

An Exegesis of Advanced Normalization Tools (*ANTs*)

Brian B. Avants, Nicholas J. Tustison, Hans J. Johnson & the ITK and registration community

Contents

.....	3
Big picture	3
.....	3
Powerful, general-purpose automated or semi-automated registration and segmentation.	4
Fine-grained and flexible maps	4
Anatomical quantification	5
Differentiable maps with differentiable inverse + <i>statistics in these spaces</i>	6
Brain images as <i>high-dimensional data</i>	6
Political activism	6
Customizable multivariate segmentation & registration	6
Mammalian cortical thickness computed with ANTs	6
General theory tunable to specific domains: <i>no-free lunch</i>	7
Agnostic data integration: prediction, decoding, diagnosis	8
$ITK+ANTs+R = ANTsR$	8
Agnostic statistics	8
How do we quantify <i>life span</i> brain health in individuals and in populations?	8
ANTS $R \rightarrow$ new insight via quantification	8
Solid Core Permits Broad Applications	9
ANTs was begotten in SyN	9
Community response	9
SyN formulation	9
SyN for optimization symmetry	10
Minimizing interpolations	10

Library for multivariate image registration, segmentation & statistics	11
Anatomical quantification	11
Modality integration	12
ANTs v Freesurfer	12
SyN Example	12
Theory + evaluation + reproducibility	13
Founding Developers	13
A long history of research	13
Rigorous transformation definition is key	13
Merit Badges	14
.	14
open source	14
competitions	14
competitions	15
papers	16
The stories behind ANTs development	16
Registration	16
Questions about evolution	17
N4	17
N4 Introduction	18
Nonparametric nonuniform intensity normalization (N3)	18
Code	19
Atropos	19
Atropos components	20
KellySlater -> KellyKapowski	20
Atropos + KK Example	20
Putting it all together	21
The ANTs cortical thickness pipeline	21
Basic components of the pipeline	21
Template building	21
Template building (cont.)	21
Brain extraction	22
Brain segmentation	22
Cortical thickness estimation	22
Data inspection w/ <i>ANTsR</i>	22

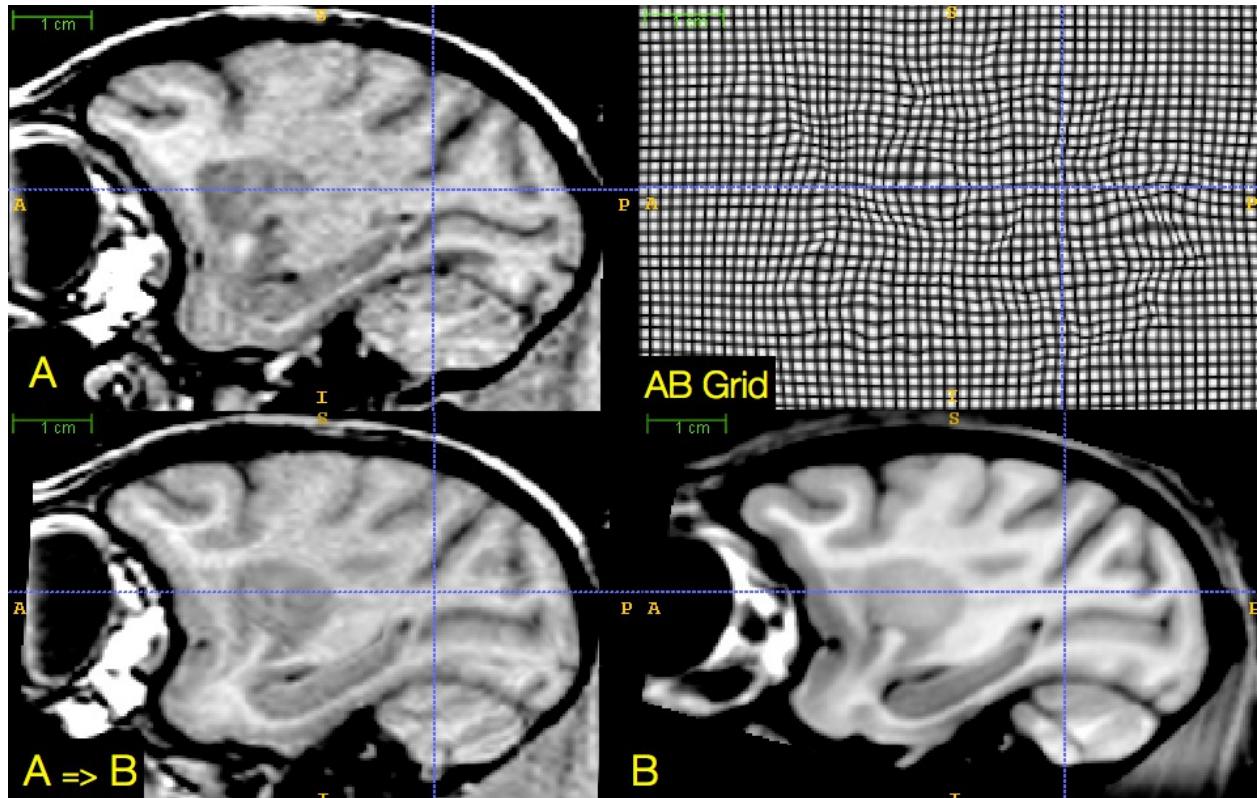
Evaluation	22
A couple notes on usage	22
Components	22
Software engineering	22
Analysis philosophy and published opinions	23
What is and <i>is not</i> image registration	23
Voodoo in voxel-based analysis	23
Instrumentation bias in the use and evaluation of software	23
Discussion	23
Problems	23
Strengths	23
Tools you can use for imaging science	24
References	24

Big picture

- Powerful, general-purpose automated or semi-automated registration and segmentation.
- Differentiable maps with differentiable inverse + *statistics in these spaces*
- Customizable multivariate segmentation & registration
- Borg philosophy: from I/O, to processing to statistical methods
- Open source, testing, many examples, consistent style, multiple platforms, active community support
...
- Integration with *R* + novel tools for prediction, decoding, high-to-low dimensional statistics.
- Collaborations with [neurodebian](#), [slicer](#), [brainsfit](#), [nipype](#), [itk](#) and more ...

Powerful, general-purpose automated or semi-automated registration and segmentation.

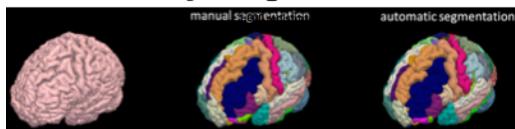
Fine-grained and flexible maps



Anatomical quantification

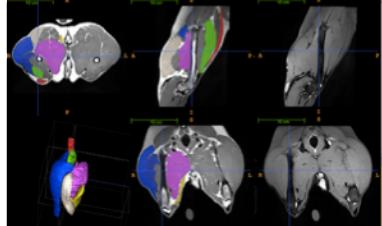
SATA Challenge MICCAI 2013: ANTs Provides Standardized Registration

1600 “easy” registrations



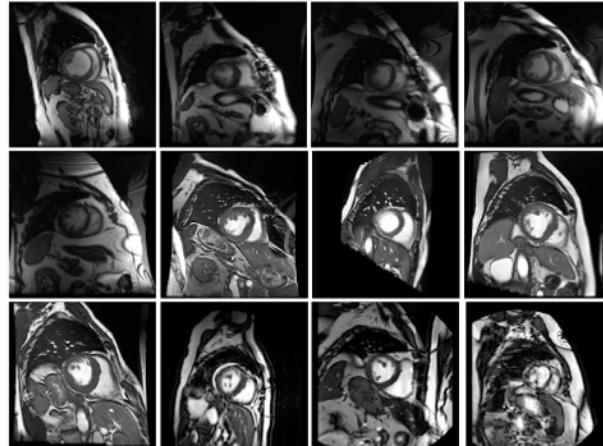
I wouldn't be surprised if getting "reasonable" & "consistent" registrations for this data is difficult or **impossible**.

7,000+ multivariate
registrations (challenging)



**Expected Results for
challenge participants**
~ Dice = 0.8+

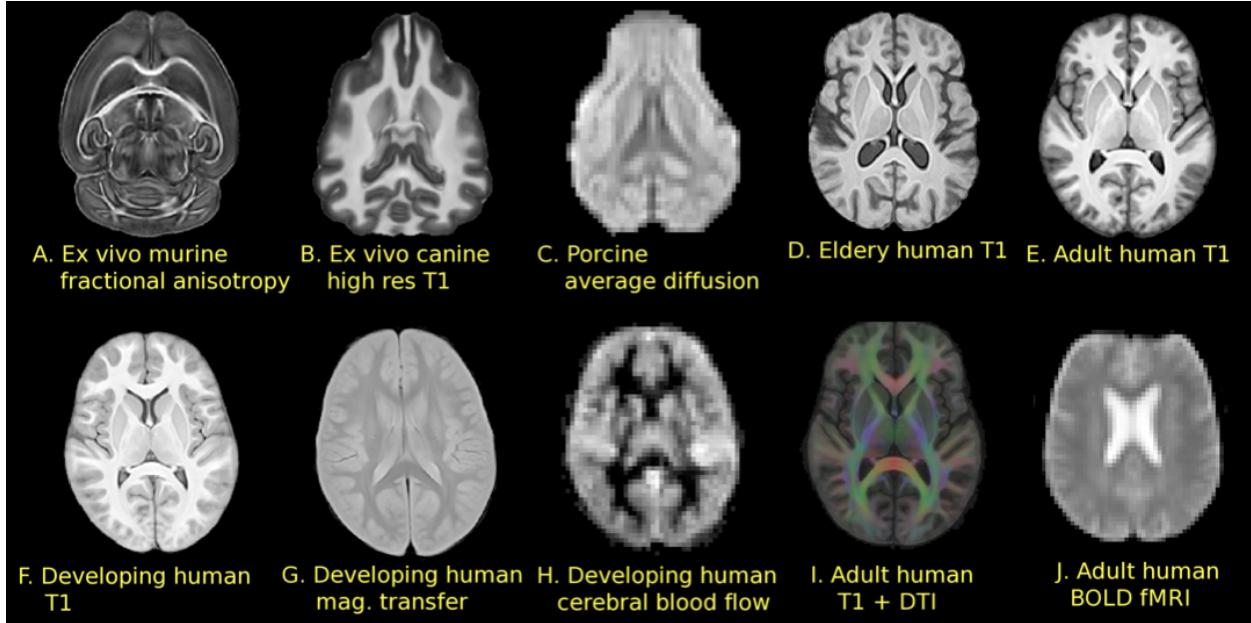
13,000+ 4D registrations
(very challenging)



Heart orientation varies between individuals

Differentiable maps with differentiable inverse + *statistics in these spaces*

Brain images as *high-dimensional data*

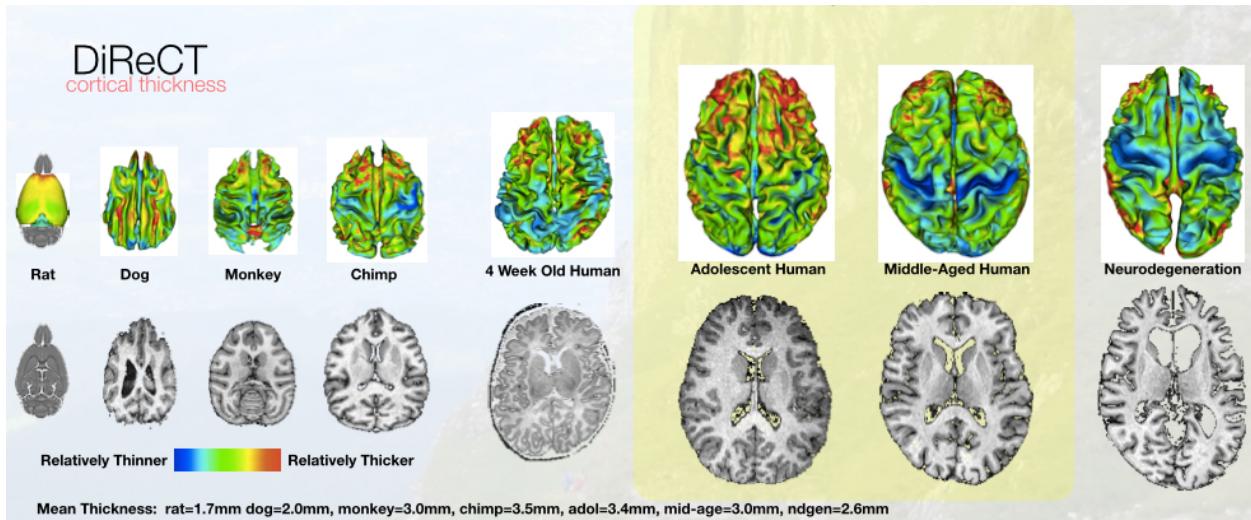


Political activism

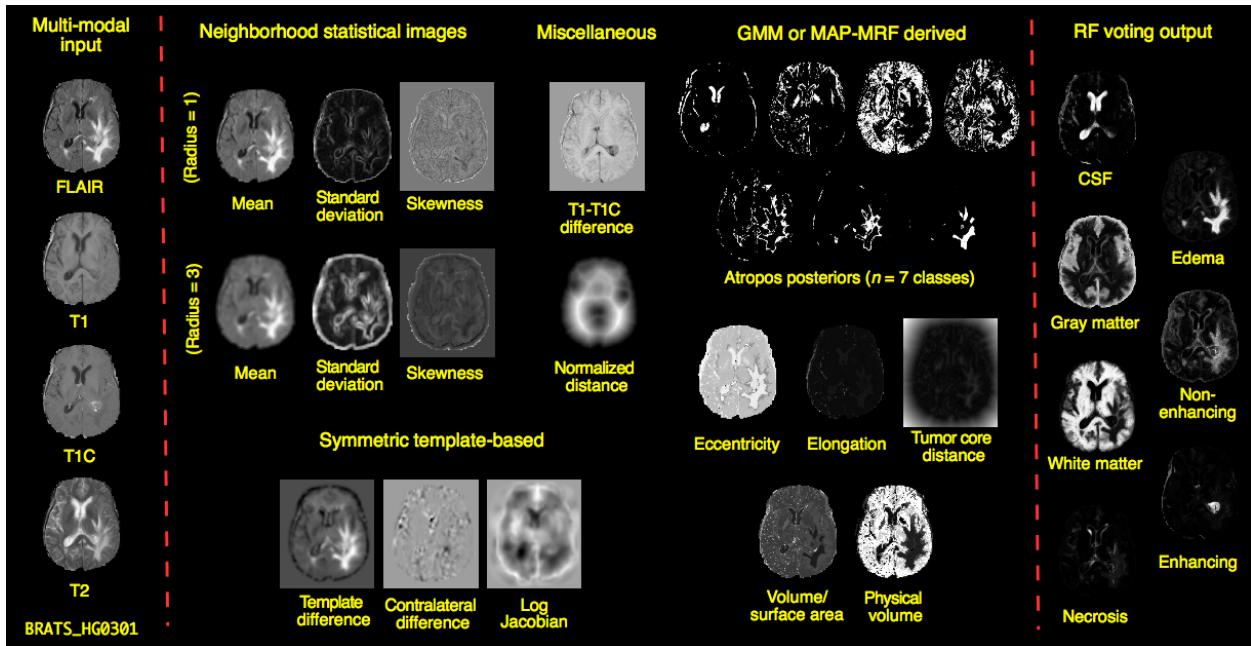
[congress commentary](#)

Customizable multivariate segmentation & registration

Mammalian cortical thickness computed with ANTs



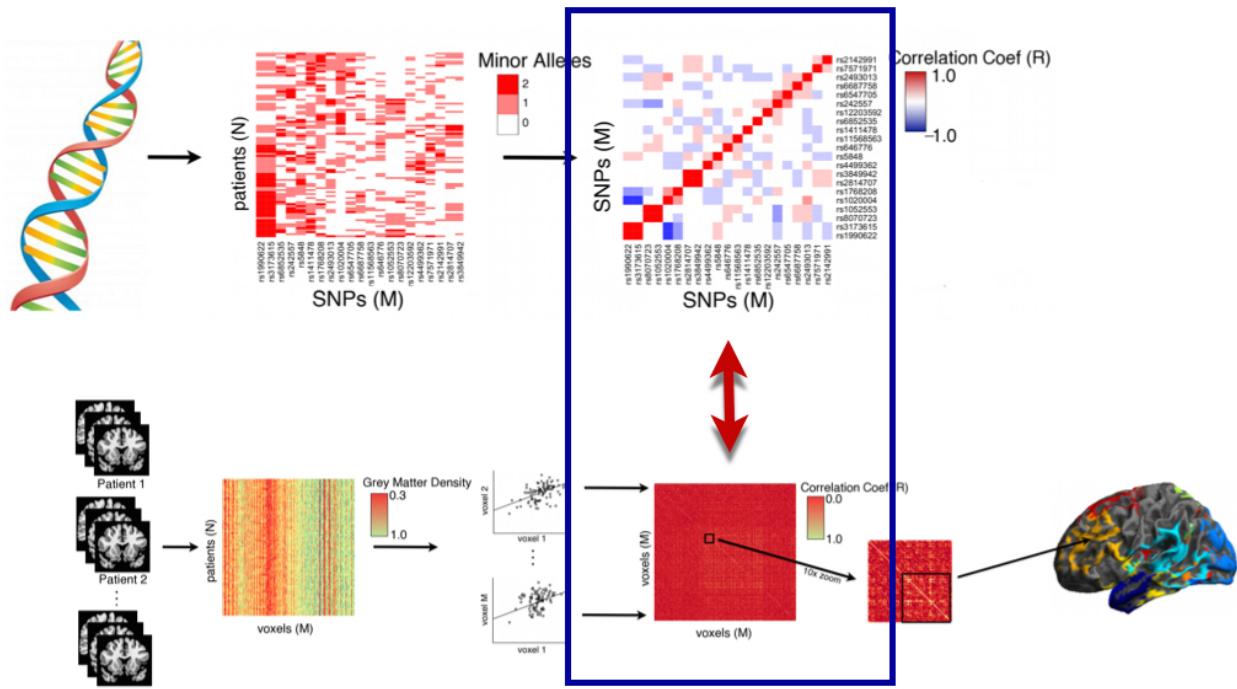
General theory tunable to specific domains: *no-free lunch*



Agnostic data integration: prediction, decoding, diagnosis

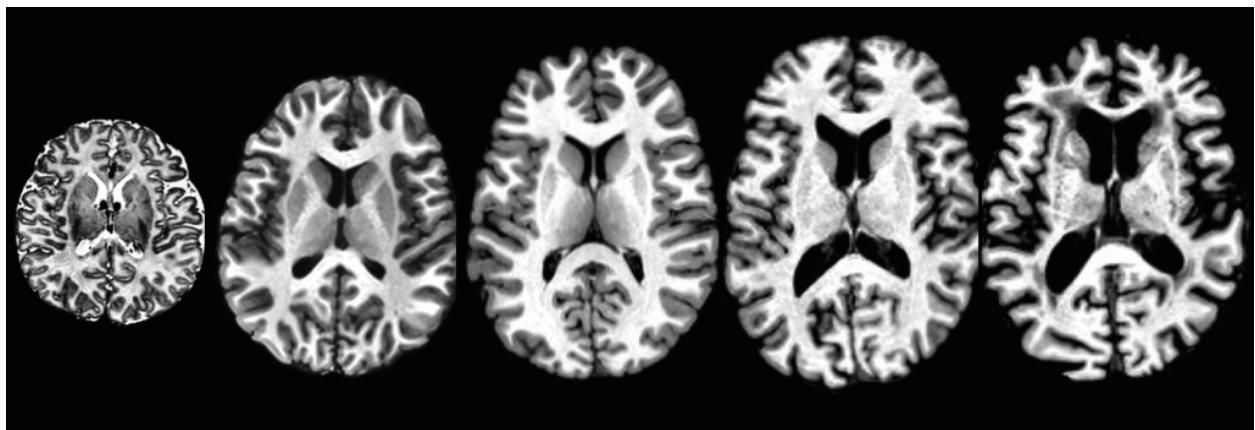
$$ITK + ANTs + R = ANTsR$$

Agnostic statistics



(McMillan et al, NoA, 2014)

How do we quantify *life span* brain health in individuals and in populations?



ANTsR → new insight via quantification

Good software should fade into the background ... however ...

As is common in science, the first big breakthrough in our understanding ... [came from] an improvement in measurement.

> Daniel Kahnemann, *Thinking, Fast and Slow* (2011)

Solid Core Permits Broad Applications

- microscopy
- satellite imagery
- interactive registration (see *Slicer*) and segmentation (see *ITK-SNAP*)

ANTs was begotten in SyN

Community response

- Just wanted to let you know—ANTS worked for me! I seem to be getting interesting results along the line_s of what I was predicting my study would find in specific parts of the striatum. —*Patryk from Ireland*
- To put not to fine a point on it—ANTS rocks. —*Craig from California*
- The ANTS package is very, very cool! —*Jaime from Wisconsin*
- Thank you for sharing this work, and congratulations for a job well done. —*Alexandra from North Carolina*

SyN formulation

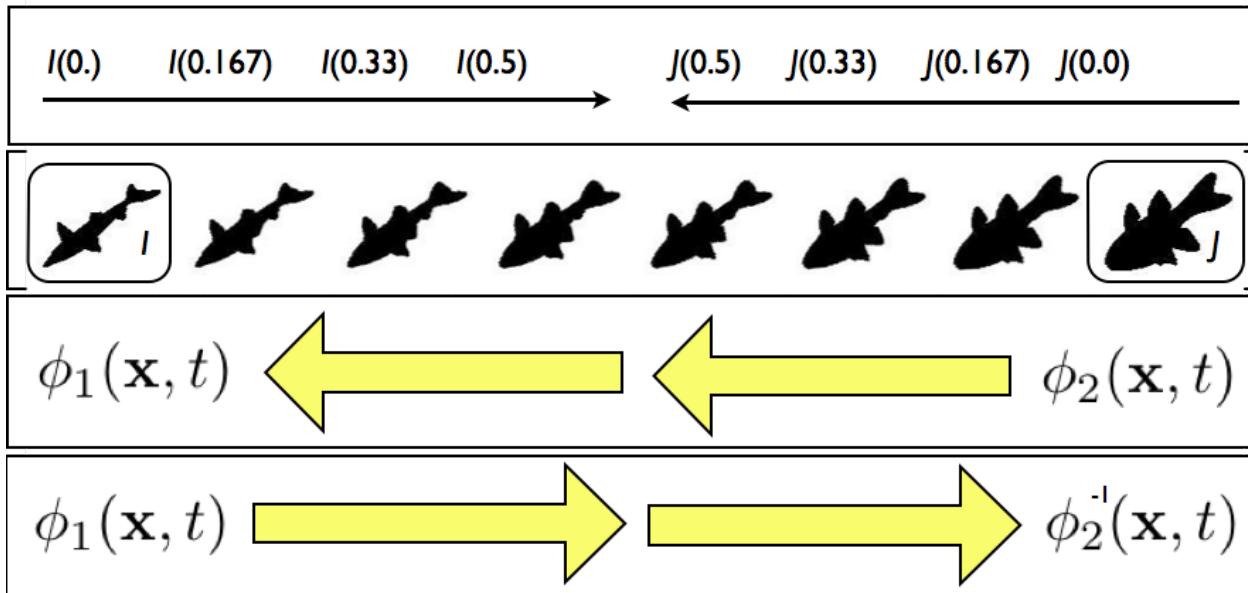
$$\inf_{\phi_1} \inf_{\phi_2} \left[\int_0^{0.5} (\|v_1(t)\|_L^2 + \|v_2(t)\|_L^2) dt + \int_{\Omega} \Pi_{\sim} (I \circ \phi_1^{-1}(\mathbf{x}, 0.5), J \circ \phi_2^{-1}(\mathbf{x}, 0.5)) d\Omega \right]$$

where

$$\frac{d\phi_i(\mathbf{x}, t)}{dt} = v_i(\phi_i(\mathbf{x}, t), t), \quad \phi_i(\mathbf{x}, 0) = \mathbf{Id}, \quad i \in \{1, 2\}$$

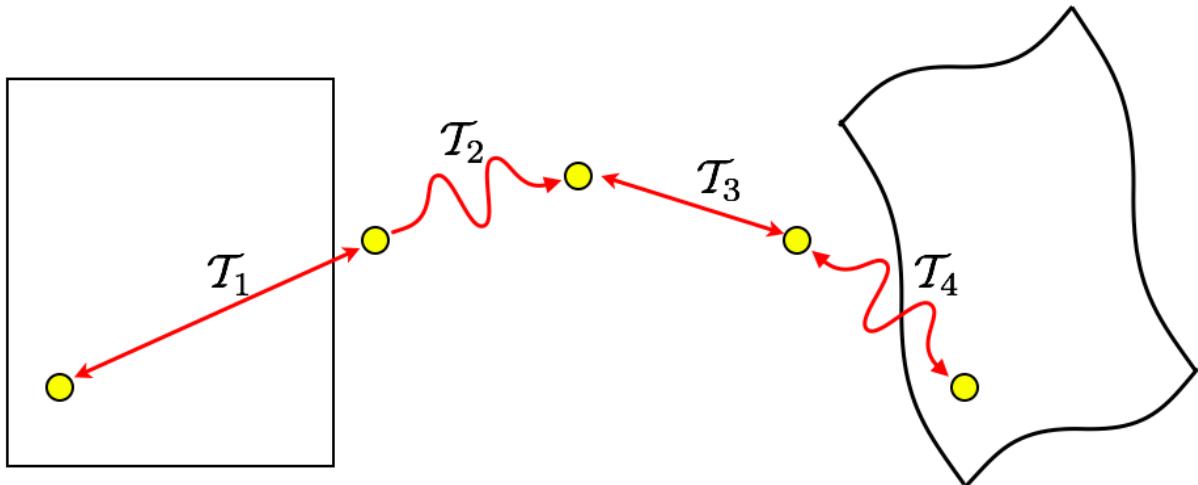
and Π_{\sim} is an arbitrary similarity metric (or metrics).

SyN for optimization symmetry



Images deform symmetrically along the shape manifold. This eliminates bias in the measurement of image differences.

Minimizing interpolations

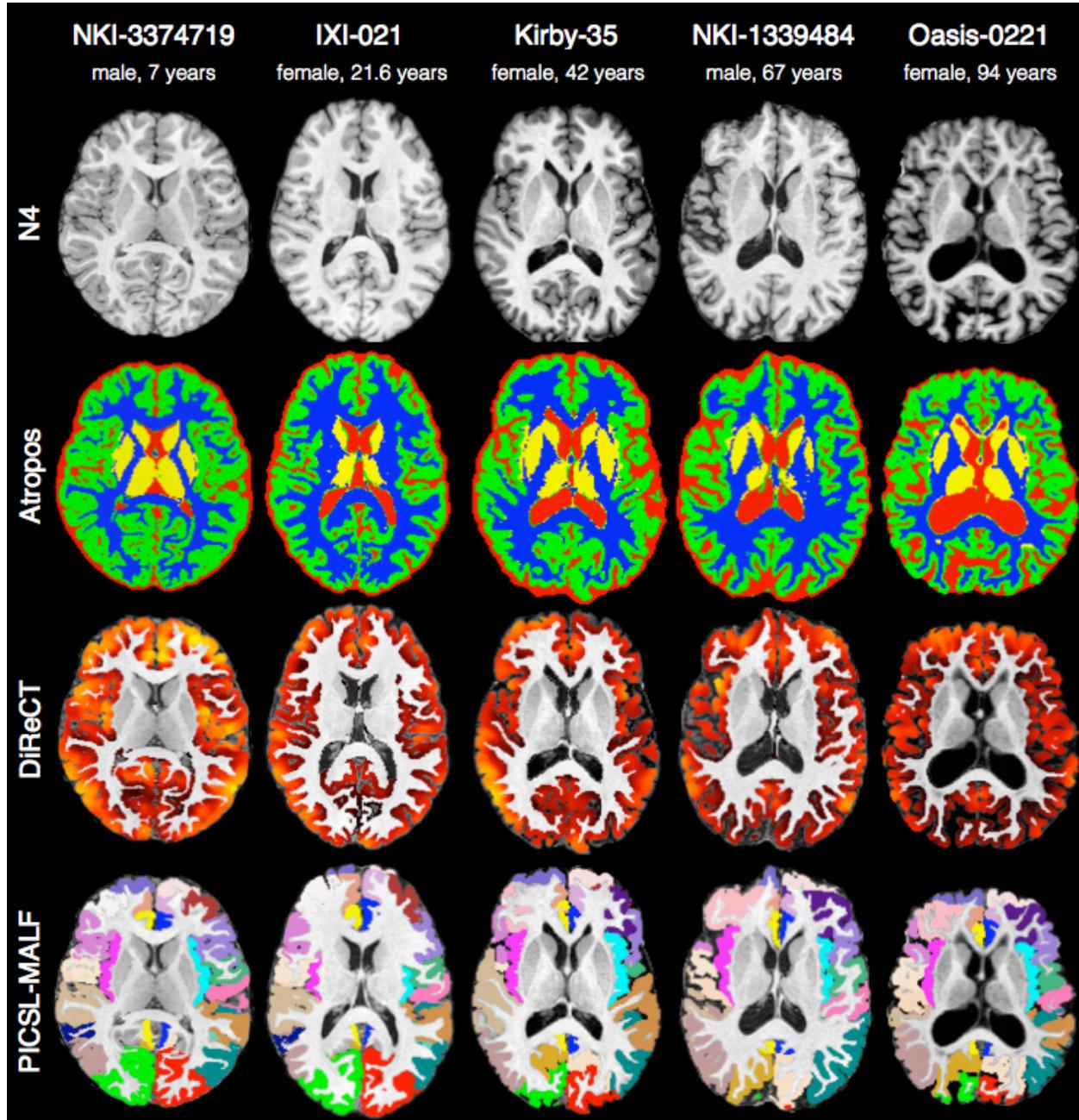


$$\mathcal{T}_{total} = \mathcal{T}_1 \circ \mathcal{T}_2 \circ \mathcal{T}_3 \circ \mathcal{T}_4$$

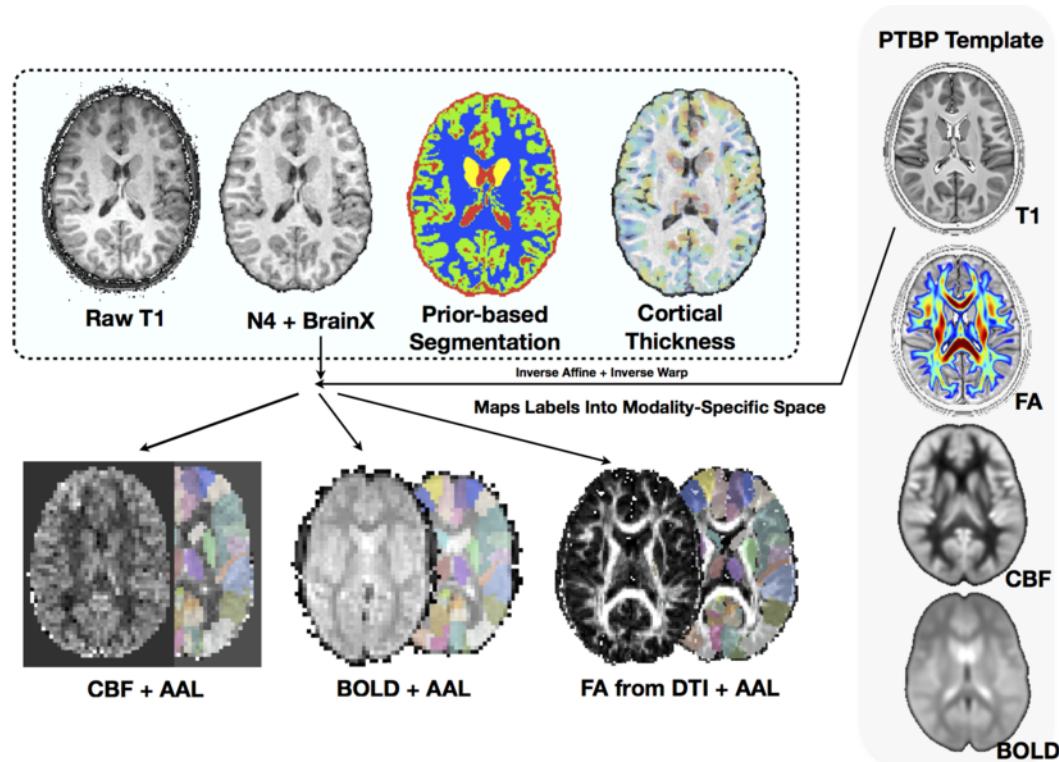
To avoid compounding interpolation error with the concatenation of transformations, ANTs never uses more than a single interpolation.

Library for multivariate image registration, segmentation & statistics

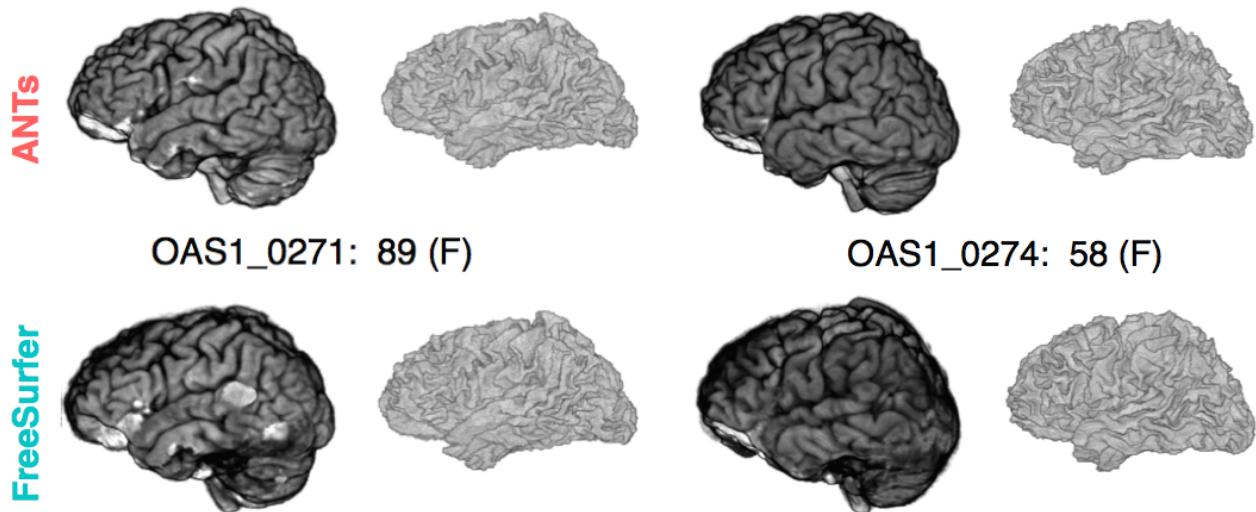
Anatomical quantification



Modality integration



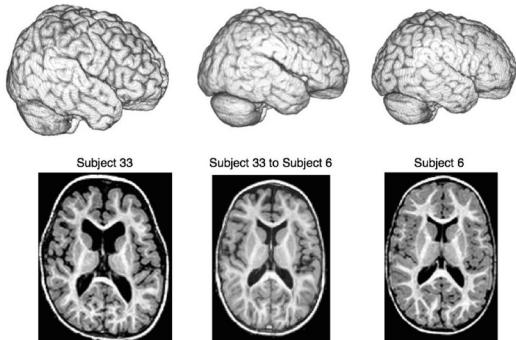
ANTs v Freesurfer



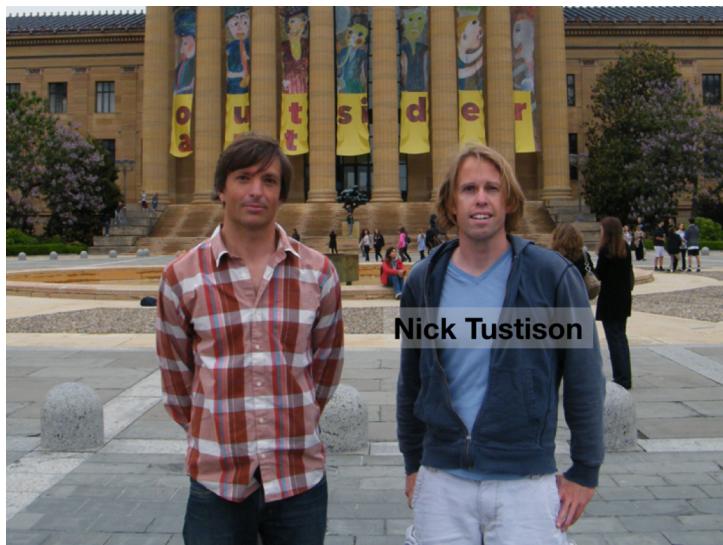
SyN Example

[SyN video](#)

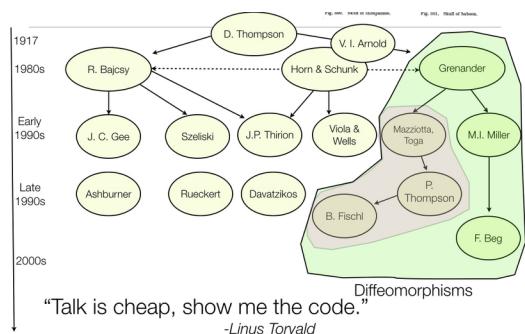
Theory + evaluation + reproducibility



Founding Developers



A long history of research



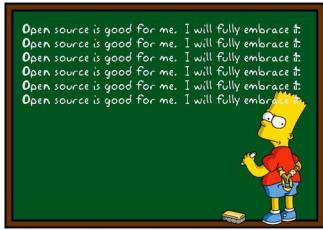
Rigorous transformation definition is key

ANTs and ITK are developed together: see B. B. Avants et al. (2014), Tustison and Avants (2013), Tustison et al. (2010) and more ...

Key definitions

- physical space
- transformation definition aware of physical space
- optimization space consistent with above
- unit testing

Merit Badges



open source

built on ITK—probably the most well-vetted medical image analysis package in the world B. B. Avants and Tustison (2014)

competitions

- Klein 2009: Brain Registration (ANTs)
- Murphy 2010: Lung Registration (ANTs)
- SATA 2012: Multi-Atlas Segmentation (ANTs+JointLabelFusion)
- SATA 2013: Multi-Atlas Segmentation (ANTs+JointLabelFusion)
- BRATS 2013: Multivariate Brain Segmentation (ANTsR)
- Yushkevich's Hipp Atlas: (hippocampusubfield.com)
- TBA: BOLD decoding (ANTsR)
- Substantial work with DTI (Camino developer in house)
- STACOM2014 ?

competitions



papers

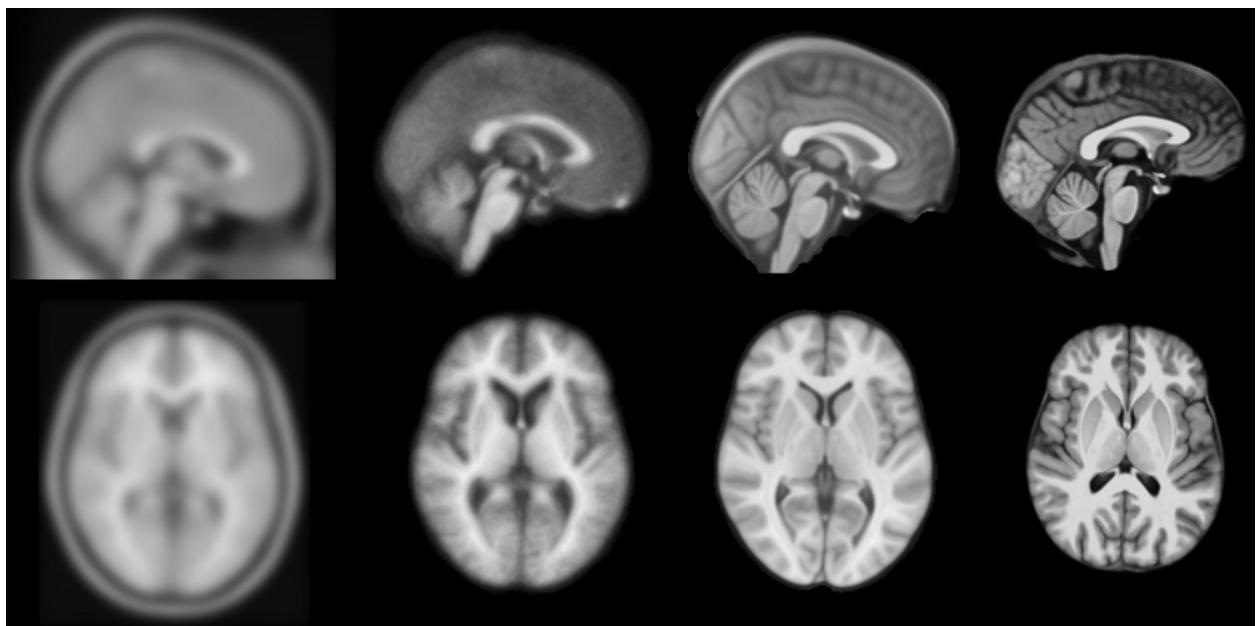
- registration : ANTs vs. everything else Klein et al. (2009)
- segmentation : Atropos vs. SPM, etc.
- bias correction : N4 vs N3
- cortical thickness : ANTs vs. FreeSurfer Tustison et al. (2014)
- compatibility with R

The stories behind ANTs development

(we can comment this out but I think it would be good to make some notes for posterity)

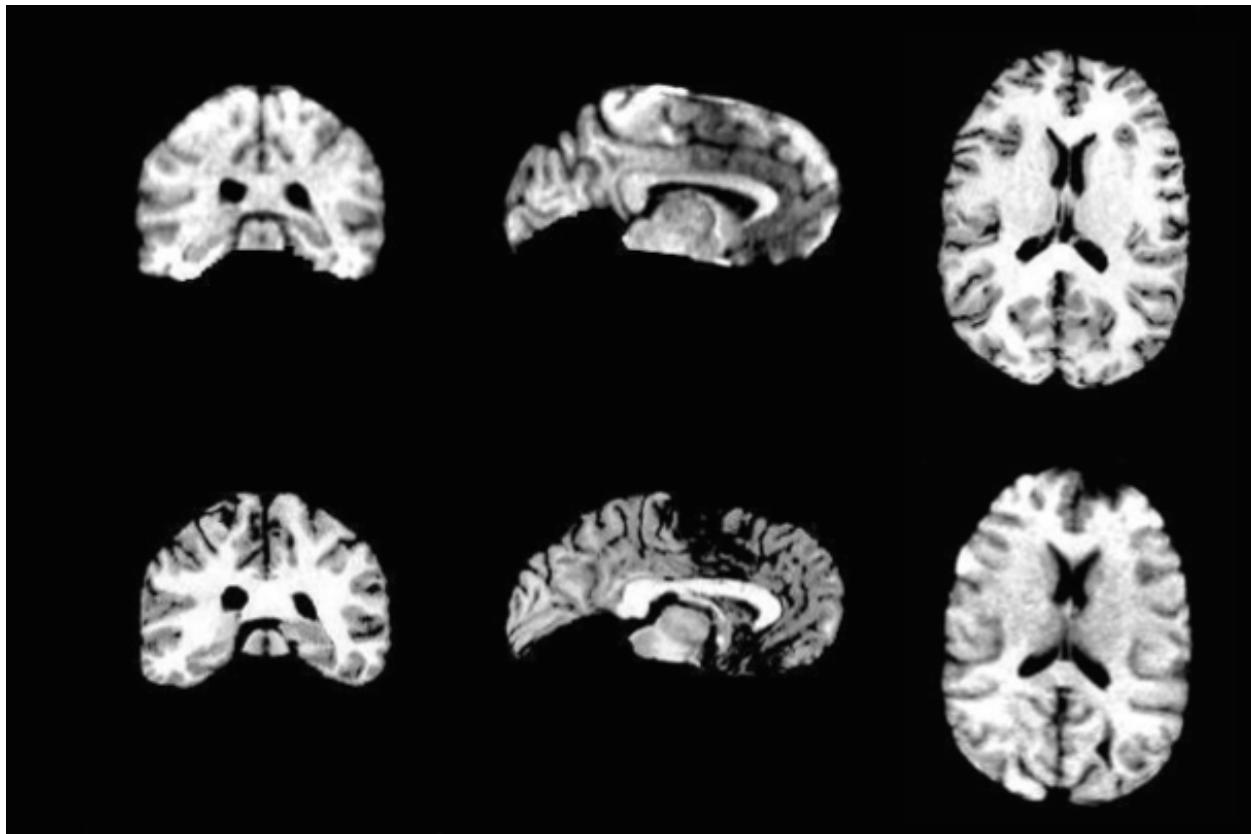
Registration

“I think these images could be better aligned.”



from [miykael](#)

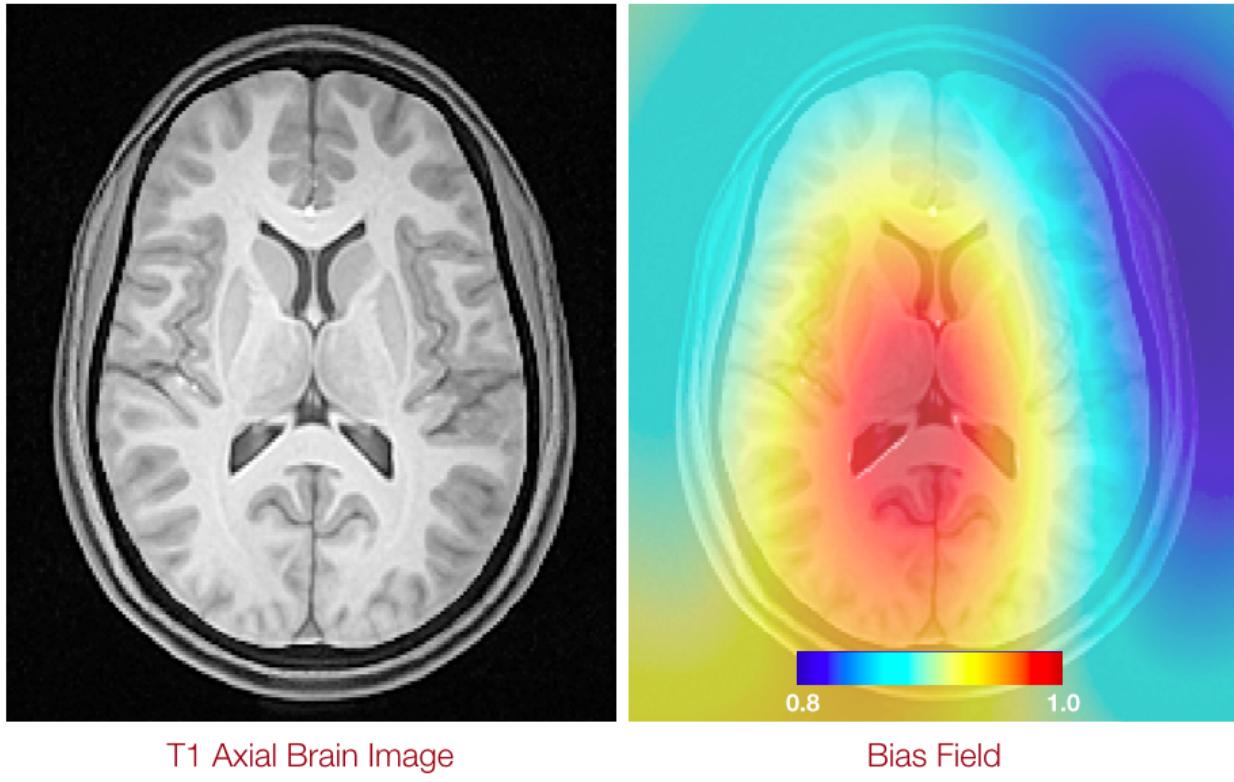
Questions about evolution



N4

- N3 (developed at the Montreal Neurological Institute) has been the gold standard for bias correction—used in important projects such as ADNI
- N3 is a set of perl scripts that works natively with the MINC file format which we tried to incorporate into an ANTs processing pipeline.
- We had so much trouble converting back and forth between ITK-compatible Nifti format and MINC that Brian suggested we try to implement N3 in ITK.
- I had some experience with B-splines and added some other tweaks giving birth to N4.

N4 Introduction



Nonparametric nonuniform intensity normalization (N3)

Sled et al., “A nonparametric method for automatic correction of intensity nonuniformity in MRI Data,” *IEEE-TMI*, 17(1), 1998.

Boyes et al., “Intensity non-uniformity correction using N3 on 3-T scanners with multichannel phased array coils,” *NeuroImage*, 39(4), 2008.

In a comparison of several correction techniques N3 performed well (Arnold et al., 2001). Also, the algorithm and software are in the public domain (<http://www.bic.mni.mcgill.ca/software/N3/>) and is probably the most widely used non-uniformity correction technique in neurological imaging.

Zheng et al., “Improvement of brain segmentation accuracy by optimizing non-uniformity correction using N3,” *NeuroImage*, 48(1), 2009.

Among existing approaches, the nonparametric non-uniformity intensity normalization method N3 (Sled et al., 1998) is one of the most frequently used... High performance and robustness have practically turned N3 into an industry standard.

Vovk et al., “A Review of Methods for Correction of Intensity Inhomogeneity in MRI,” *IEEE-TMI*, 26(3), 2007.

A well-known intensity inhomogeneity correction method, known as the N3 (nonparametric nonuniformity normalization), was proposed in [15]... Interestingly, no improvements have been suggested for this highly popular and successful method... The nonparametric nonuniformity normalization (N3) method [15] has obviously become the standard method against which other methods are compared.

Code

COMMAND:

```
N4BiasFieldCorrection
```

OPTIONS:

```
-d, --image-dimensionality 2/3/4
-i, --input-image inputImageFilename
-x, --mask-image maskImageFilename
-w, --weight-image weightImageFilename
-s, --shrink-factor 1/2/3/4/...
-c, --convergence [<numberOfIterations=50x50x50x50>, <convergenceThreshold=0.0>]
-b, --bspline-fitting [splineDistance, <splineOrder=3>]
    [initialMeshResolution, <splineOrder=3>]
-t, --histogram-sharpening [<FWHM=0.15>, <wienerNoise=0.01>, <numberOfHistogramBins=200>]
-o, --output correctedImage
    [correctedImage, <biasField>]
-h
--help
```

Talk is cheap, show me the code.

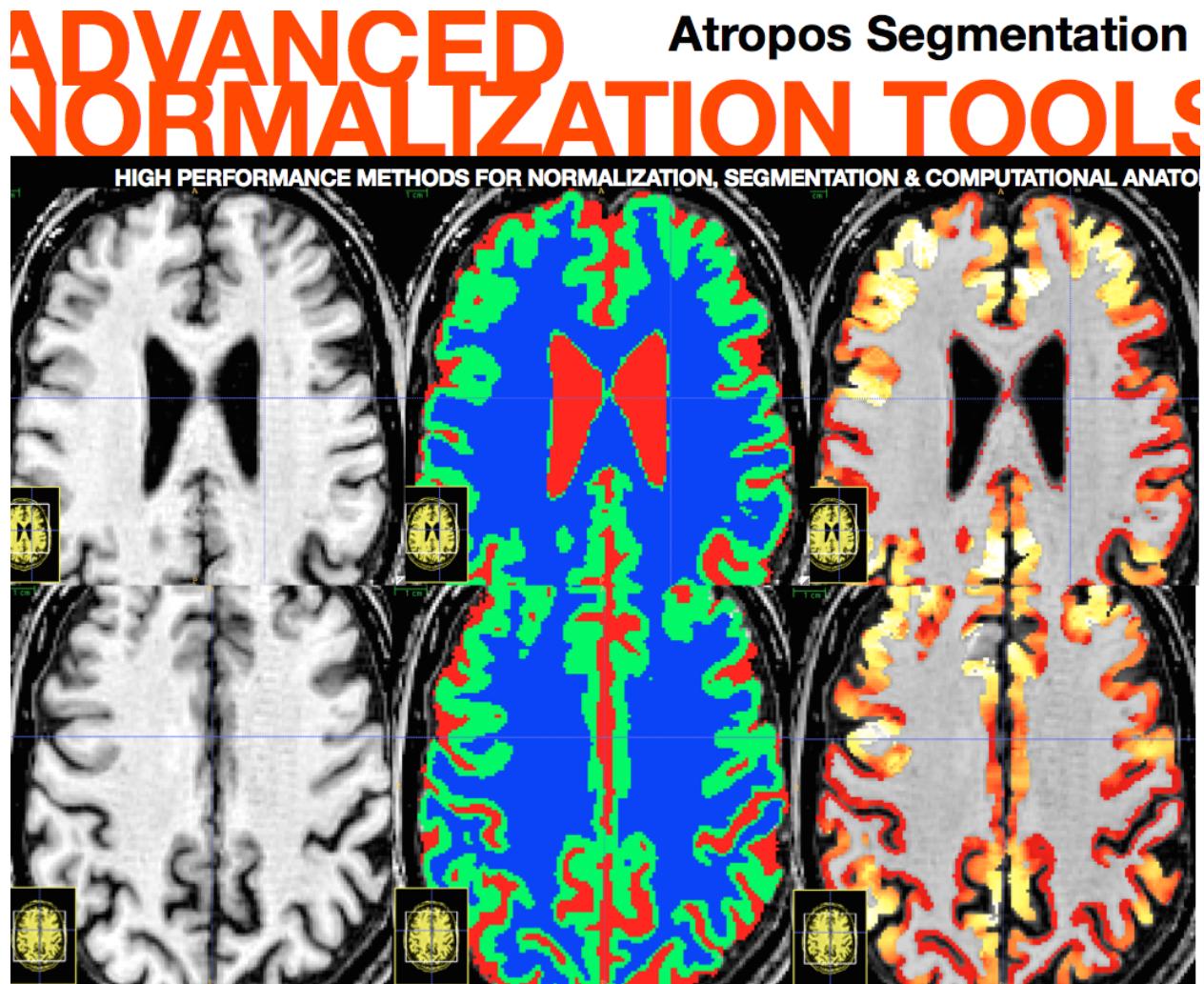
Atropos

- Similar to our experience with N3, we tried to incorporate FAST (from the FMRIB at Oxford) into an ANTs processing pipeline.
- Phil (?) pointed out the difficulty of incorporating priors into FAST.
- Related, Brian went to a segmentation-related worksop at MICCAI and aired publicly his disappointment that so much of what had been developed in the community over the last 20+ years has not been made publicly available. “What’s wrong with you people!”
- 3-tissue algorithm in ImageMath —> multivariate, n-class Atropos

Atropos components

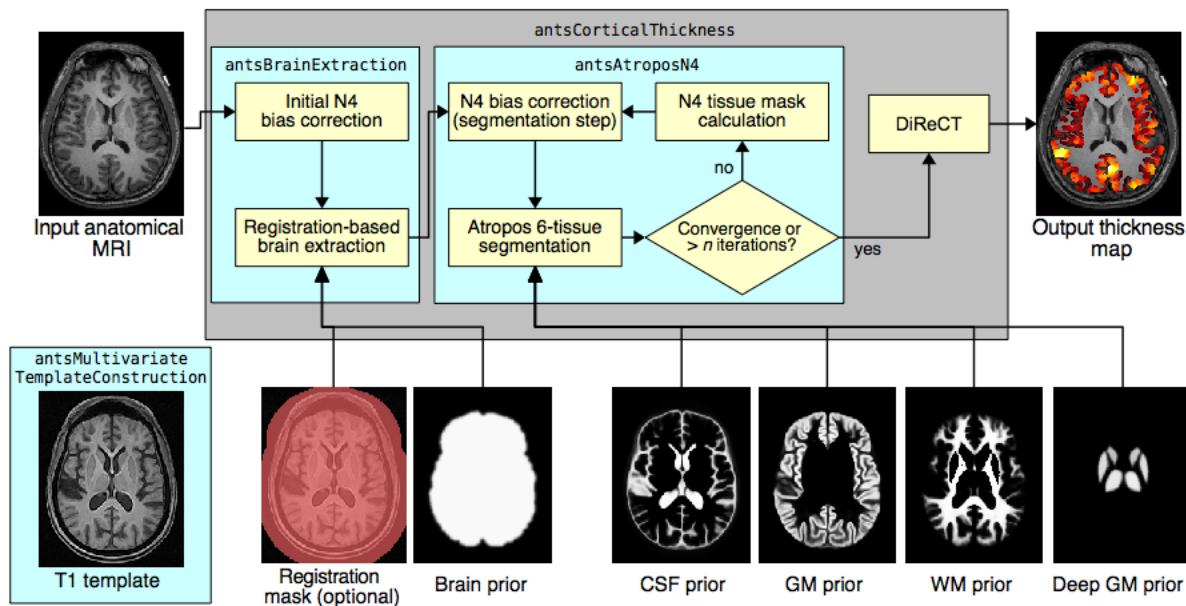
KellySlater → KellyKapowski

Atropos + KK Example



Putting it all together

The ANTs cortical thickness pipeline



*Large-scale evaluation of ANTs and FreeSurfer cortical thickness measurements, NeuroImage 2014.**

All software components are open source and part of the Advanced Normalization Tools (ANTs) repository.

Basic components of the pipeline

1. template building (offline)
2. brain extraction
3. cortical thickness estimation
4. cortical parcellation

Template building

Tailor data to your specific cohort

- Templates representing the average mean shape and intensity are built directly from the cohort to be analyzed, e.g. pediatric vs. middle-aged brains.
- Acquisition and anonymization (e.g. defacing) protocols are often different.

Template building (cont.)

Each template is **processed** to produce auxiliary images which are used for brain extraction and brain segmentation.

Brain extraction

Comparison with de facto standard FreeSurfer package. Note the difference in separation of the gray matter from the surrounding CSF. (0 failures out of 1205 scans)

Brain segmentation

Randomly selected healthy individuals. Atropos gets good performance across ages.

Cortical thickness estimation

In contrast to FreeSurfer which warps coupled surface meshes to segment the gray matter, ANTs diffeomorphically registers the white matter to the combined gray/white matters while simultaneously estimating thickness.

Data inspection w/*ANTsR*

spider plots

Evaluation

Question: In the absence of ground truth, how do we evaluate performance?

Answer: We use demographic information with known thickness sensitivity and correlate with measured thickness quantities. One of the most well-known, most easily obtained, and most confident measures available is “age.” So we take 50% of the thickness data to train a model (e.g. linear regression) and then calculate the model’s age prediction error on the other 50%. We do this for n=1000 permutations to build a distribution. Similarly, we can do this for gender.

A couple notes on usage

- Out of the many cortical thickness algorithms that have been proposed, FreeSurfer dominates. And rightfully so, because it works well and has been the only publicly available tool (until recently).
- In the same spirit, we have made our tools publicly available. Usage is similar to that of FreeSurfer (see below). We also make several templates available.

“Talk is cheap, show me the code.” —Linus Torvald

Components

Software engineering

[ants gource](#)

Analysis philosophy and published opinions

What is and *is not* image registration

not registration

Voodoo in voxel-based analysis

logical circularity VBA

Instrumentation bias in the use and evaluation of software

Instrumentation bias in the use and evaluation of software

Discussion

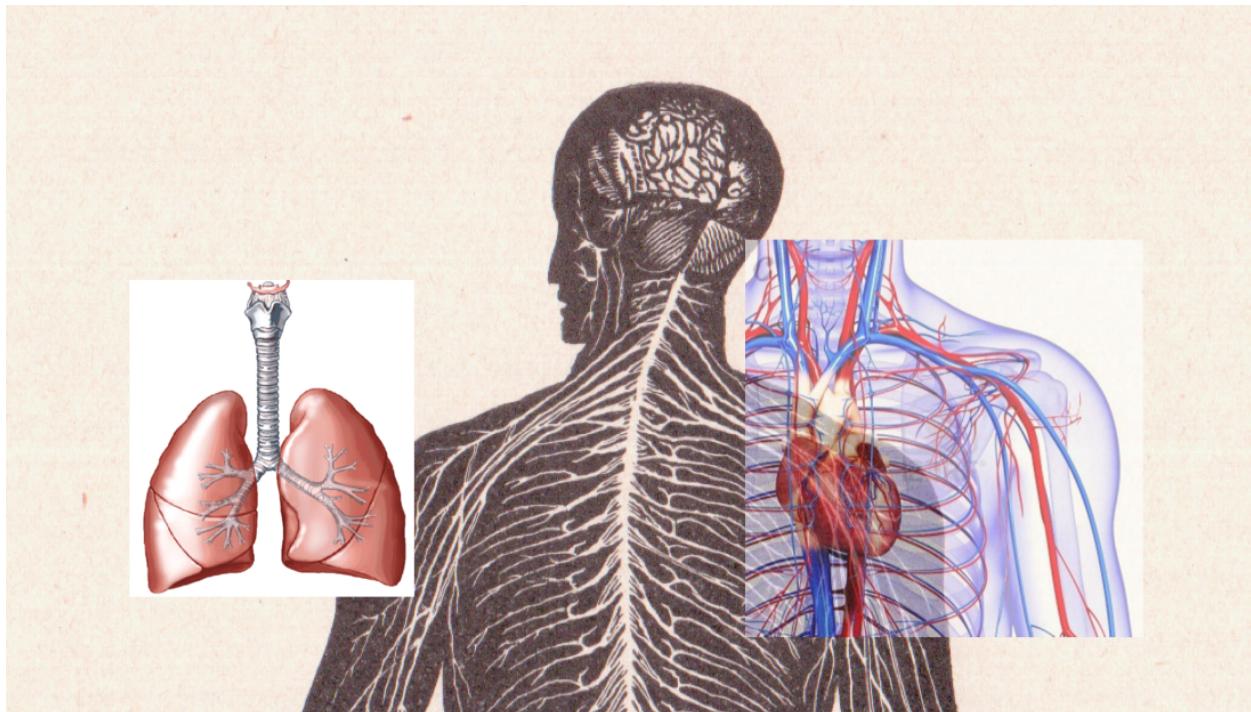
Problems

There are several problems and shortcomings to this analysis.

- Customizable for specific problems but not too specific
- Latest theoretical advances in registration not yet wrapped for users

Strengths

Some strengths include relatively few assumptions, a flexible implementation and open-science approach.



Tools you can use for imaging science

- Core developers: *B. Avants, N. Tustison, H. J. Johnson, J. T. Duda*
- Many contributors, including users ...
- Multi-platform, multi-threaded C++ stnava.github.io/ANTs
- Developed in conjunction with <http://www.itk.org/>
- R wrapping and extension stnava.github.io/ANTsR
- rapid development, regular testing + many eyes → bugs are shallow

References

- Avants, Brian B., and Nicholas J. Tustison. 2014. “The ITK Image Registration Framework.” *Front Neuroinform* 7. Penn Image Computing; Science Laboratory, Department of Radiology, University of Pennsylvania Philadelphia, PA, USA.: 39. doi:[10.3389/fninf.2013.00039](https://doi.org/10.3389/fninf.2013.00039).
- Avants, Brian B., Nicholas J. Tustison, Michael Stauffer, Gang Song, Baohua Wu, and James C. Gee. 2014. “The Insight ToolKit Image Registration Framework.” *Front Neuroinform* 8. Penn Image Computing; Science Laboratory, Department of Radiology, University of Pennsylvania Philadelphia, PA, USA.: 44. doi:[10.3389/fninf.2014.00044](https://doi.org/10.3389/fninf.2014.00044).
- Klein, Arno, Jesper Andersson, Babak A Ardekani, John Ashburner, Brian Avants, Ming-Chang Chiang, Gary E Christensen, et al. 2009. “Evaluation of 14 Nonlinear Deformation Algorithms Applied to Human Brain MRI Registration.” *Neuroimage* 46 (3). New York State Psychiatric Institute, Columbia University, NY, NY 10032, USA. arno@binarybottle.com: 786–802. doi:[10.1016/j.neuroimage.2008.12.037](https://doi.org/10.1016/j.neuroimage.2008.12.037).
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- Tustison, Nicholas J., Brian B. Avants, Philip A. Cook, Yuanjie Zheng, Alexander Egan, Paul A. Yushkevich, and James C. Gee. 2010. “N4ITK: Improved N3 Bias Correction.” *IEEE Trans Med Imaging* 29 (6). Department of Radiology, University of Pennsylvania, Philadelphia, PA 19140, USA. ntustison@wustl.edu: 1310–20. doi:[10.1109/TMI.2010.2046908](https://doi.org/10.1109/TMI.2010.2046908).
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