

//3D PRINT YO BRAIN

//By Keiland C

//nov16 vs

//Programs needed:

//Freesurfer -open source mri software

//MRICron (needs NIFTI format) or matlabs version -easily convert images

//meshlab or autodesk -open source still viewer and manipulator

//links

https://en.wikipedia.org/wiki/Magnetic_resonance_imaging

<http://freesurfer.net/>

***<http://freesurfer.net/fswiki> //this was the most helpful

<https://surfer.nmr.mgh.harvard.edu/fswiki/recon-all> //this details the all in one step, it describes in more detail what it is doing, perfect if you want to customize a step

<http://people.cas.sc.edu/rorden/mricron/index.html>

<https://www.mathworks.com/matlabcentral/fileexchange/42997-dicom-to-nifti-converter--nifti-tool-and-viewer>

<http://meshlab.sourceforge.net/>

<http://www.autodesk.com/>

//Convert the images, freesurfer gets weird with dicoms for some-reason

//test out different micron settings, this will depend on the type of scanner used

//I think i ended up going with **SFL/SPM8** , I need to read up

more on this though

//something funny happens with dti vs normal imaging
around this or the next step

//freesurfer code:

//I had to move the free surfer install to the documents folder
due to permissions errors

//sudo wouldn't fix it...

//use the mail archives to fix errors

<https://mail.nmr.mgh.harvard.edu/pipermail/freesurfer/2012-July/024750.html>

//make sure the tutorials work on your machine, this could
save a lot of time (trust me)

//recon-all is amazing and will automate pretty much
everything,

//especially the skull stripping stage

//(it seems mri scientists are fairly lazy as well)

//check the docs if need to segment out or edit the process

**recon-all -s {your name here} -all -i {path to your newly
converted file}**

//this will process for awhile (no really, took like 16+ hours on
my mbp)

//and then save a folder that you named after yourself,

//into the subjects folder

//to convert pial to stl etc.

**mrisc_convert {path to freesurfer}/subjects/mybrain/surf/
rh.pial rh.stl**

//use the same code with the lh.pial to lh.stl

//import the stl's into meshlab or whatever software you choose

//run a smoothing algorithm on it and maybe reconstruct if the file is unseemly large

//It's best to print hemisphere by hemisphere, then glue

//TODO

//figure out how to print the cerebellum with free surfer,
//(i don't want the 3d me to be shaky)

<https://mail.nmr.mgh.harvard.edu/pipermail/freesurfer/2016-May/045371.html>

//use mri_tessellate?

https://surfer.nmr.mgh.harvard.edu/fswiki/mri_tessellate

<https://surfer.nmr.mgh.harvard.edu/fswiki/tessellate>

//figure out the problems with the DTI anatomy images,
<https://mail.nmr.mgh.harvard.edu/pipermail/freesurfer/2012-July/024750.html>

//print the DTI connectome?!?!?!?!?!?!?