

3dQwarp and Its *Nwarp* Friends

Or, How I Learned to Stop Worrying and
Love Getting My Datasets all Warped

Linear and Nonlinear Warping

- The Central Equation:
 - $S(x)$ = source image $B(x)$ = base image
 - $S(W(x)) \approx B(x)$ where $W(x)$ = desired warp function
= shows where each point x in B maps to in S
- 3dAllineate: $W(x) = \mathbf{M}x$ where $\mathbf{M} = 4 \times 3$ matrix
 - M has 12 parameters to optimize
- 3dQwarp: $W(x) = W_1(W_2(\dots W_{n-1}(W_n(x))) \dots)$
 - Each $W_k(x)$ is a 3D polynomial function over a “patch” that covers part of the 3D brain volume
 - Patches start big [$W_1(x)$] and shrink and shrink
 - Cubic patch = 24 parameters ; Quintic = 81 params
 - By the end, 1000s of parameters have been used

The Good and The Ugly

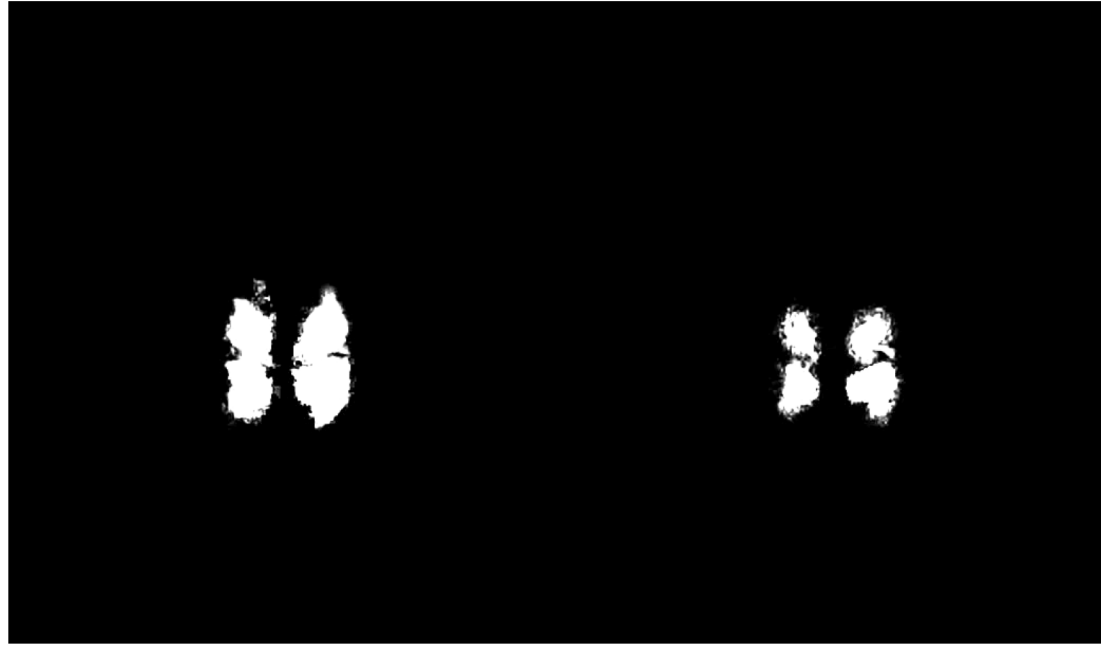
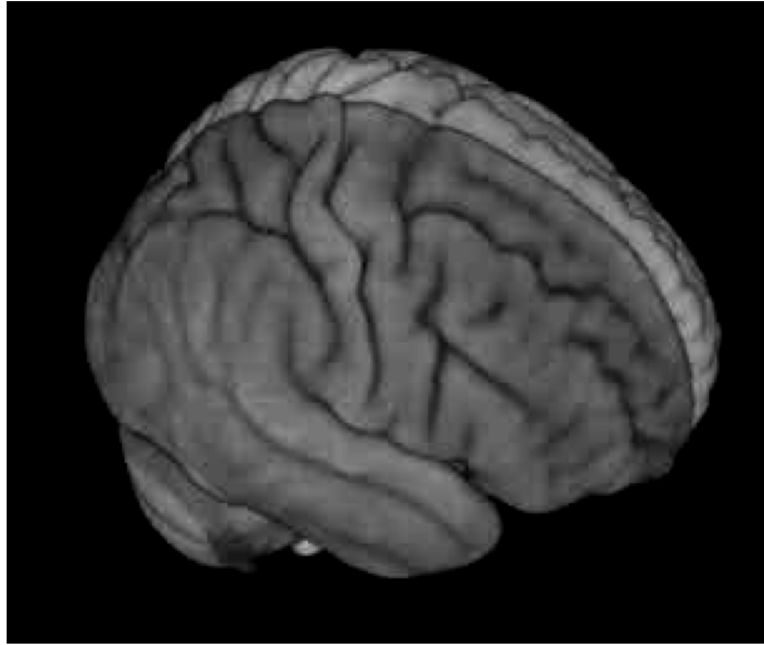
- the Good:

- Nonlinear warping can match anatomical structures between subjects more closely than linear transformation
- Can also be used for intra-subject warping for high accuracy matching (e.g., pre- and post-surgery)

- the Ugly:

- Nonlinear warping can seriously distort when it tries to match in regions that don't really "fit together" (e.g., 2 gyri in one person, 1 gyrus in another)
- Extraneous small features can drive warping in strange ways (unlike linear transformation)
- Partial brain coverage is a problem

Start: Looking Good



Compare FSL *FNIRT* vs AFNI *3dQwarp*
Average of 101 brain volumes warped to template

Good Matches to Anatomical Labels

Align MindBoggle 101 T₁ Datasets to Separate Template:
Overlap Probability Maps for 3 of the Labeled Regions

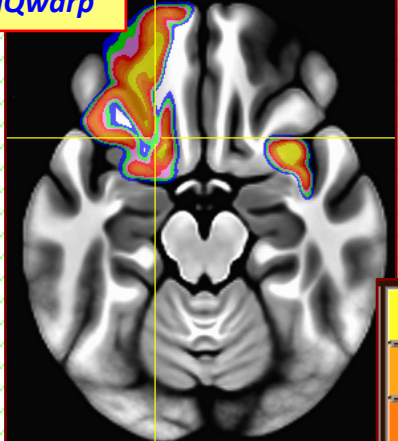
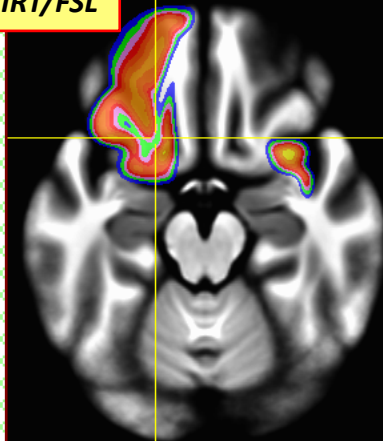
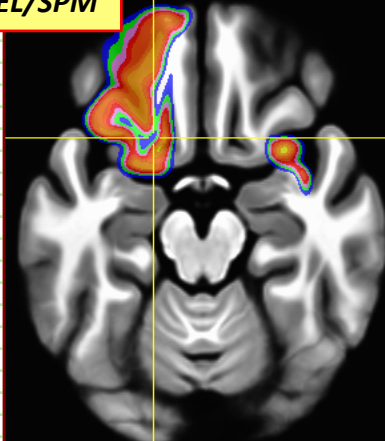
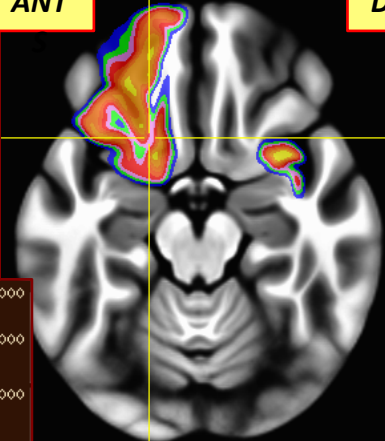
LH: lateral orbital frontal
RH: caudal anterior cingulate
RH: insula

ANT

DARTEL/SPM

FNIRT/FSL

3dQwarp

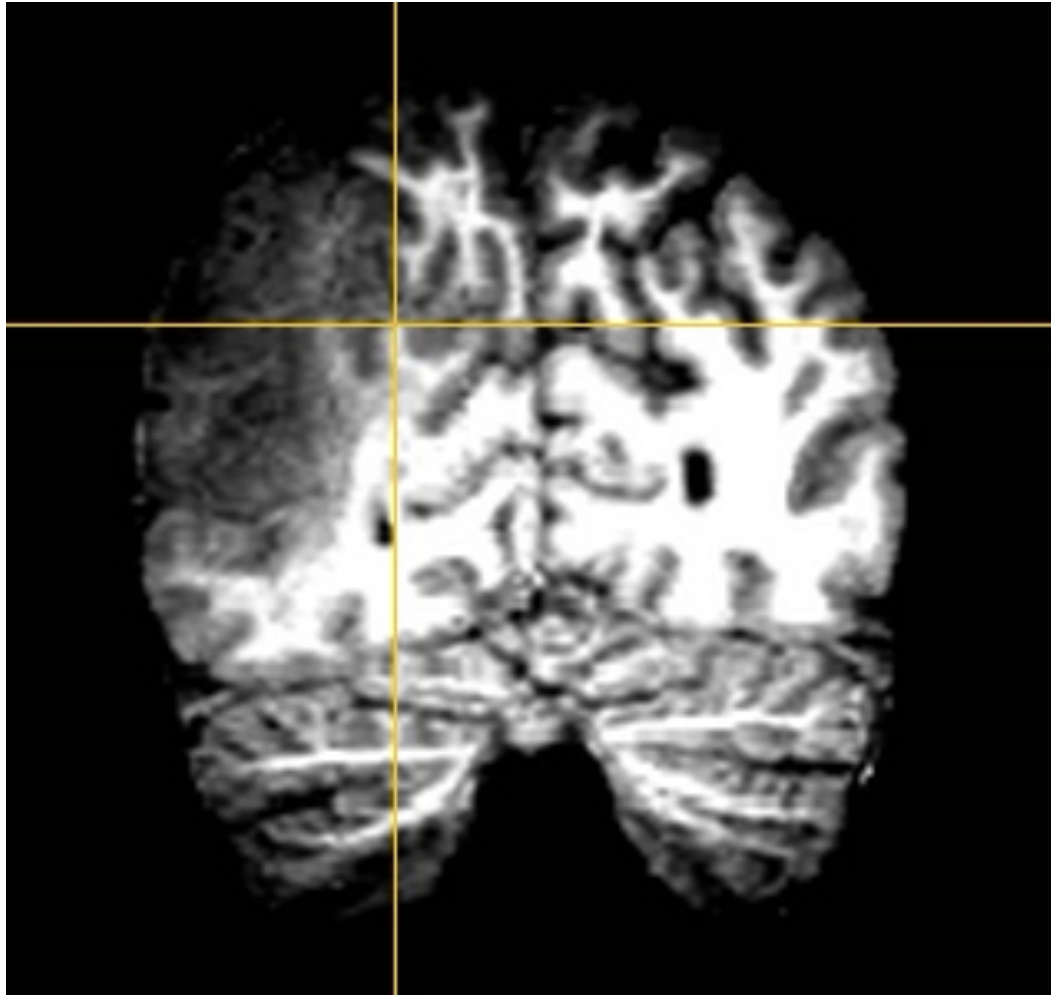


ANTs & DARTEL
& FNIRT
run with default
settings

3dQwarp distribution of overlap
probabilities is 2nd order stochastically
dominant in a majority of 62 labeled regions

More yellow in the
overlay means more
90+% overlap in labels

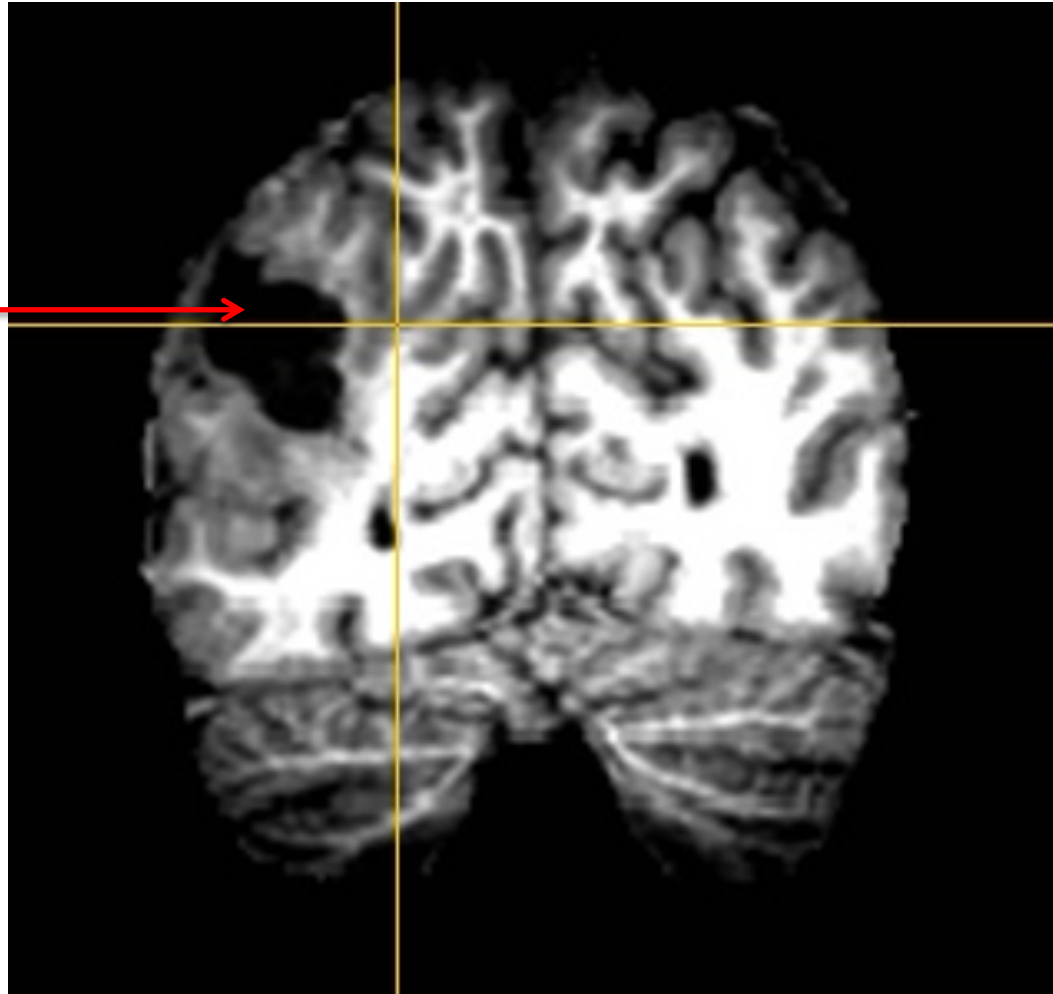
Maybe Even Useful: Neurosurgery



Pre-surgical volume

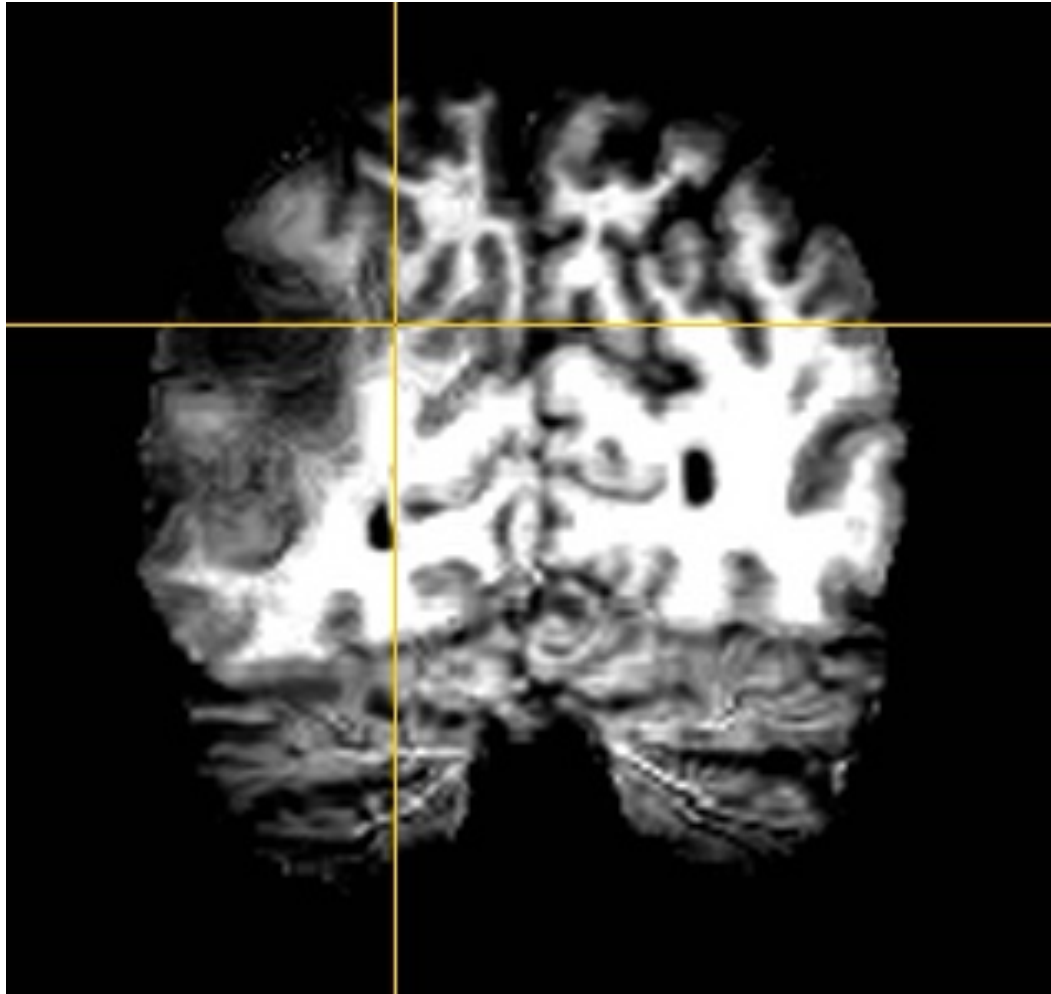
Neurosurgery

Manually
drawn
"exclusion
Mask"



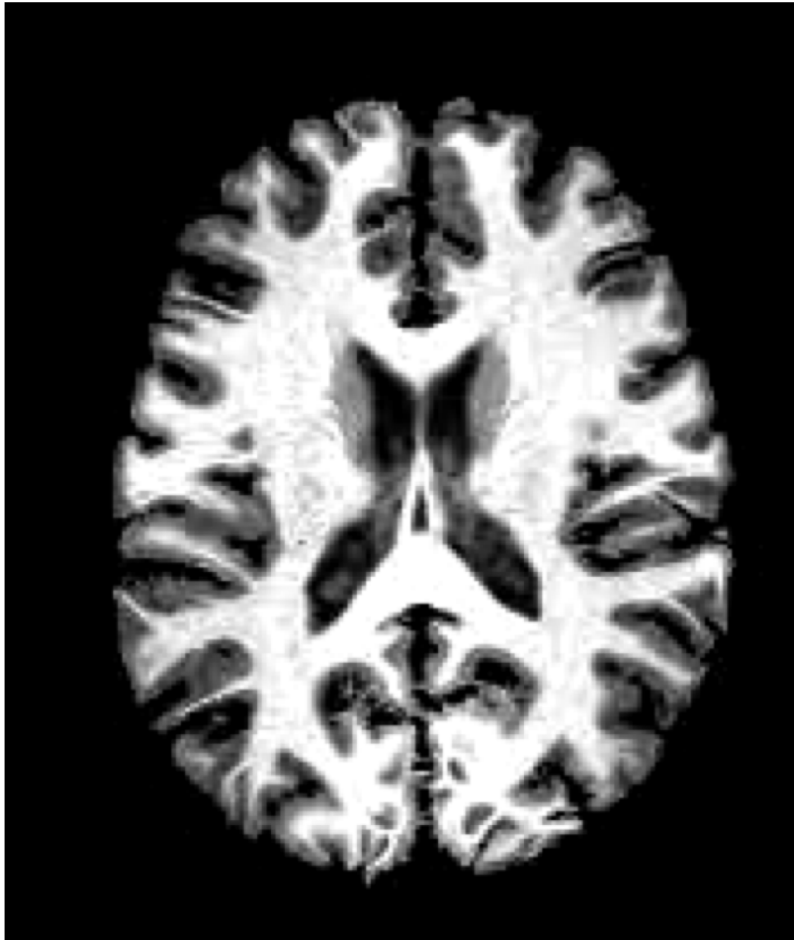
Post-surgical volume

Neurosurgery

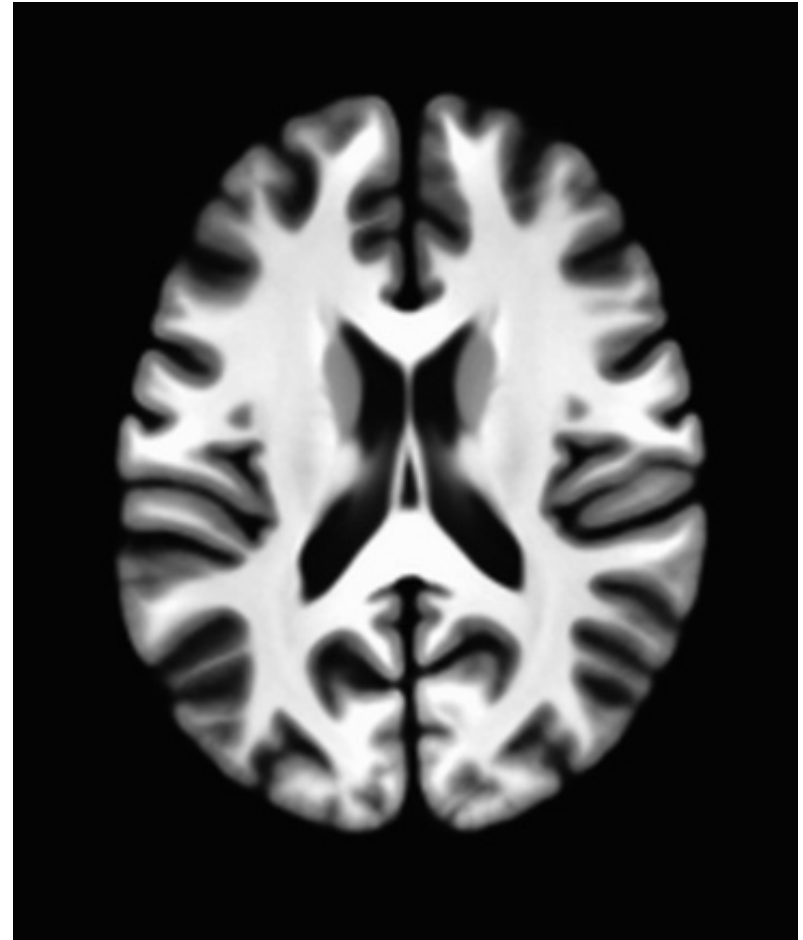


Pre-surgical volume
3dQwarp-aligned to Post-surgical volume

But ... Some Ugly



All 101 Volumes
After Warping



Mean of 101 Volumes
After Warping


How to Make a Template

- Given a collection of skull-stripped structural (T_1 -weighted) datasets
- Script *@toMNI_Awarp* pre-processes each dataset (*3dUnifize* and *@auto_tlrc*)
 - *3dUnifize* make the image intensity more uniform over the volume
- Script *@toMNI_Qwarp* runs *3dQwarp* to collectively warp them together over finer and finer patch levels
- Has been used to create Haskins pediatric brain atlas (now distributed with AFNI)

What Else to Do with a Warp?

- Warp another dataset the same way
 - *3dNwarpApply* (e.g., carry EPI to template)
- Warp some discrete points the same way
 - *3dNwarpXYZ* (e.g., eCog electrode locations)
- Compute voxel-wise functions of a warp
 - *3dNwarpFuncs* (e.g., volume distortion)
- Compose multiple warps together
 - *3dNwarpCat* and *3dNwarpCalc*
- Can compute inverse warp $W^{-1}(x)$, to map locations in $S(x)$ to matching locations in B
 - $S(W(x)) \approx B(x) \rightarrow S(x) \approx B(W^{-1}(x))$

How to Use *3dQwarp*

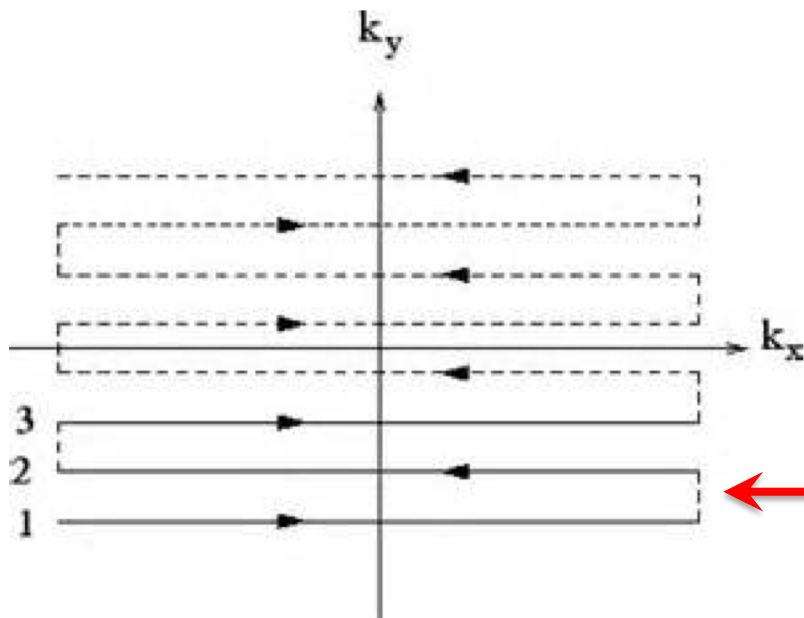
- Run it yourself (the “old school” or “real man” way)
- *auto_warp.py* (easier, less flexible)
-  Use *@SSwarper* script to **S**kull **S**trip and **W**arp to MNI template
- Use ‘*-tlrc_NL_warp*’ option in *afni_proc.py* to have transformation to template space be done via *auto_warp.py*
- Use *@toMNI_Awarp* and *@toMNI_Qwarpar* to create a study (or population) specific template
- Use ‘*-plusminus*’ option in *3dQwarp* to warp blip-up and blip-down EPI datasets to “meet in the middle” (script *unWarpEPI.py*)

Yet to Be Done

- ~~Incorporate more fully into *afni_proc.py* and *uber_subject.py*~~
 - ~~Warping to template; un-warping EPI distortions~~
- Explore how much nonlinear warping to a template can improve group analysis in functional and anatomical MRI
 - And improvements to *3dSeg* (segmentation)
- Extend matching algorithm to allow label-based matches, vs. existing intensity-based
- Speed the damn thing up!
- Write a paper about it!

Distortion Correction of EPI

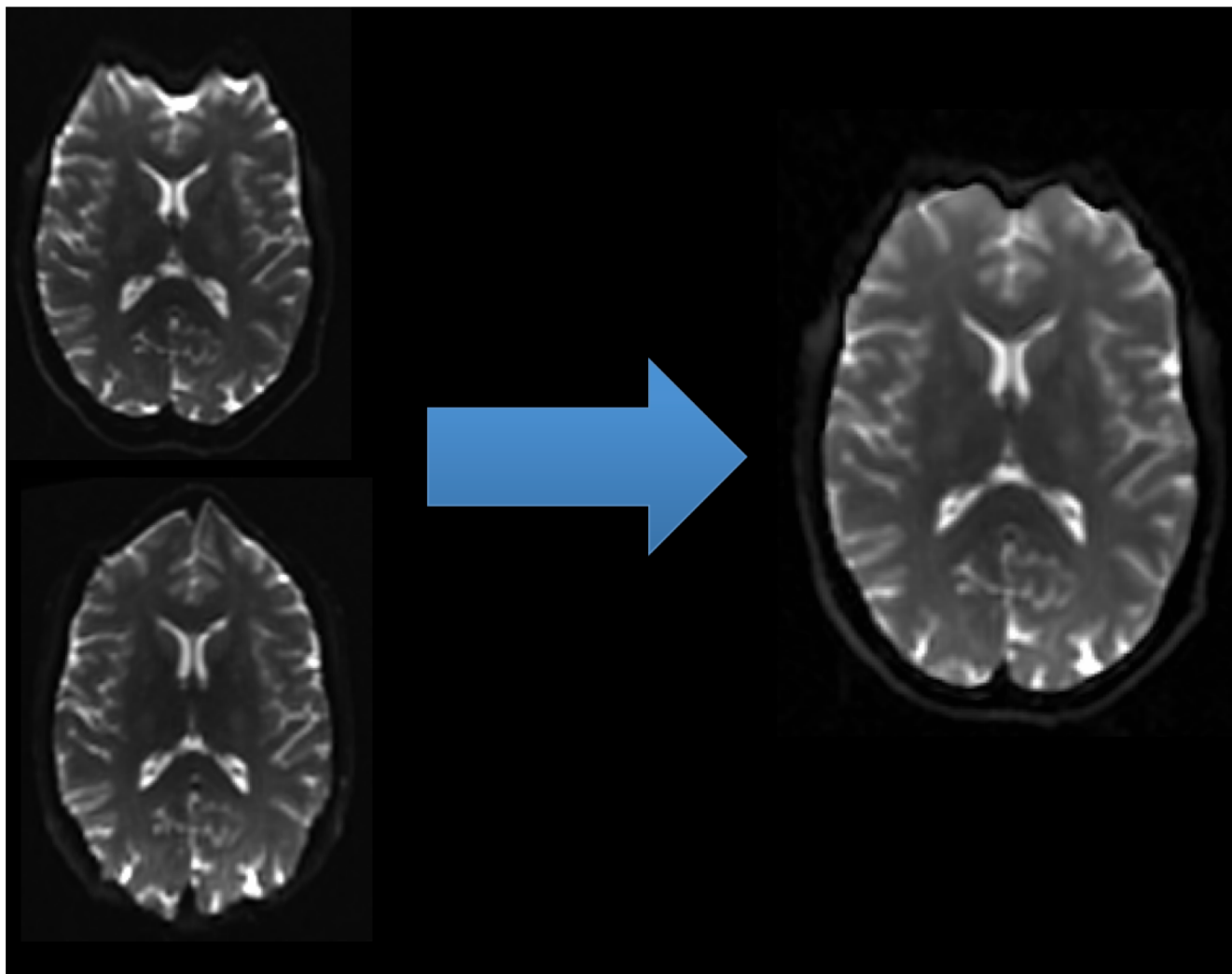
- Acquire a few 3D images with opposite phase encoding method (“blip up” and “blip down”)
 - This will reverse the distortions of the EPI data
 - You don’t need many opposite blip images
 - Ideally, also reverse the slice-selection gradient to also reverse the small slice-selection distortions



EPI scanning method
(in Fourier k-space)
"Blip up"

Distortion Correction of EPI

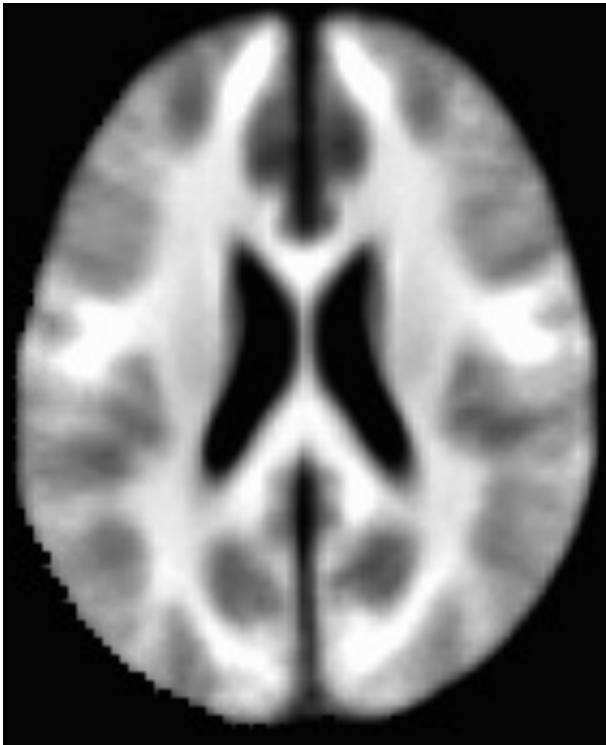
- Use *unWarpEPI.py* to fix the distortions



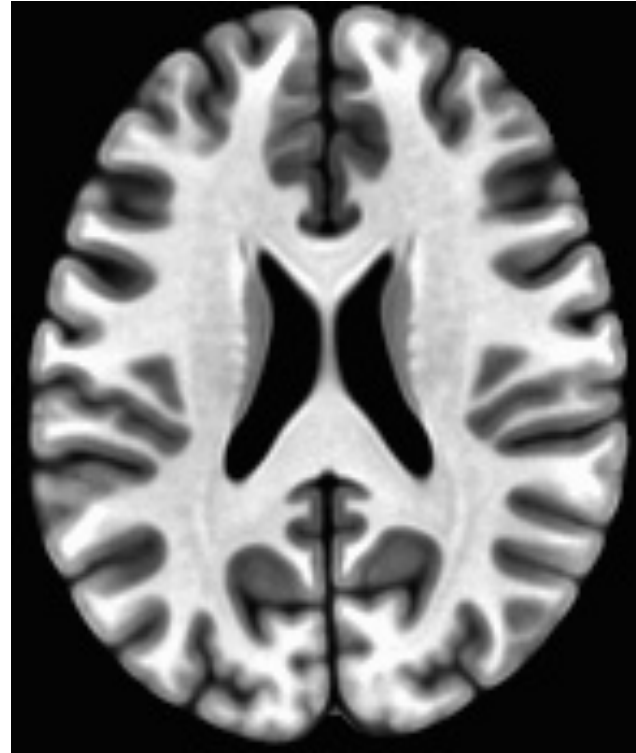
Nonlinear Warping to MNI Template

- afni_proc.py *can* do the nonlinear warping for you
 - But, nonlinear warping is slow (in fact, slowly slow)
 - If you need to re-rerun subject analysis, nonlinear warping will slow the re-run script down *a lot*
- Solution: do the nonlinear warping *before* using afni_proc.py, then supply the warping results so that afni_proc.py will skip doing the warping itself
- Mechanism: the **@SSwarper** script (tcsh)
 - Does Skull Stripping ("SS") and nonlinear warping
 - Base dataset is **MNI152_2009_template_SSW.nii.gz**
 - Nonlinearly warped, not too blurry

Two MNI Templates



MNI152_1mm_uni+tlrc
Affine alignments

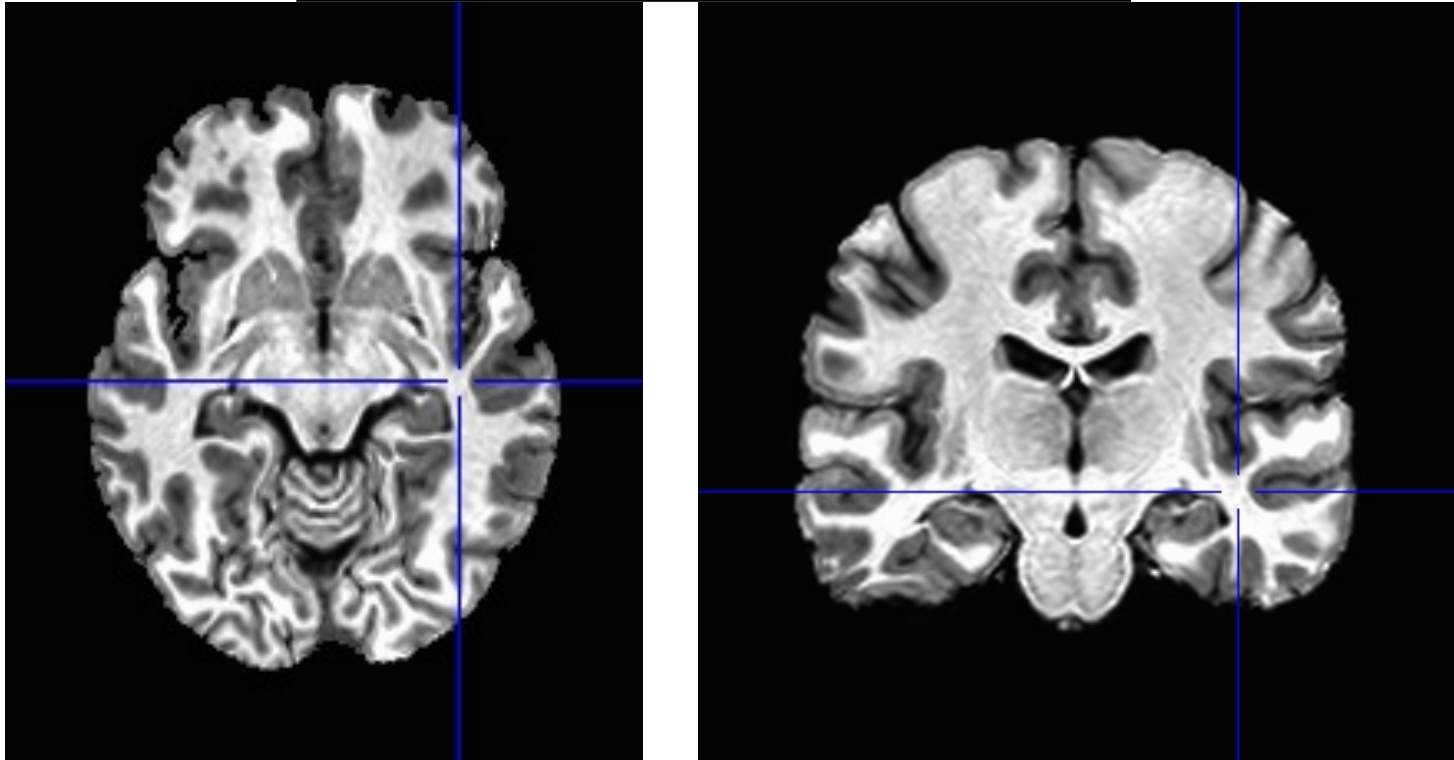


MNI152_2009_template.nii.gz
Nonlinear alignments

What @SSwarper Reads and Writes

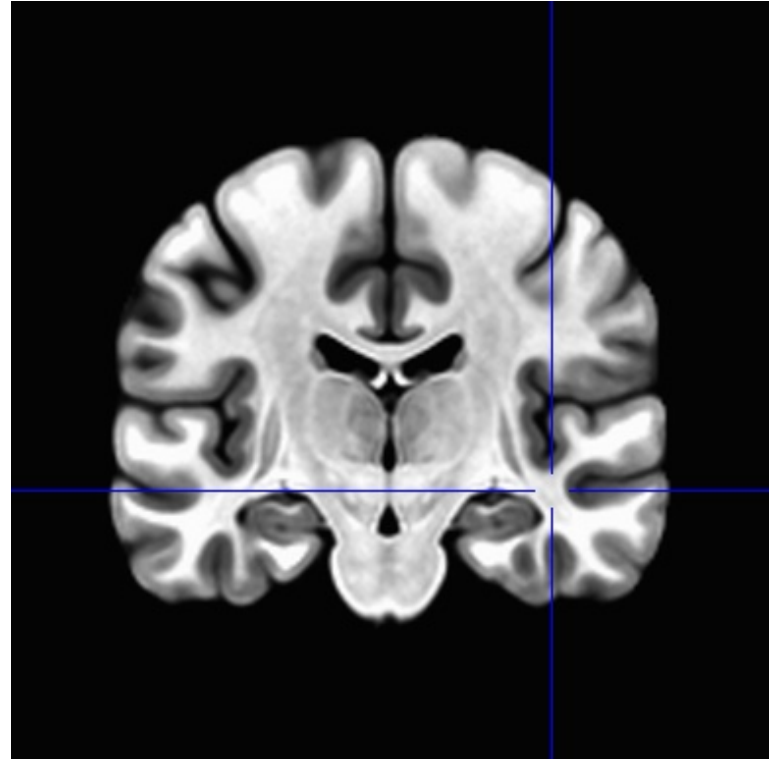
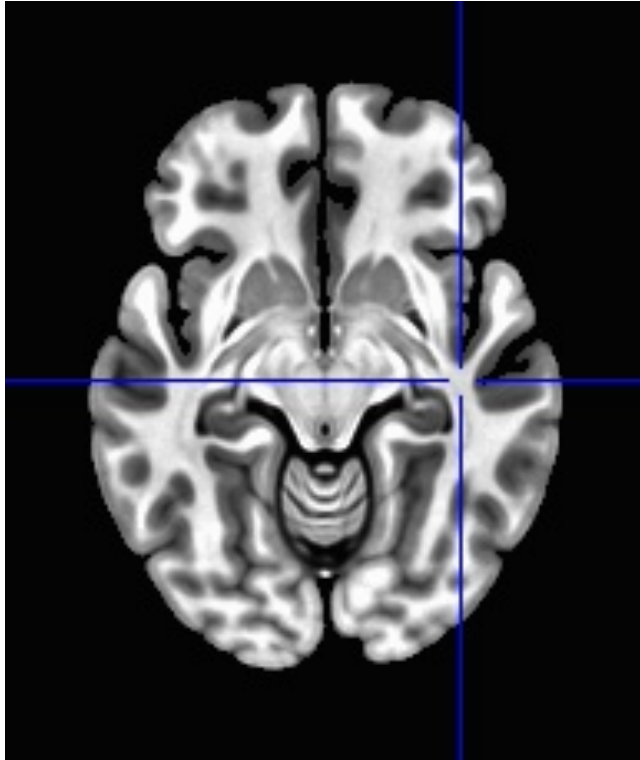
- Inputs:
 - T1-weighted anatomical image of subject (skull-on)
 - Subject ID code, for names of output files
- Outputs (subject ID = **sub007**):
 - `anatSS.sub007.nii`
 - skull-stripped dataset in original coordinates
 - `anatQQ.sub007.nii`
 - skull-stripped dataset, nonlinearly warped to MNI template
 - `anatQQ.sub007.aff12.1D`
 - affine matrix to transform original dataset to MNI template
 - `anatQQ.sub007_WARP.nii`
 - incremental warp from affine transformation to nonlinearly aligned dataset
- These files are needed for later use in `afni_proc.py`

@SSwarper Results



sub00440 from Beijing-Zang
in the FCON-1000 collection

MNI Template Slices



For comparison