

## 1 SUPPLEMENTARY DATA

### Data augmentation ranges for Dense-Unet training

Rotation: [-10, 10]

Shearing: [-0.1, 0.1]

Scaling: [0.9, 1.1]

### ANTs registration command for T1 to diffusion space using b0 and FA images.

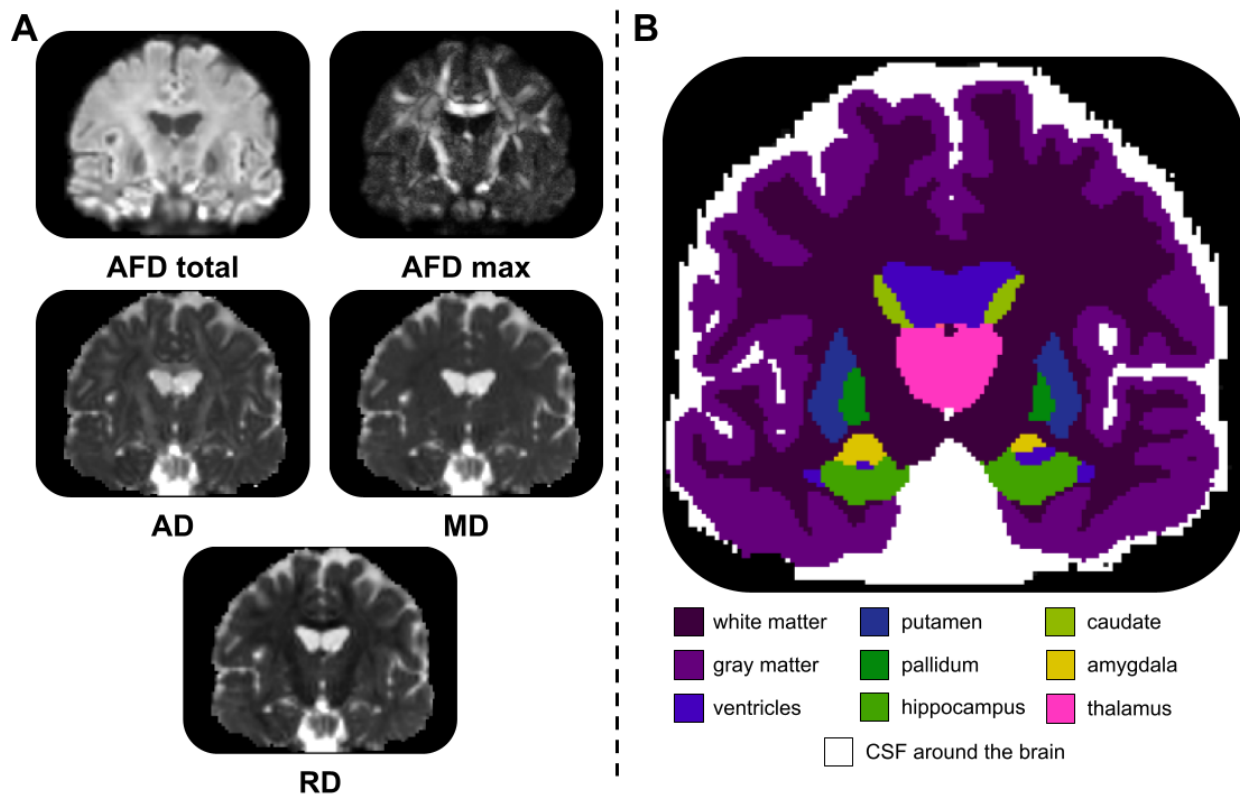
```
antsRegistration --dimensionality 3 --float 0\
  --output [output,outputWarped.nii.gz,outputInverseWarped.nii.gz]\
  --interpolation Linear --use-histogram-matching 0\
  --winsorize-image-intensities [0.005,0.995]\
  --initial-moving-transform [b0.nii.gz,t1.nii.gz,1]\
  --transform Rigid['0.2']\
  --metric MI[b0.nii.gz,t1.nii.gz,1,32,Regular,0.25]\
  --convergence [500x250x125x50,1e-6,10] --shrink-factors 8x4x2x1\
  --smoothing-sigmas 3x2x1x0\
  --transform Affine['0.2']\
  --metric MI[$b0,$t1,1,32,Regular,0.25]\
  --convergence [500x250x125x50,1e-6,10] --shrink-factors 8x4x2x1\
  --smoothing-sigmas 3x2x1x0\
  --transform SyN[0.1,3,0]\
  --metric MI[b0.nii.gz,t1.nii.gz,1,32]\
  --metric CC[fa.nii.gz,t1.nii.gz,1,4]\
  --convergence [50x25x10,1e-6,10] --shrink-factors 4x2x1\
  --smoothing-sigmas 3x2x1

antsApplyTransforms -v -d 3 -n NearestNeighbor -i wmparc.nii.gz\
  -r b0.nii.gz -t output1Warp.nii.gz -t output0GenericAffine.mat\
  -o wmparc_registered.nii.gz

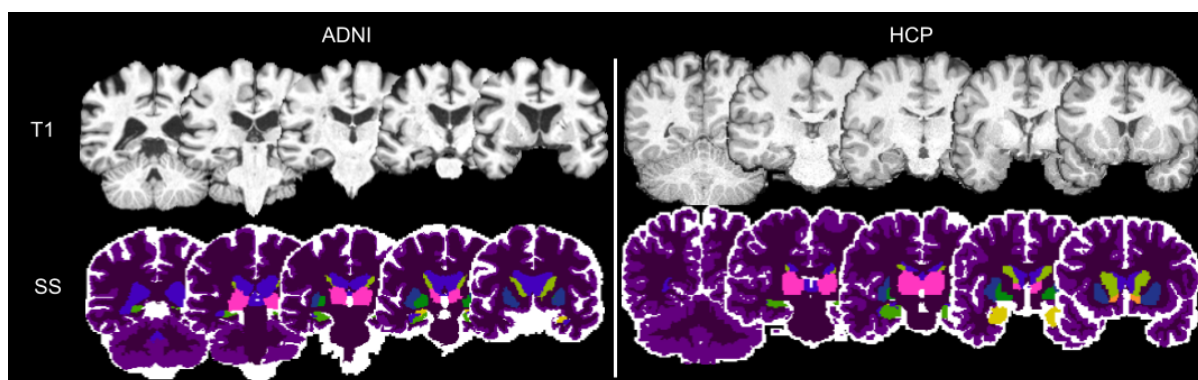
antsApplyTransforms -v -d 3 -n NearestNeighbor -i aparc+aseg.nii.gz\
  -r b0.nii.gz -t output1Warp.nii.gz -t output0GenericAffine.mat\
  -o aparc_aseg_registered.nii.gz
```

This command is based on the one used in TractoFlow.

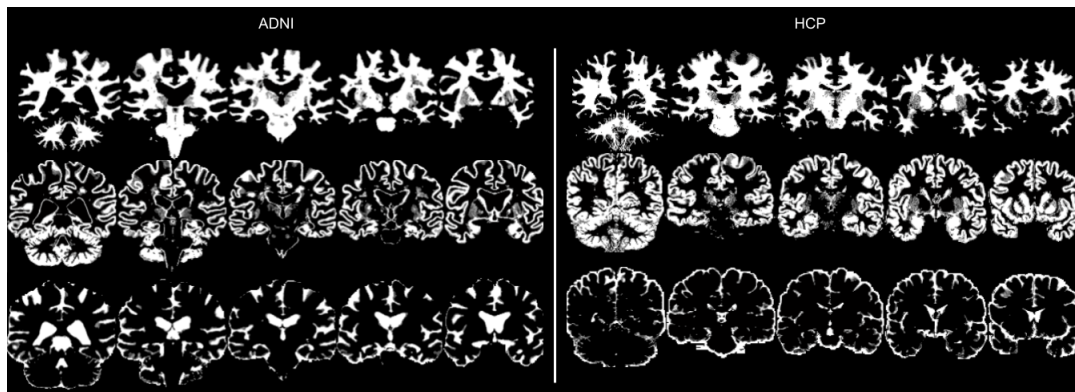
## 2 SUPPLEMENTARY TABLES AND FIGURES



**Figure S1.** In A, the 5 images used as input channel. In B, the 10 tissue class segmentation.



**Figure S2.** Silver standard image from Freesurfer on ADNI and HCP subjects.



**Figure S3.** ADNI and HCP subjects segmentation from FSL-Fast



**Figure S4.** Nuclei mosaic for ADNI and HCP subjects from DORIS.

**Table S1.** Related t-test for all label volumes between DORIS and the silver standard.

| Label       | T                  | p-value              |
|-------------|--------------------|----------------------|
| WM          | -20.619754316231   | 1.87002304545186E-31 |
| GM          | 5.32398808768571   | 1.16487986447687E-06 |
| Ventricles  | -16.3983266370104  | 1.12776835129094E-25 |
| Putamen     | 7.24984563457834   | 4.36694047443652E-10 |
| Pallidum    | -17.4565807508637  | 3.30797195260101E-27 |
| Hippocampus | -0.819219532230748 | 0.415441646579405    |
| Caudate     | -22.1378407131987  | 2.48624959519249E-33 |
| Amygdala    | -1.4838719980424   | 0.142331888075401    |
| Thalamus    | -6.06896720284198  | 5.91698289223593E-08 |

**Table S2.** Volume (in mm<sup>3</sup>) of smallest labels for DORIS and the silver standard

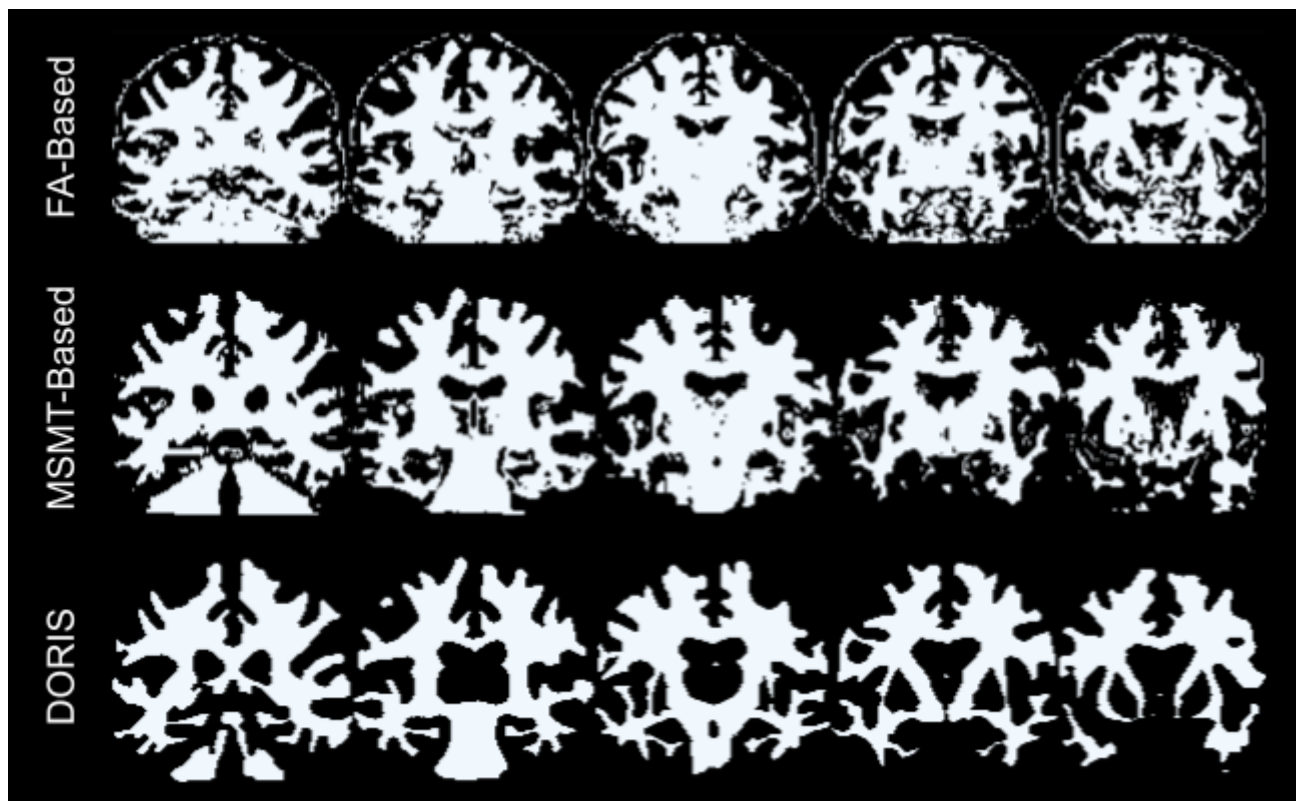
| type            | label       | volume       |
|-----------------|-------------|--------------|
| DORIS           | Amygdala    | 3814 (303)   |
| DORIS           | Caudate     | 7563 (750)   |
| DORIS           | Hippocampus | 8432 (578)   |
| DORIS           | Pallidum    | 3100 (559)   |
| DORIS           | Putamen     | 9616 (1137)  |
| DORIS           | Thalamus    | 14137 (1173) |
| DORIS           | Ventricles  | 22908 (4876) |
| silver standard | Amygdala    | 3741 (452)   |
| Silver standard | Caudate     | 6628 (750)   |
| Silver standard | Hippocampus | 8389 (809)   |
| Silver standard | Pallidum    | 2322 (358)   |
| Silver standard | Putamen     | 10040 (1162) |
| Silver standard | Thalamus    | 13652 (1068) |
| Silver standard | Ventricles  | 20477 (5003) |

**Table S3.** Related t-test for DORIS volumes in test-retest

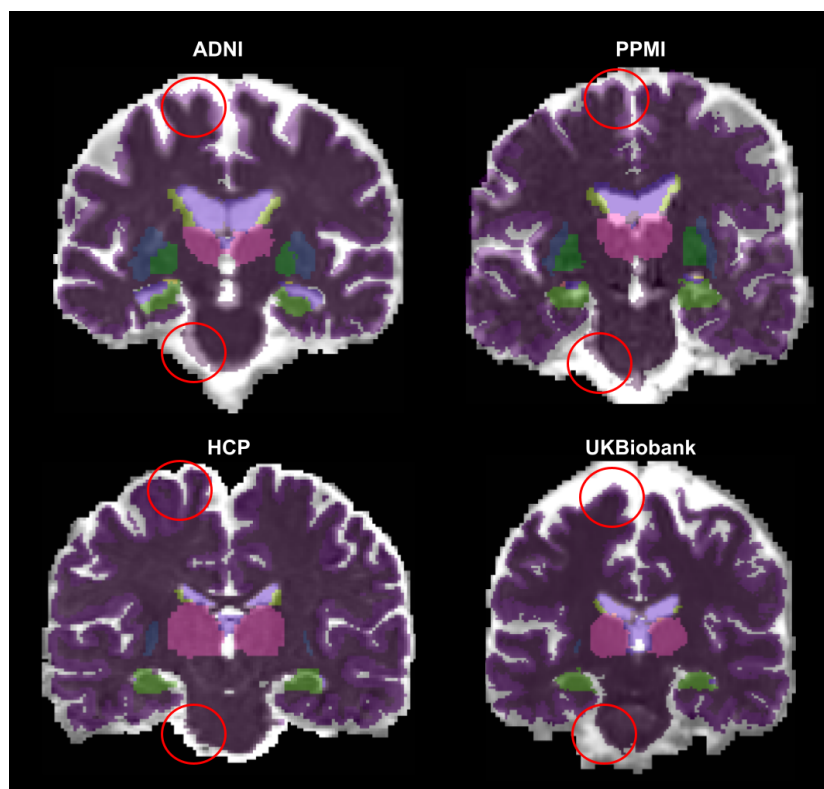
| Label       | T                  | p-value           |
|-------------|--------------------|-------------------|
| WM          | 0.272086349464586  | 0.790594903814357 |
| GM          | -0.606080243692412 | 0.556761393593017 |
| Ventricles  | -0.684811358828159 | 0.50763586210935  |
| Putamen     | 0.724749503407634  | 0.483737254912877 |
| Pallidum    | 0.885461971013694  | 0.394851868599488 |
| Hippocampus | -0.539277357153973 | 0.600440444292768 |
| Caudate     | 0.530965542463882  | 0.605997081871768 |
| Amygdala    | -0.346666373623647 | 0.735382752665228 |
| Thalamus    | 1.74974259161594   | 0.107968596843501 |

**Table S4.** Related t-test for the silver standard volumes in test-retest

| Label       | T                  | p-value            |
|-------------|--------------------|--------------------|
| WM          | 0.0864547032459438 | 0.932658612248051  |
| GM          | -2.05317553446496  | 0.0646211460904481 |
| Ventricles  | -1.07355243429012  | 0.306009446133064  |
| Putamen     | 1.03950537071394   | 0.320879581411232  |
| Pallidum    | -1.11117350802823  | 0.290191085096696  |
| Hippocampus | -0.175439878929938 | 0.86392179513451   |
| Caudate     | -0.39284293561879  | 0.701941462917964  |
| Amygdala    | -1.6724325678476   | 0.122609136897853  |
| Thalamus    | 1.22045802906405   | 0.24780879896461   |



**Figure S5.** WM mask extracted from FA, MSMT signal fraction and DORIS from a Penthera 3T subject.



**Figure S6.** Red circles show segmentation errors in the silver standard that are not present in other subjects. Given a large number of such examples in the training set, one can imagine that the Dense-Unet model will learn to avoid these errors.