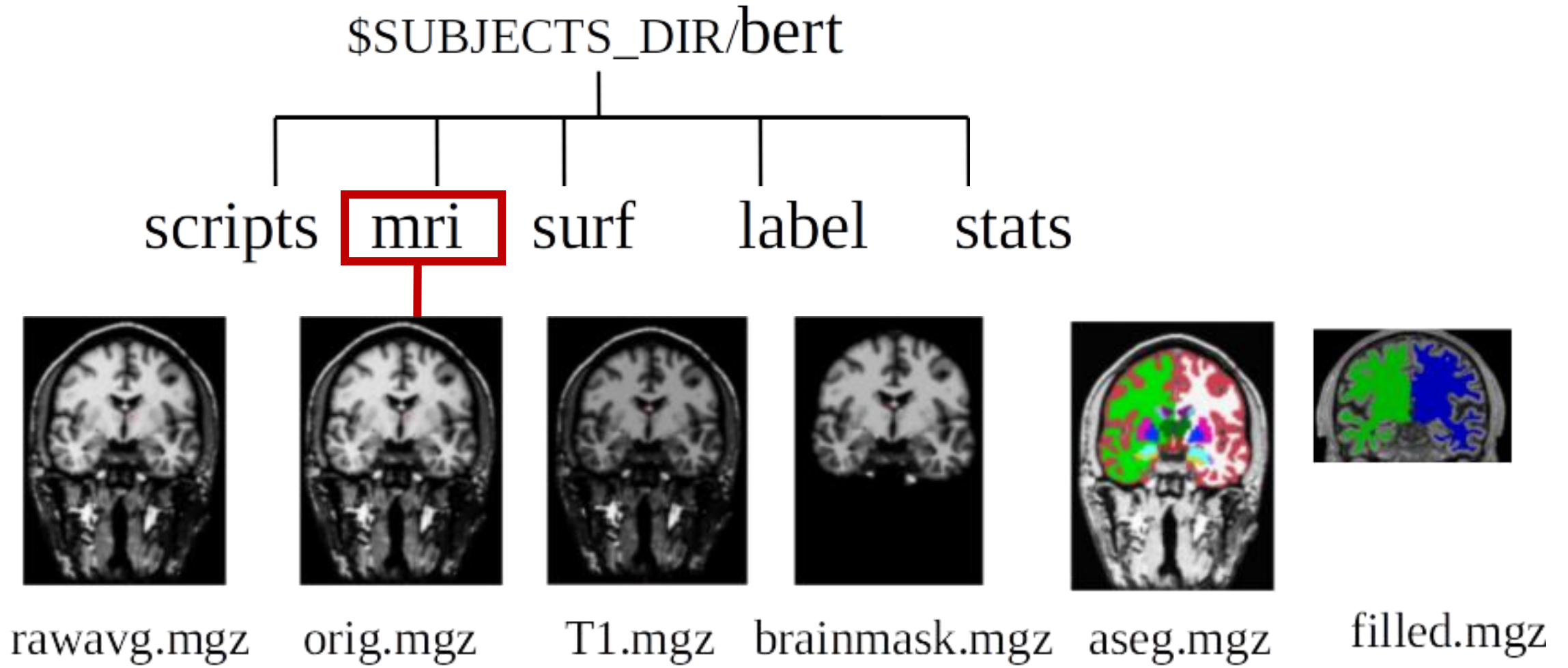
Mem:      total      used      free      shared  buff/cache  available
Swap:    16678908    339968    16338940    218268    36074164    36764268

#####
program versions used
$Id: recon-all,v 1.580.2.16 2017/01/18 14:11:24 zkaufman Exp $
$Id: mri_motion_correct.fsl,v 1.15 2016/02/16 17:17:20 zkaufman Exp $
mri_convert.bin -all-info
ProgramName: mri_convert.bin ProgramArguments: -all-info ProgramVersion: $Name: stable6 $ TimeStamp: 2021/08/05-08:41:38-GMT BuildTimeStamp: Jan 18 2021

3732 2252 4770 1.950 0.500 0.128 0.230 95 5.7 V1_exvivo
4505 2659 5723 1.980 0.457 0.136 0.039 68 7.2 V2_exvivo
315 203 703 2.726 0.322 0.127 0.029 4 0.4 MT_exvivo
355 237 1469 4.212 0.737 0.087 0.018 2 0.3 perirhinal_exvivo
224 157 611 3.152 0.634 0.122 0.023 2 0.2 entorhinal_exvivo

Started at Thu Aug 5 14:11:38 IST 2021
Ended at Fri Aug 6 00:43:21 IST 2021
#@## recon-all-run-time-hours 10.529
recon-all -s 011_S_4075_tp0 finished without error at Fri Aug 6 00:43:21 IST 2021
```

Upon successful completion of recon-all



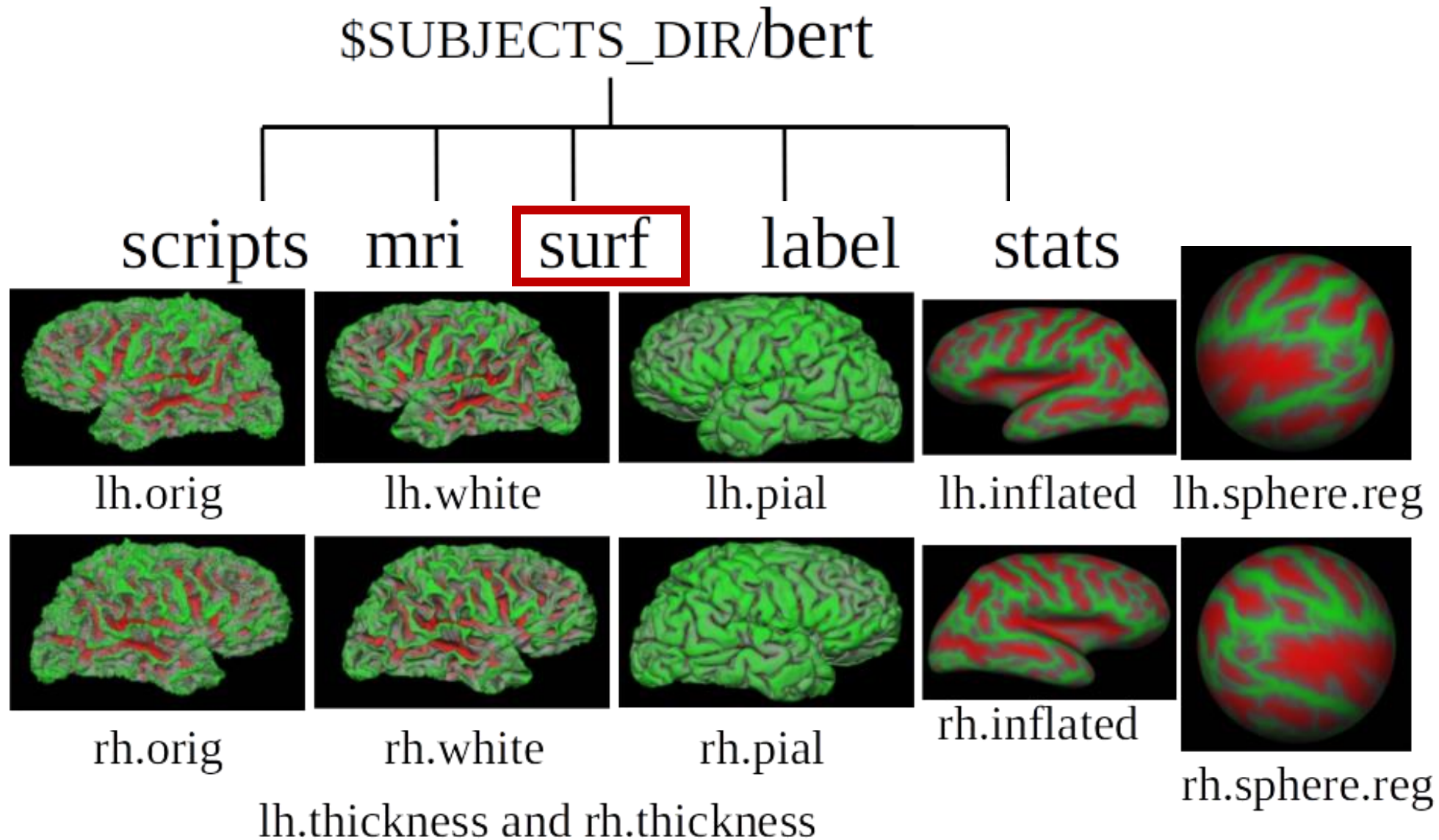
Viewing volumes with Freeview

```
set $SUBJECTS_DIR"
```

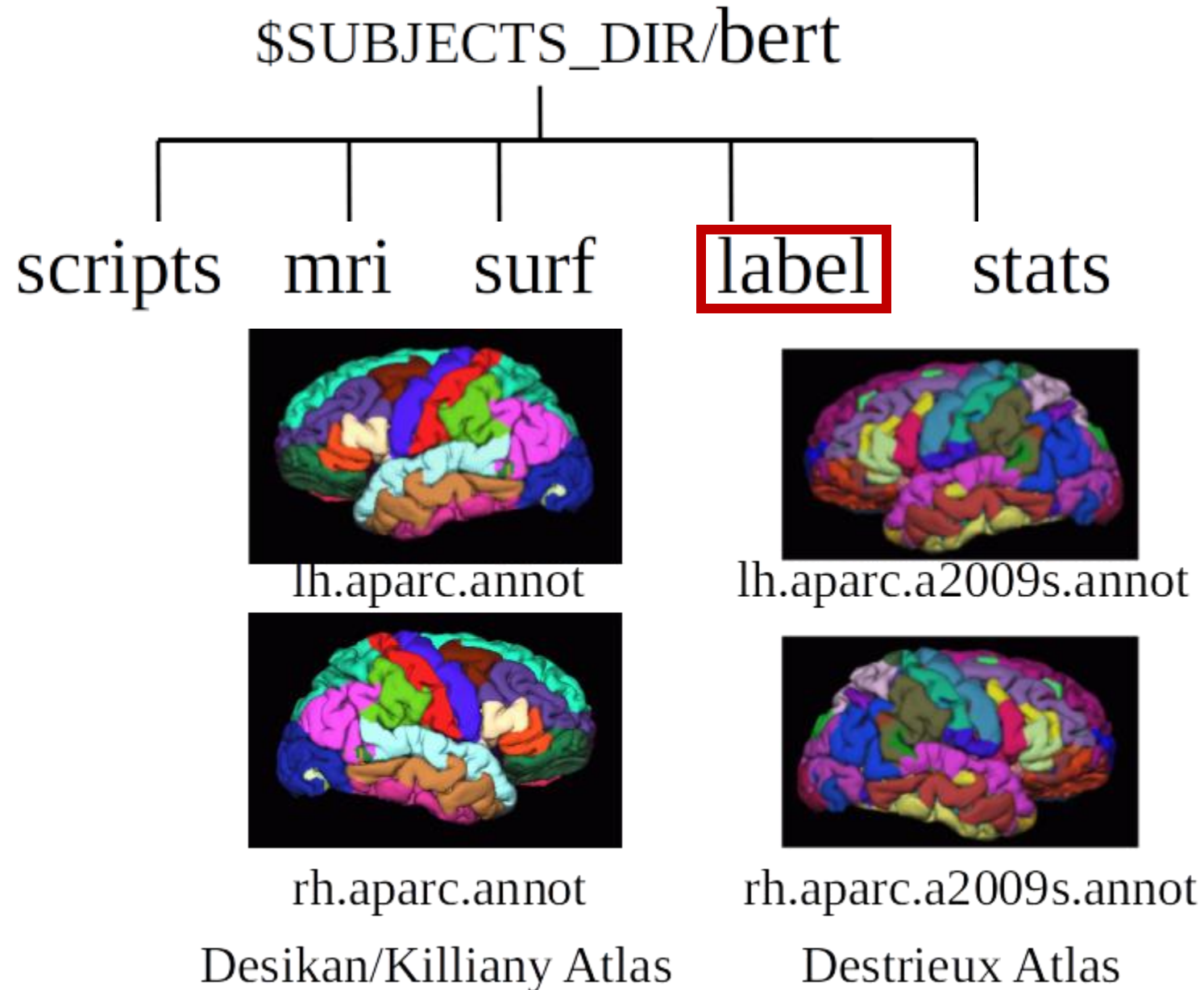
```
freeview -v \  
011_S_4075_tp0/mri/T1.mgz \  
011_S_4075_tp0/mri/wm.mgz \  
011_S_4075_tp0/mri/brainmask.mgz \  
011_S_4075_tp0/mri/aseg.mgz:colormap=lut:opacity=0.2 \  
-f 011_S_4075_tp0/surf/lh.white:edgecolor=blue \  
011_S_4075_tp0/surf/lh.pial:edgecolor=red \  
011_S_4075_tp0/surf/rh.white:edgecolor=blue \  
011_S_4075_tp0/surf/rh.pial:edgecolor=red
```

https://surfer.nmr.mgh.harvard.edu/fswiki/FsTutorial/OutputData_freeview

Upon successful completion of recon-all



Upon successful completion of recon-all



Viewing surface atlases with freeview

```
set $SUBJECTS_DIR"
```

```
freeview -f
```

```
011_S_4075_tp0/surf/lh.pial:annot=aparc.annot:name=pial_a  
parc:visible=0 \
```

```
011_S_4075_tp0/surf/lh.pial:annot=aparc.a2009s.annot:nam  
e=pial_aparc_des:visible=0 \
```

```
011_S_4075_tp0/surf/lh.inflated:overlay=lh.thickness:overlay  
_threshold=0.1,3::name=inflated_thickness:visible=0 \
```

```
011_S_4075_tp0/surf/lh.inflated:visible=0 \
```

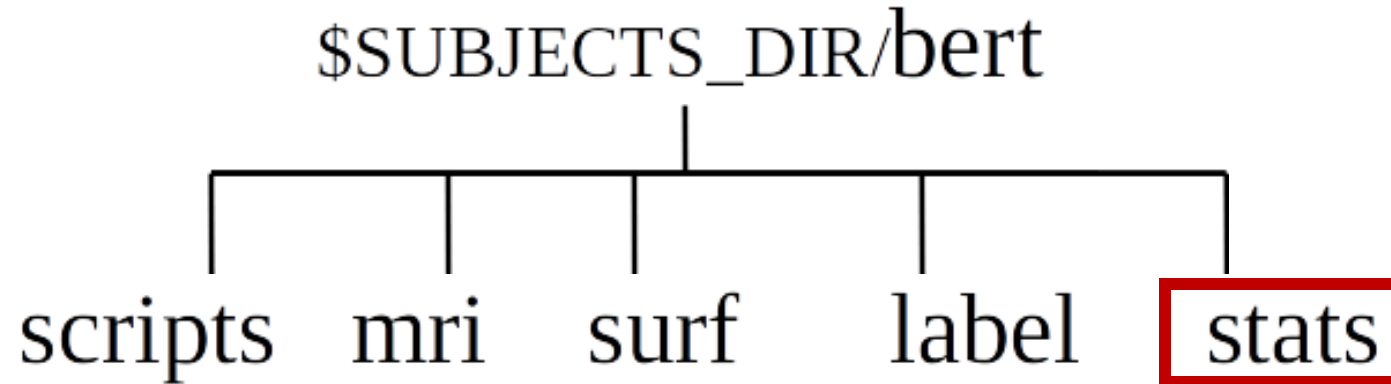
```
011_S_4075_tp0/surf/lh.white:visible=0 \
```

```
011_S_4075_tp0/surf/lh.pial \
```

```
--viewport 3d
```

https://surfer.nmr.mgh.harvard.edu/fswiki/FsTutorial/OutputData_freeview

Upon successful completion of recon-all



- aseg.stats -- subcortical volumetric stats
- wmparc.stats -- white matter segmentation volumetric stats
- lh/rh.aparc.stats -- left/right hemi Desikan Killiany surface atlas
- lh/rh.aparc.a2009.stats -- left/right hemi Destrieux

ROI summaries – stats folder

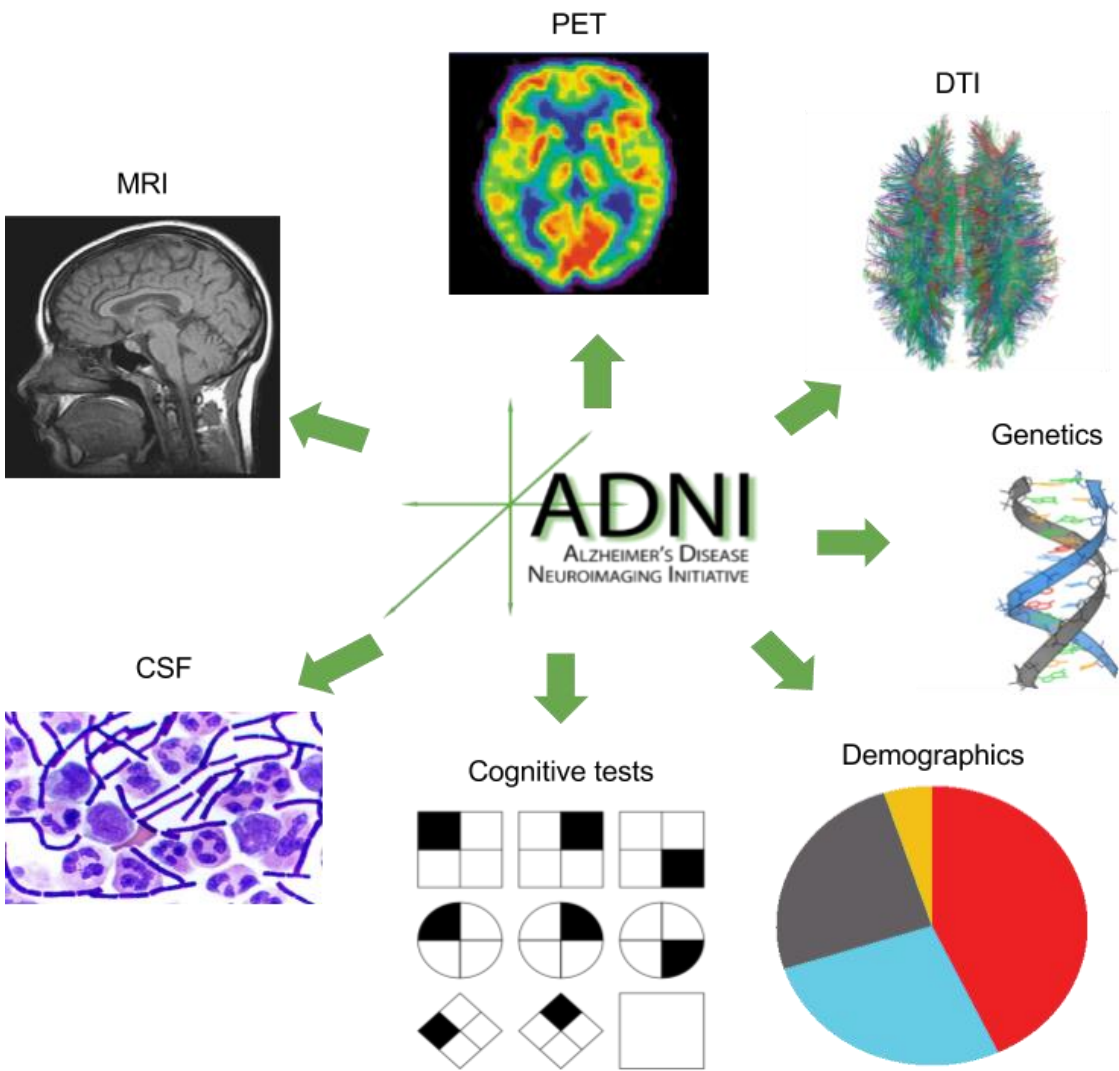
Index	SegId	NVoxels	Volume_mm3	StructName	Mean	StdDev	Min	Max	Range
1	4	7230	7553.6	Left-Lateral-Ventricle	29.2	12.5	11.0	91.0	80.0
2	5	267	311.1	Left-Inf-Lat-Vent	51.3	11.5	14.0	84.0	70.0
3	7	10858	11506.0	Left-Cerebellum-White-Matter	86.8	6.7	24.0	106.0	82.0
4	8	50468	50270.0	Left-Cerebellum-Cortex	63.6	11.0	5.0	95.0	90.0
5	10	6625	6379.0	Left-Thalamus-Proper	91.6	8.8	40.0	111.0	71.0
6	11	3335	3407.1	Left-Caudate	81.2	7.1	50.0	100.0	50.0
7	12	4369	4385.7	Left-Putamen	87.2	5.2	57.0	106.0	49.0
8	13	1943	1923.6	Left-Pallidum	98.7	3.9	65.0	112.0	47.0
11	16	17918	17745.1	Brain-Stem	85.7	10.5	25.0	108.0	83.0
12	17	4014	3842.1	Left-Hippocampus	71.1	7.7	26.0	101.0	75.0
13	18	1598	1559.4	Left-Amygdala	72.8	6.4	27.0	97.0	70.0

Routines to generate spread sheets for multiple subjects

- `asegstats2table --help`
- `aparcstats2table --help`

Alzheimer's Disease Neuroimaging Initiative

- ADNI



	Screening/ baseline visit	Month 6	Month 12	Month 18	Month 24	Month 30	Month 36
Normal							
Clinical	X	X	X		X		X
MRI 1.5	X	X	X		X		X
MRI 3	X	X	X		X		X
PET							
CSF							
Blood							
Telephone							
Mild cognitive impairment							
Clinical							
MRI 1.5							
MRI 3							
PET	X	X	X	X	X		X
CSF	X		X				
Blood	X	X		X	X		X
Telephone	X	X		X	X		X
Alzheimer disease							
Clinical	X	X	X		X		
MRI 1.5	X	X	X		X		
PET	X	X	X		X		
CSF	X		X				
Blood	X	X	X		X		
Telephone						X	

Ø2000 subjects
 ➤8000 scans

This session

- ADNI
 - Baseline visit
 - Number of subjects: 98
 - Age matched: 75/73 (CN/AD)
 - Gender: M/F (45/53)
 - Cognitively Normal : 51 -- M/F (21/30)
 - Alzheimer's Disease : 47 – M/F (24/23)
 - MMSE : 19-30
 - Severe cognitive impairment: 0-17
 - Mild cognitive impairment: 18-23
 - No cognitive impairment: 24-30

