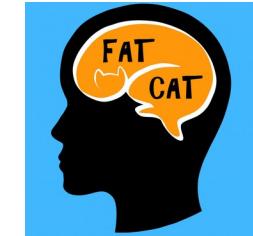


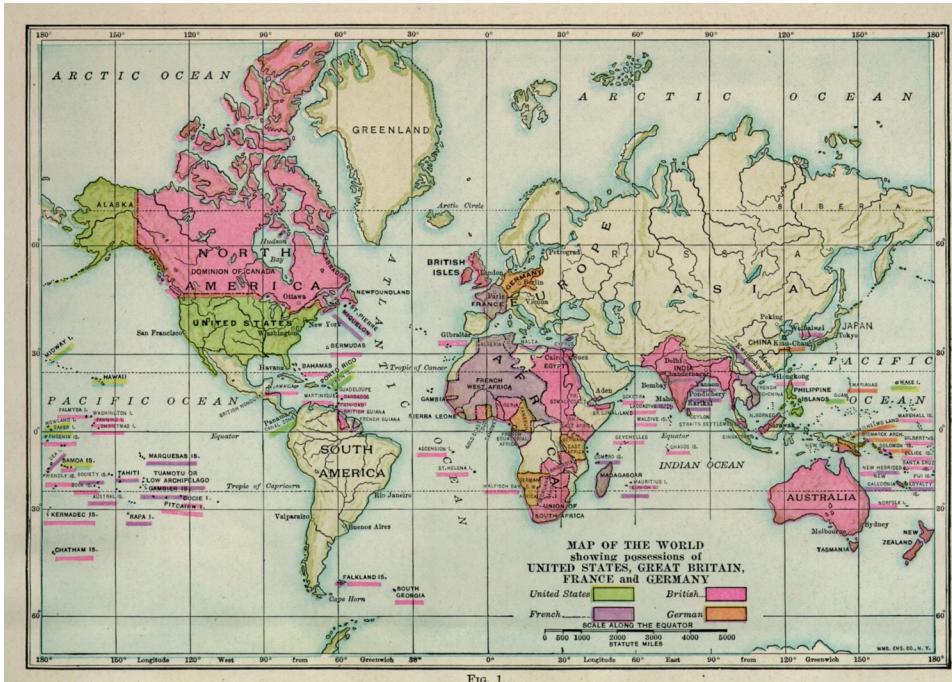


# AFNI



## Didactics and Demonstrations

### ATLASES AND TEMPLATES

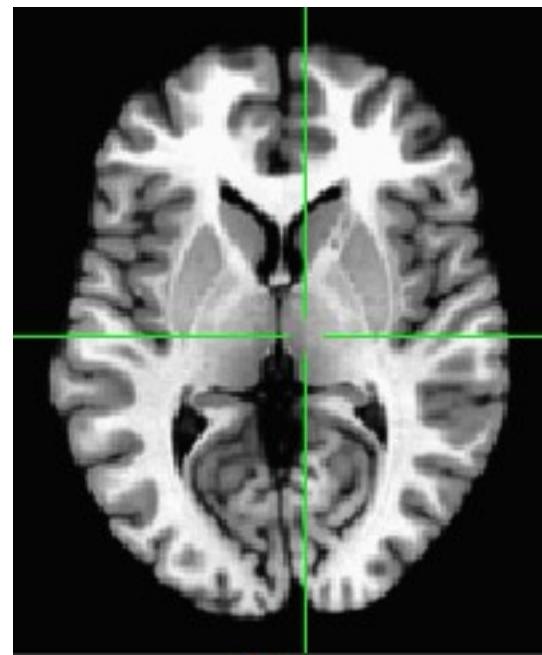


## Definitions

### Template

A reference dataset (typically whole brain) to which we align our subjects' data. We report coordinates of results relative to this template.

*Examples:* TT\_N27+tlrc, MNI152\_2009c+tlrc,  
HaskinsPeds\_NL\_template1.0\_SSW.nii.gz, D99\_macaque+tlrc



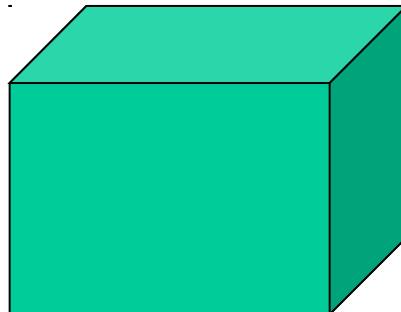
TT\_N27+tlrc

## Definitions

### Template Space

An (x, y, z) coordinate system shared by many datasets in alignment with a template.

Ex: TLRC (Talairach-Tournoux), MNI, MNI\_ANAT, ORIG.



To see what “space” a dset is in, type:

```
3dinfo -space DSET_NAME
```

## Definitions

**A note on AFNI usage of “tlrc”:**

AFNI format “+tlrc” view - means in a "standard" space -  
e.g. Talairach-Tournoux, MNI, pediatric template,  
macaque, ...

anat+tlrc.HEAD, anat+tlrc.BRIK

**Space names**

MNI, TLRC, ....

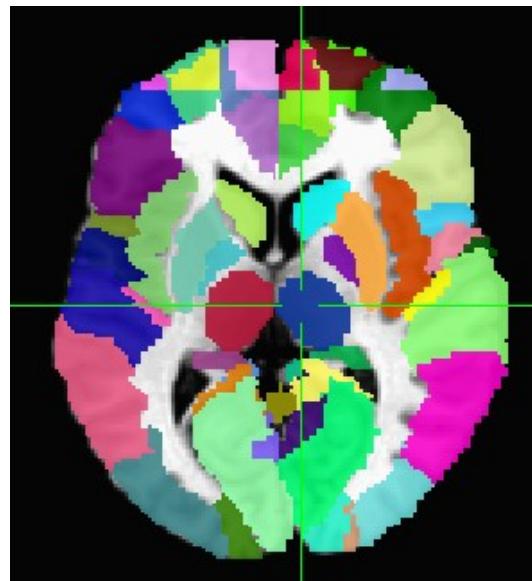
Lots of variants - specify the template explicitly in your  
papers

## Definitions

### Atlas

A dataset containing segmentation or parcellation information. It can be considered a “map” of ROIs: each ROI is defined as a set of voxels with a certain integer value (and a string label can be attached to each ROI).

Ex: TTatlas+tlrc, TT\_N27\_EZ\_ML+tlrc, my\_roidset+orig.

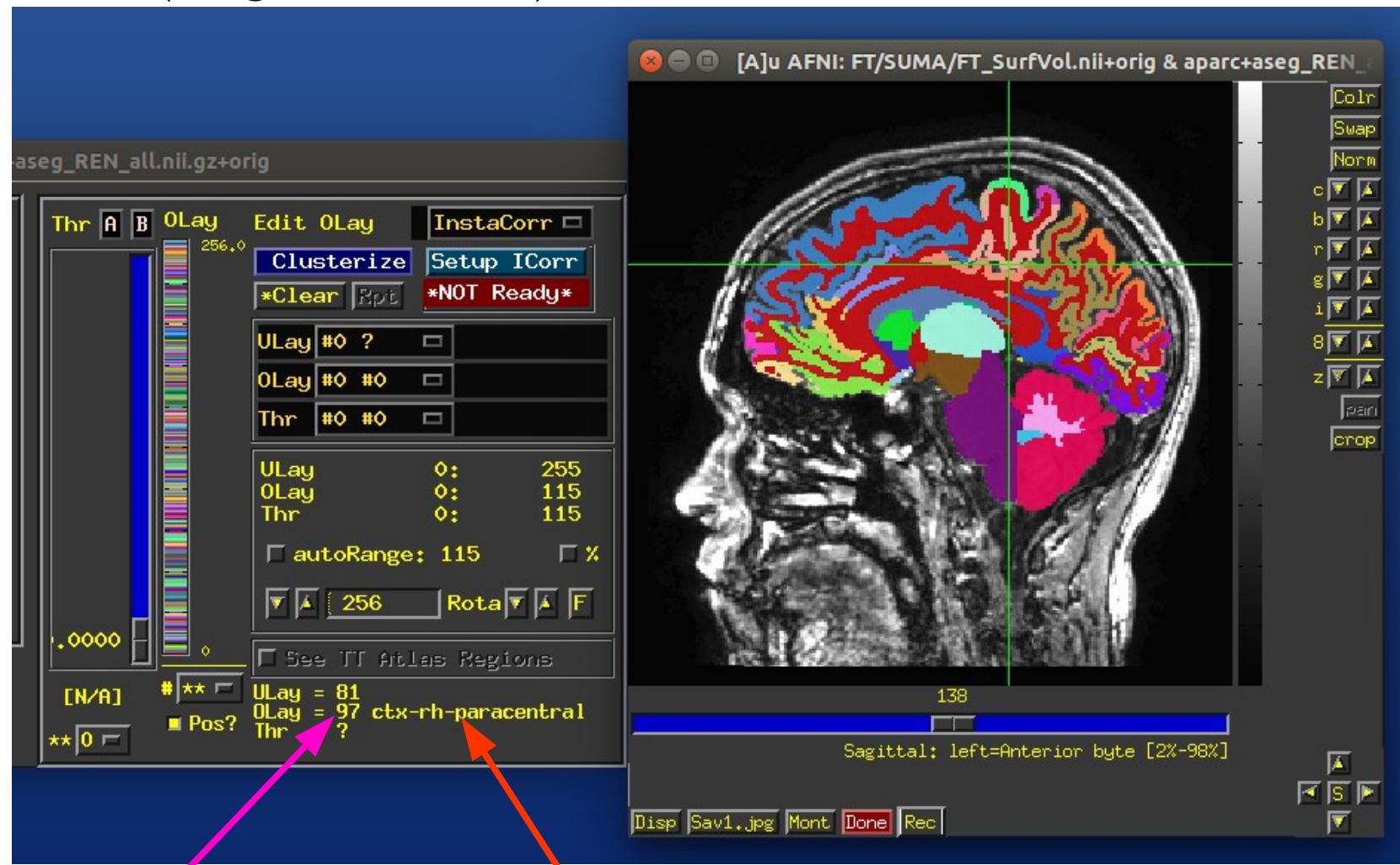


TT\_N27\_EZ\_ML+tlrc

*See more description about templates+atlases (including making your own) on the AFNI website:  
[https://afni.nimh.nih.gov/pub/dist/doc/html/doc/template\\_atlas/framework.html](https://afni.nimh.nih.gov/pub/dist/doc/html/doc/template_atlas/framework.html)*

# Definitions

## Atlas (or general ROI) label in GUI

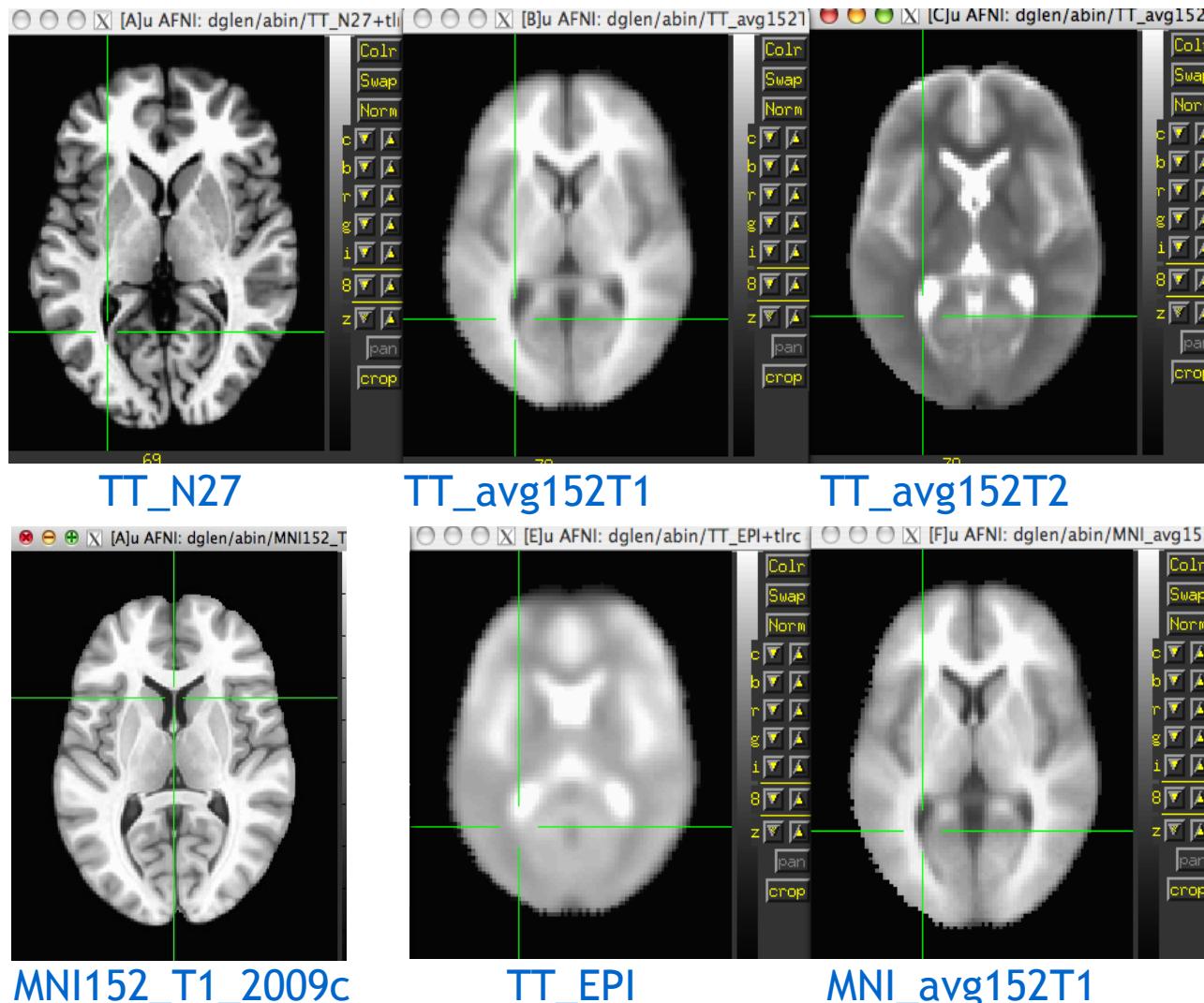


integer value of ROI

associated string label

## Templates included with AFNI

After default AFNI installation, these templates (and others) would be in `~/abin/`:



*And a quick question: what important properties does each dset here have?*

## Templates included with AFNI

A helpful note on viewing templates (or any dsets) **each time** you open up the AFNI GUI, regardless of directory!

Set the **AFNI\_GLOBAL\_SESSION** variable in your `~/.afnirc` file, e.g.,:

```
AFNI_GLOBAL_SESSION = /home/nmandela/abin
```

All dsets there will appear in your “Underlay” or “Overlay” menu in the AFNI GUI.  
(Use full path of directory, no “~” or “\$HOME”.)

## Templates included with AFNI

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```
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```

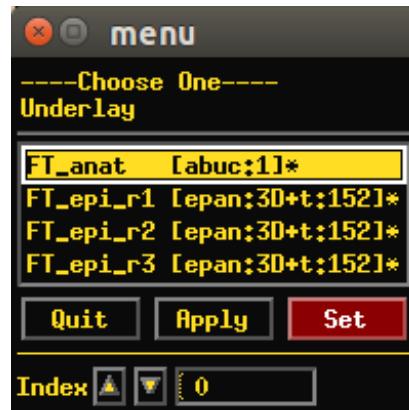
All dsets there will appear in your “Underlay” or “Overlay” menu in the AFNI GUI.  
(Use full path of directory, no “~” or “\$HOME”.)

*Ex:* then open AFNI GUI in some directory, say:

```
~/AFNI_data6/FT_analysis/FT/
```

*after* →

*before* →



## Standard spaces

**Reasons to use a standard template space:**

- Compare across subjects and groups easily for every voxel in the brain
- Standardize coordinates with others
- Know where a voxel is automatically from an atlas
- Mostly automated and no specific ROI drawing required

**Reasons not to use a standard template space:**

- Inconsistency among subjects
- Inconsistency among groups - elderly versus younger
- Use consistent anatomical ROIs with good anatomical knowledge
- Lower threshold for multiple comparison adjustments

## "Standard" spaces - Some History

Talairach - Tournoux - 1988 (1957)

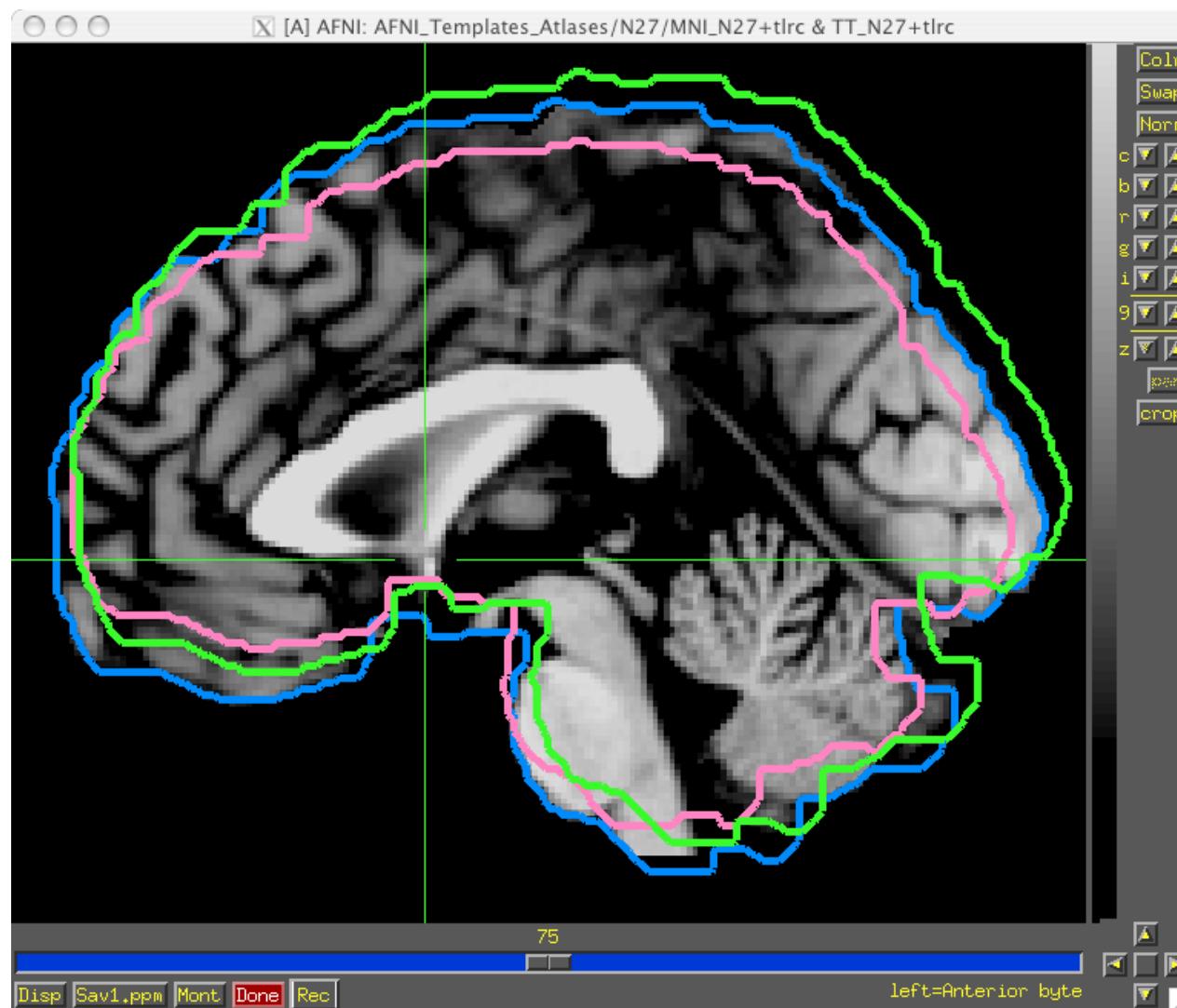
- Stereotaxic system - no MRI template
- One of the first uses of AFNI ! (1994)
- Brodmann atlas (1907-09) UTHSCSA - Talairach Daemon (1997)

MNI - 1992 - 2009

- MNI-305 1992 - Approximately Talairach
- MNI-N27 1998 - Single brain affinely aligned to MNI-305
- MNI-152 - 2001, linear affine (ICBM-UCLA, MNI, UTHSCSA)
- MNI-152 - 2006, nonlinear, symmetric/asymmetric (FSL and maybe SPM too use asymmetric version)
- MNI-152 - 2009, nonlinear 1mm, 0.5mm
  - a,b,c x2 (symmetric/asymmetric) AFNI includes 2009c-asymmetric 1mm3 (unifized and original)
- MNI-Anat - 2005, SPM Anatomy toolbox - Eickhoff-Zilles Cytoarchitectonic Atlas and Macrolabel atlas. Shifted coordinate system for N27. 2020 update, atlases for version 3.0 moves to MNI152-2009?

## Template spaces differ in origin

TLRC, MNI, and MNI-Anat



## Template spaces differ in size

The **MNI** brain is larger than the **TLRC** brain.



## From space to space

### For going between TLRC and MNI:

Approximate equation

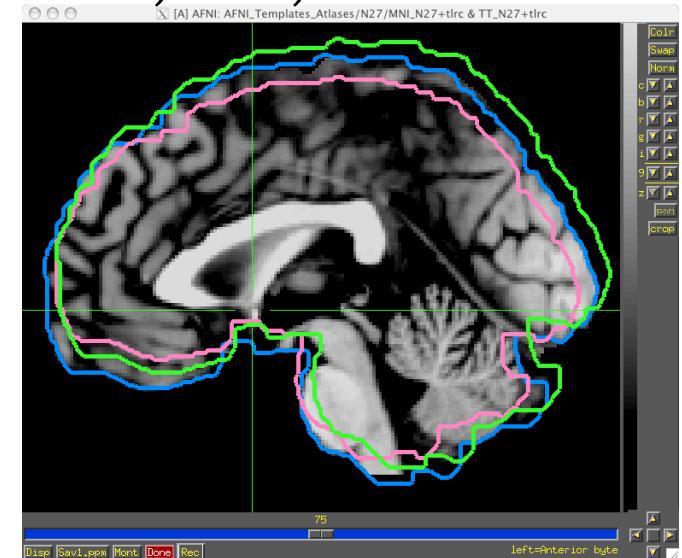
→ used by **whereami** and **3dWarp**

A manual TLRC transformation of MNI template to TLRC space

→ used by **whereami** (as precursor to MNI Anat.), based on N27 template

Multiple space coordinates reported in **whereami** output  
**(AFNI\_ATLAS\_TEMPLATE\_SPACE\_LIST)**

### TLRC, MNI, and MNI-Anat



### For going between MNI and MNI Anat (Eickhoff et al., 2005):

MNI + (0, 4, 5) = MNI Anat. (in RAI coordinate system)

### Going between TLRC and MNI Anat (as practiced in whereami):

Go from TLRC (TT\_N27) to MNI via  
manual transform of N27 template

Add ( 0, 4, 5 )

## Choosing a template

Try to pick a template that...

- is similar to the subject group: neonates, pediatric, young adults, elderly, macaque, rabbit...
- is of the same modality and coverage as your data sets
- has a relevant atlas segmentation.

## Choosing a template

Try to pick a template that...

- is similar to the subject group: neonates, pediatric, young adults, elderly, macaque, rabbit...
- is of the same modality and coverage as your data sets
- has a relevant atlas segmentation.

You can also make your own template (and maybe an atlas too):

- Scripts/commands exist in AFNI
  - Ex. Haskins pediatric atlas
  - several methods tested
  - best approach: *iterative nonlinear alignment*
    - `make_template_dask.py`
    - **Previously @toMNI\_Awarp,  
@toMNI\_Qwarpar**

# Finding templates and atlases

Good places to find templates and atlases

AFNI!- abin directory

AFNI! - <http://afni.nimh.nih.gov:/pub/dist/atlas>

NITRC - <https://www.nitrc.org/>

PRIME-RE - <https://prime-re.github.io/>

Allen Brain Project -

<https://portal.brain-map.org/>

BrainMaps.org - <http://brainmaps.org/>

MNI - <http://nist.mni.mcgill.ca>

Scalable Brain -

<https://scalablebrainatlas.incf.org>

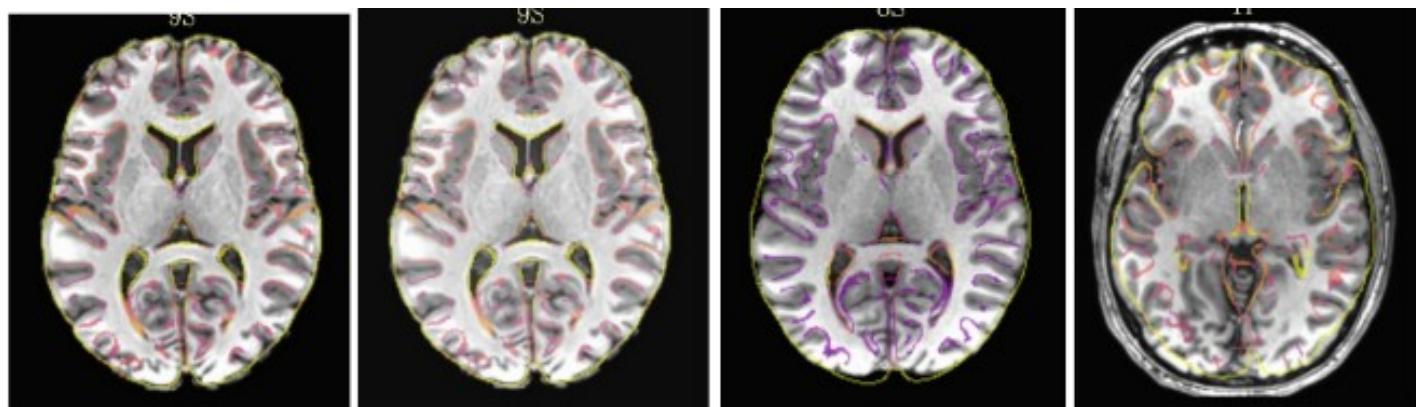
UCLA - <http://www.bmap.ucla.edu/portfolio/atlas>

afni.nimh.nih.gov:/pub/dist/atlas	
dir	2017.11.21.2129
dir	2019.04.11.1640
dir	2019.04.07.2131
file	2012.09.04.1626
dir	2019.09.24.1357
dir	2020.01.24.1623
dir	2019.04.05.1617
file	2019.03.14.1818
dir	2017.11.21.2129
file	2014.10.22.1800
dir	2017.11.21.2129
dir	2020.05.05.1745
dir	2017.10.25.1407
dir	2019.07.22.1638
dir	2020.04.29.1719
dir	2017.11.21.2129
dir	2017.03.27.1023
dir	2013.04.16.1734
file	2013.04.16.1738
dir	2019.04.19.1846
dir	2017.11.21.2129
dir	2019.04.03.1224
dir	2019.01.24.1816
dir	2019.12.12.2304
dir	2019.07.30.1723
dir	2020.04.30.2132
dir	2017.11.21.2129
dir	2019.07.30.1718
dir	2017.11.21.2129
dir	2017.11.21.2130
	<a href="#">bnsl</a>
	<a href="#">brainnetome</a>
	<a href="#">caez</a>
	<a href="#">caez_18_all_atlases.tgz</a>
	<a href="#">cat</a>
	<a href="#">current</a>
	<a href="#">current_pmaps</a>
	<a href="#">current.tgz</a>
	<a href="#">elsedemo</a>
	<a href="#">Haskins_MNI152.jpg</a>
	<a href="#">HaskinsPeds</a>
	<a href="#">IBT</a>
	<a href="#">keukens</a>
	<a href="#">macaque</a>
	<a href="#">marmoset</a>
	<a href="#">MNI</a>
	<a href="#">mni_acpc</a>
	<a href="#">MNI_atlasses</a>
	<a href="#">MNIatlases.tgz</a>
	<a href="#">MNI_HCP</a>
	<a href="#">MNI_VmPFC</a>
	<a href="#">mouse</a>
	<a href="#">Pauli_subcortical</a>
	<a href="#">Princeton_visual</a>
	<a href="#">rat</a>
	<a href="#">SchaeferYeo</a>
	<a href="#">SUIT_Cerebellum</a>
	<a href="#">UNC_Infant_Atlases</a>
	<a href="#">VmPFC</a>
	<a href="#">waxrat</a>

## Trading Spaces

How to transform data to a standard template space

- **@SSwarper** - skullstrip and align data to some select standard spaces
- **@animal\_warper** - skullstrip, align data to animal template and bring atlas into native space
- **auto\_warp.py** - combination affine and nonlinear alignment to a template
- **@auto\_tlrc** - affine alignment to a template
- Manual talairach



@SSW

AW

@at

manual

## Nonlinear alignment to template

### Multiple nonlinear alignment tools in AFNI

**3dQwarp:** the standard nonlinear workhorse

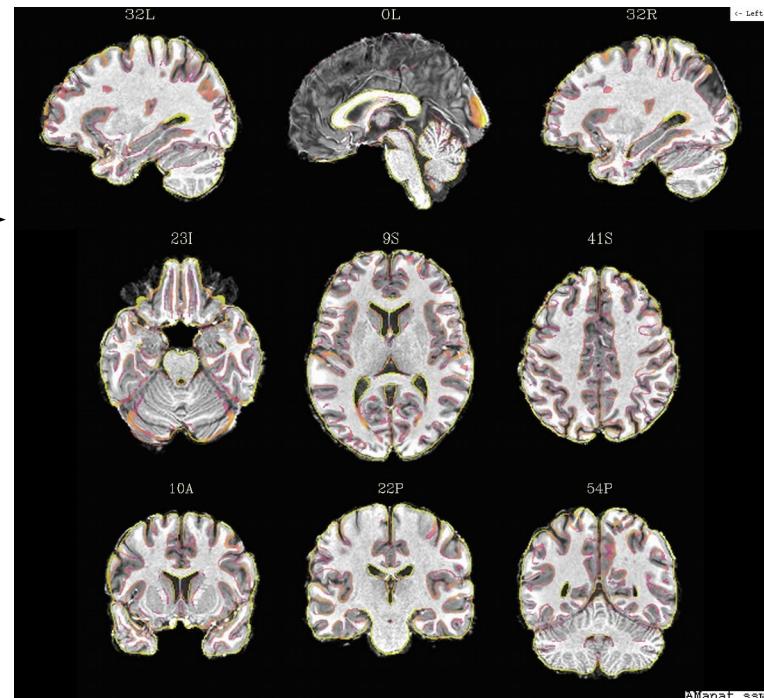
**@SSwarper:** skull stripping and alignment in one-- and bonus automatic QC images

+ @SSwarper uses a multi-volume base; several exist for standard templates already, and more can be made-- see full description for these online:

[https://afni.nimh.nih.gov/pub/dist/doc/html/doc\\_template\\_atlas/sswarper\\_base.html](https://afni.nimh.nih.gov/pub/dist/doc/html/doc_template_atlas/sswarper_base.html)

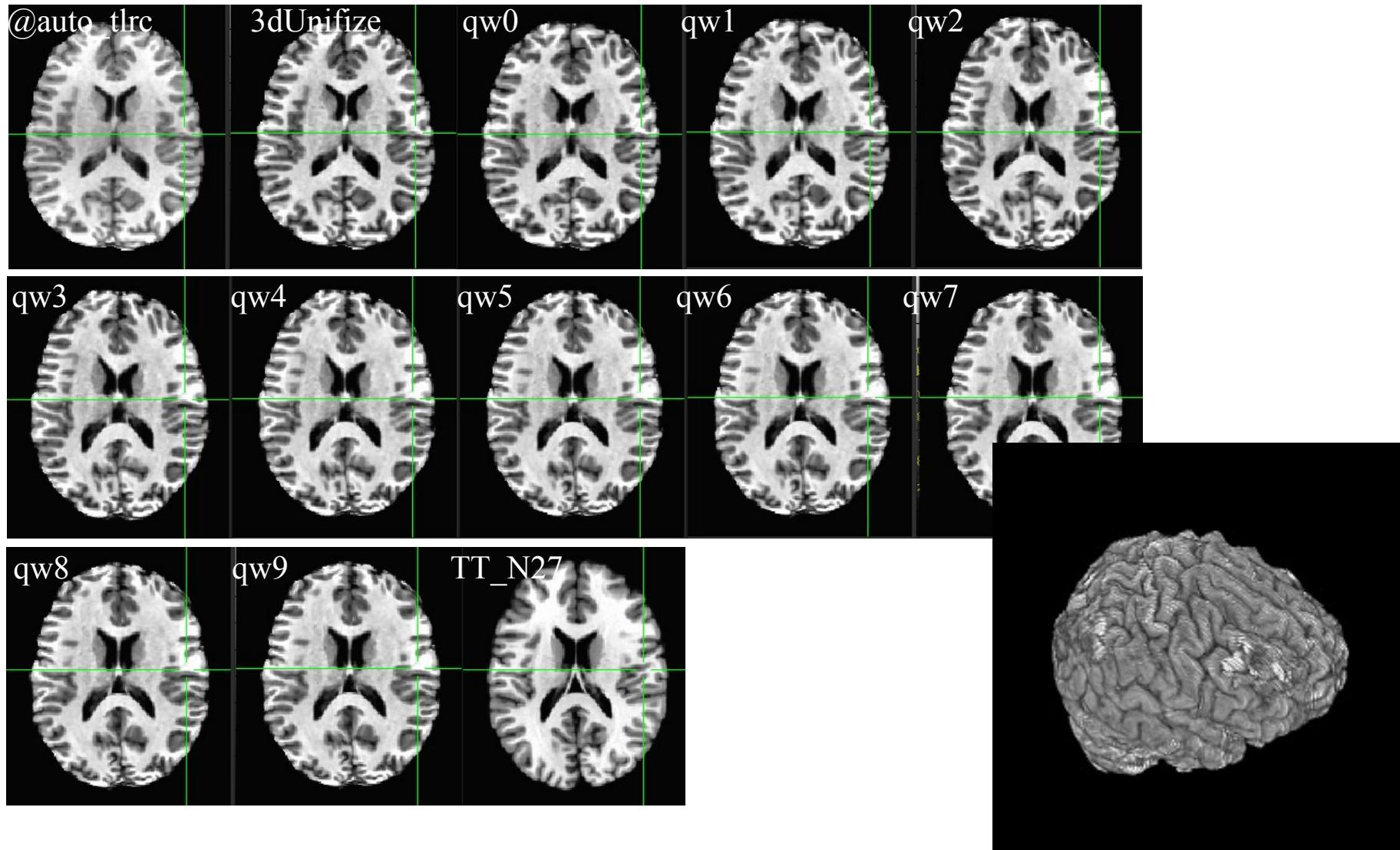
**auto\_warp.py:** wrapper for alignment functionality, ~simpler syntax (@auto\_tlrc + 3dQwarp together)

Ex: **auto\_warp.py -base MNI152\_T1\_2009c+tlrc. \ -suffix \_awarp -input striptorig.**



## Nonlinear alignment to template

3dQwarp, through multiple levels of refinement →



# Trading Spaces

## **afni\_proc.py options**

tlrc block before volreg!

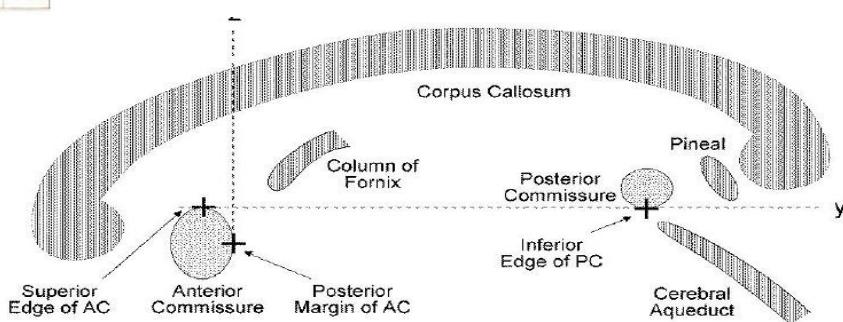
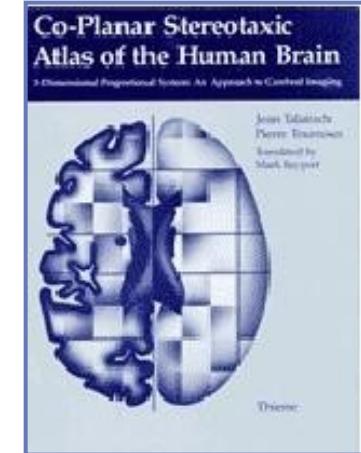
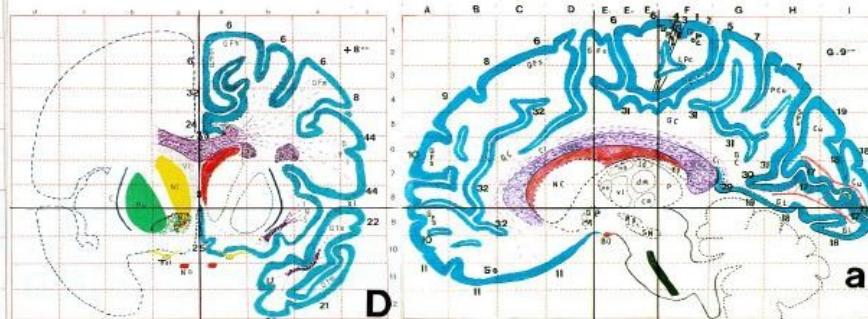
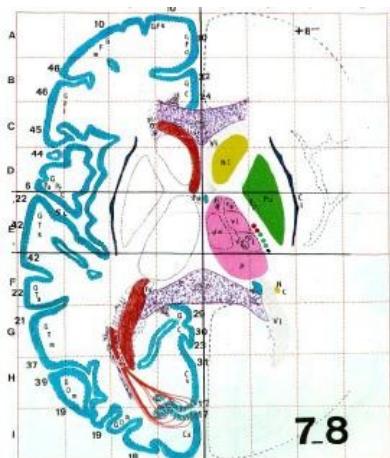
**-volreg\_tlrc\_warp**  
**-tlrc\_base** \$tpath/\$btemplate (runs @auto\_tlrc)  
**-tlrc\_NL\_warp** (runs auto\_warp.py)

For @SSwarper output, add these:

**-copy\_anat** anatSS.\${subj}.nii \\\n**-anat\_has\_skull** no \\\n**-tlrc\_NL\_warp** \\\n**-tlrc\_NL\_warped\_dsets** \\\n    anatQQ.\${subj}.nii \\\n    anatQQ.\${subj}.aff12.1D \\\n    anatQQ.\${subj}\_WARP.nii

# Space Travel - Getting to Talairach Space

- The original purpose of AFNI transform datasets to Talairach-Tournoux (stereotaxic) coordinates
- Manual mode - mark various anatomical locations, defined in  
Jean Talairach and Pierre Tournoux  
“Co-Planar Stereotaxic Atlas of the Human Brain”  
Thieme Medical Publishers, New York, 1988,  
1957 (original version introducing AC-PC stereotaxic system)



## Automatic Talairach transform (affine) with `@auto_tlrc`

- In the olden days, people would spend a lot of time transforming data to standard space by hand (see Supplement slides for how to perform the Manual TLRC transform using the AFNI GUI by setting AC-PC landmarks).
- Here, we describe how to perform a TLRC transform *automatically* using AFNI's `@auto_tlrc` (used by `afni_proc.py` for linear affine alignment to standard space).
  - ◊ Differences from Manual Transformation:
    - Instead of setting AC-PC landmarks and volume boundaries by hand, the anatomical volume is warped (using 12-parameter affine transform) to a template volume in TLRC space.
    - The Anterior Commissure (AC) center is no longer at  $xyz = (0,0,0)$ , and the size of brain box is that of the template you use.
      - ◊ For various reasons, some good and some bad, templates adopted by the neuroimaging community are not all of the same size. Be mindful when using various atlases or comparing standard-space coordinates.
    - You, the user, can choose from various templates for reference (just be consistent in your group analysis).
    - It is easy + automatic. Just check final results to make sure nothing went seriously awry.

## @auto\_tlrc example

- To run in AFNI\_data6/afni/  
Transform the subj anat to a template space (output: **anat\_TT+tlrc**):

```
@auto_tlrc  
-base TT_N27+tlrc  
-suffix _TT  
-input anat+orig
```



- Then apply the transform stored in **anat\_TT+tlrc**'s header to a “follower dset” (here, func data), specifying output resolution at 3 mm (output: **func\_slim\_TT+tlrc**):

```
@auto_tlrc  
-apar anat_TT+tlrc  
-input func_slim+orig  
-suffix _TT  
-dxyz 3
```



- Instead of **TT\_N27**, you could also use the **icbm452**, or the **mni**'s **avg152T1** template, or any other template you like  
(see **@auto\_tlrc -help** for a few good words on templates)

# That's All for Now

