



NA-MIC

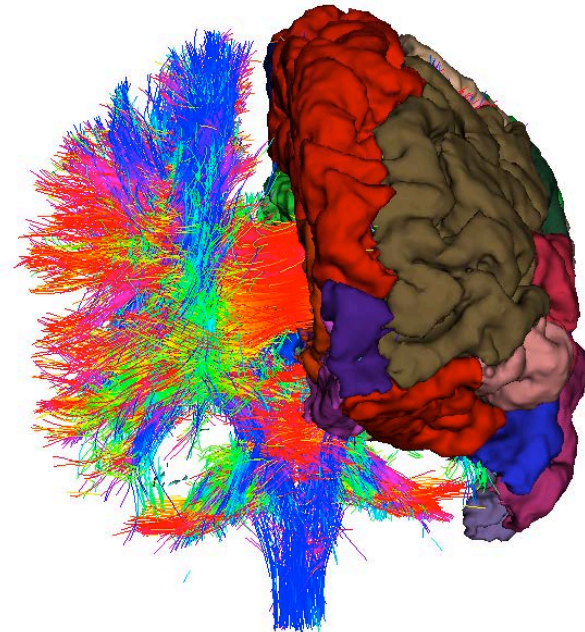
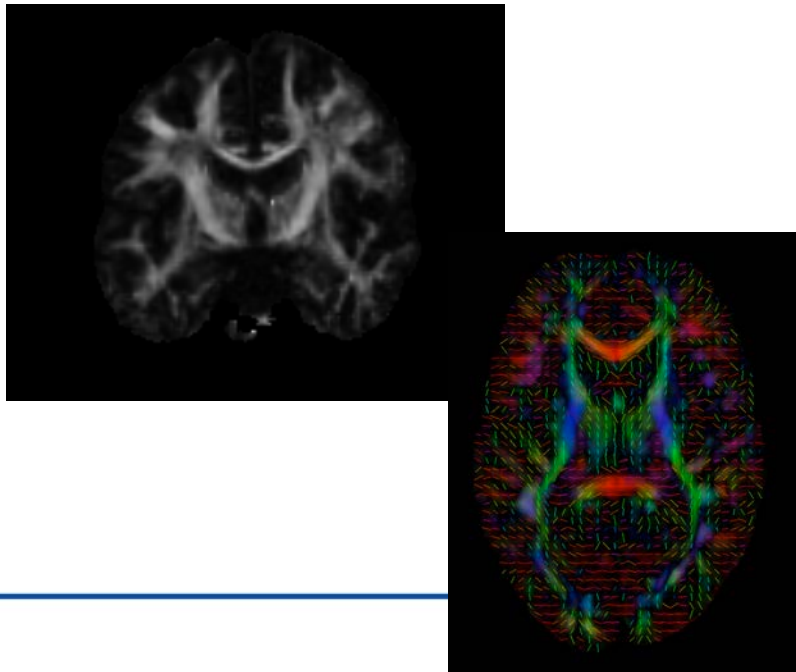
National Alliance for Medical Image Computing

<http://www.na-mic.org>

Driving Biological Problem Huntington's Disease



THE UNIVERSITY
OF IOWA



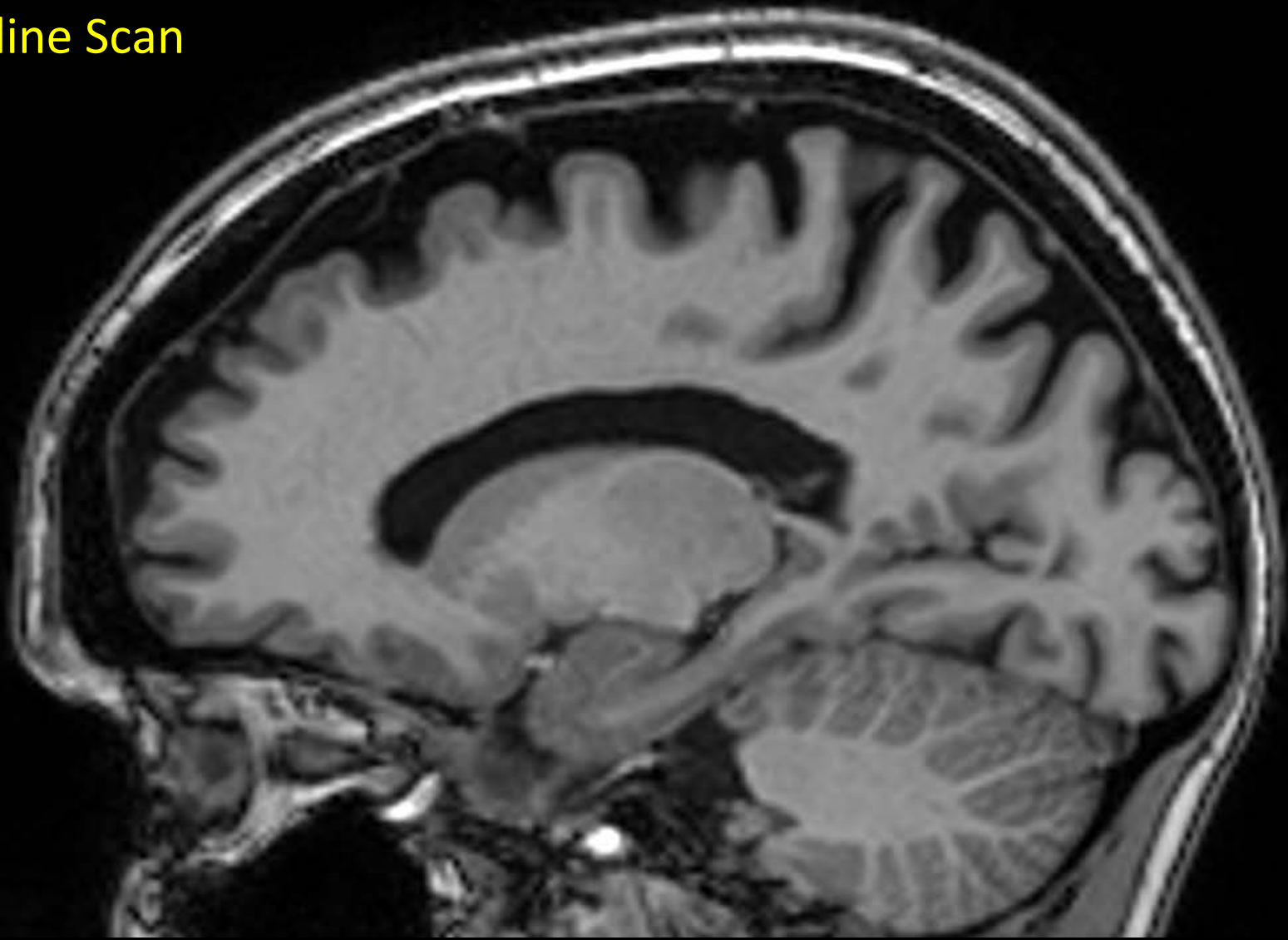


Builds on PREDICT-HD

- The NIH-funded project “Neurobiological Predictors of Huntington’s Disease” (PREDICT-HD) studies Huntington’s disease (HD), a neurodegenerative genetic disorder that affects muscle coordination, behavior, and cognitive function, and causes severe debilitating symptoms by middle age.
-

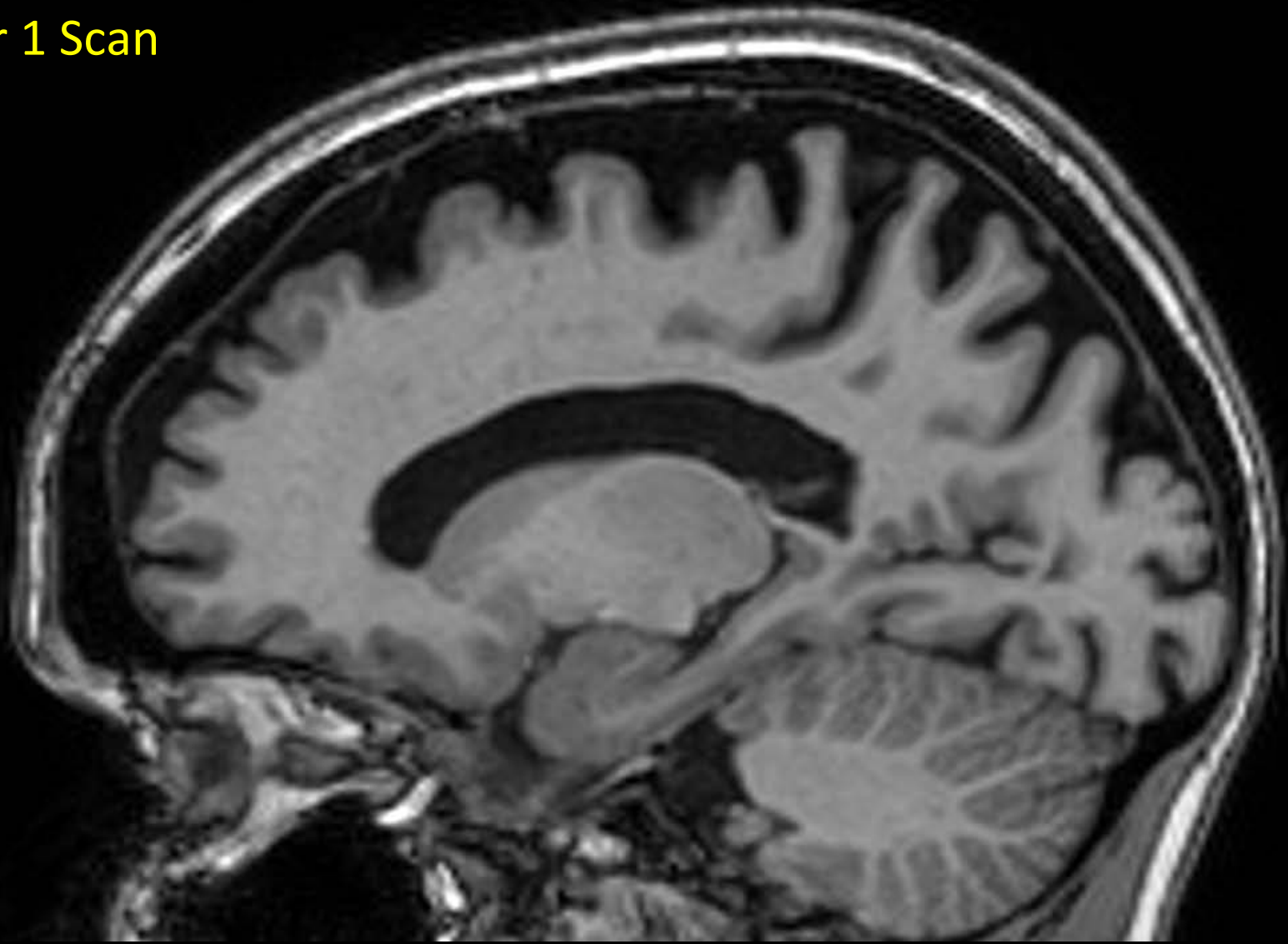
TRACK-HD Stage 1 HD Subject

Baseline Scan



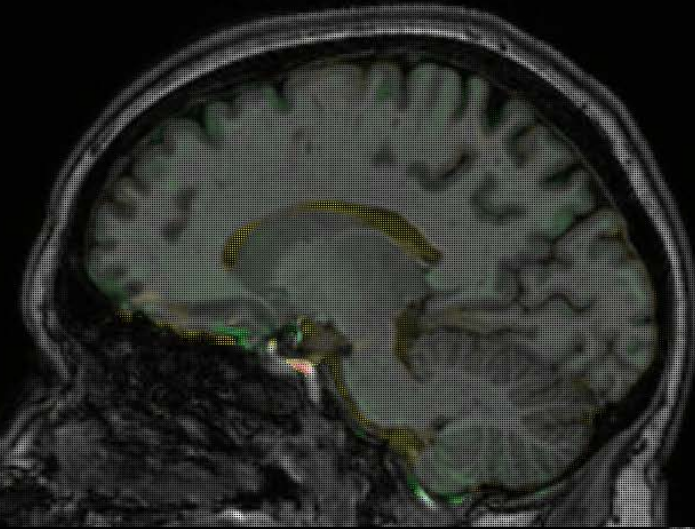
TRACK-HD Stage 1 HD Subject

Year 1 Scan

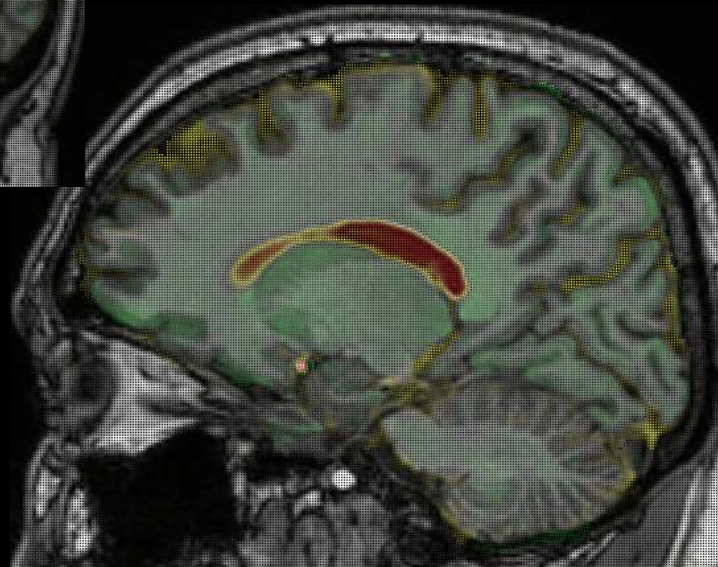


24-month voxel-compression mapping

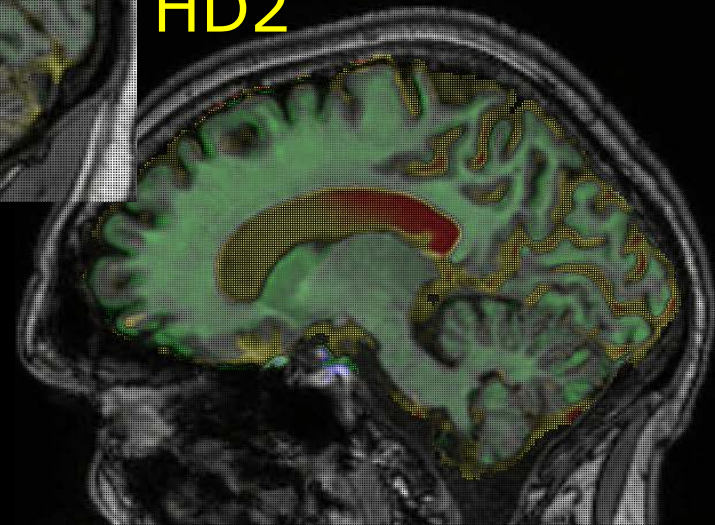
Control



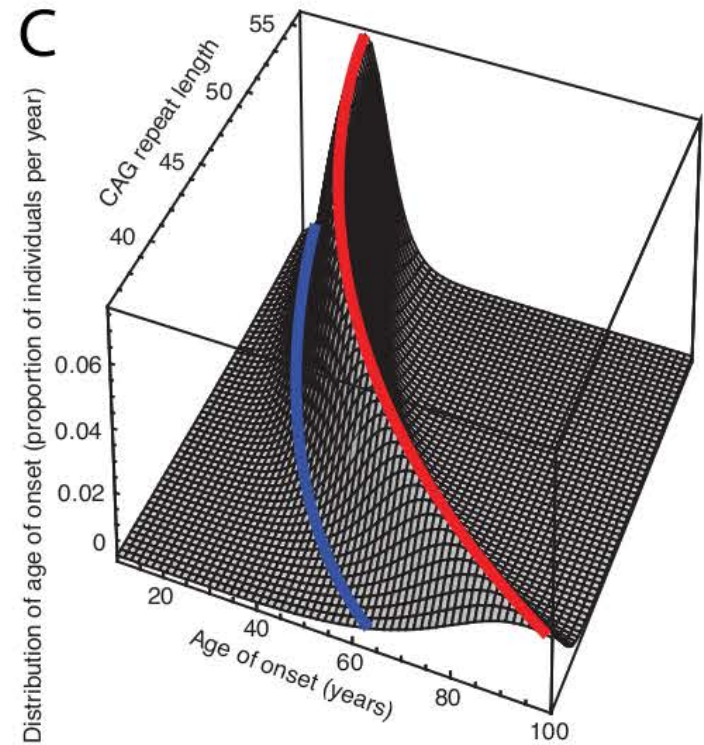
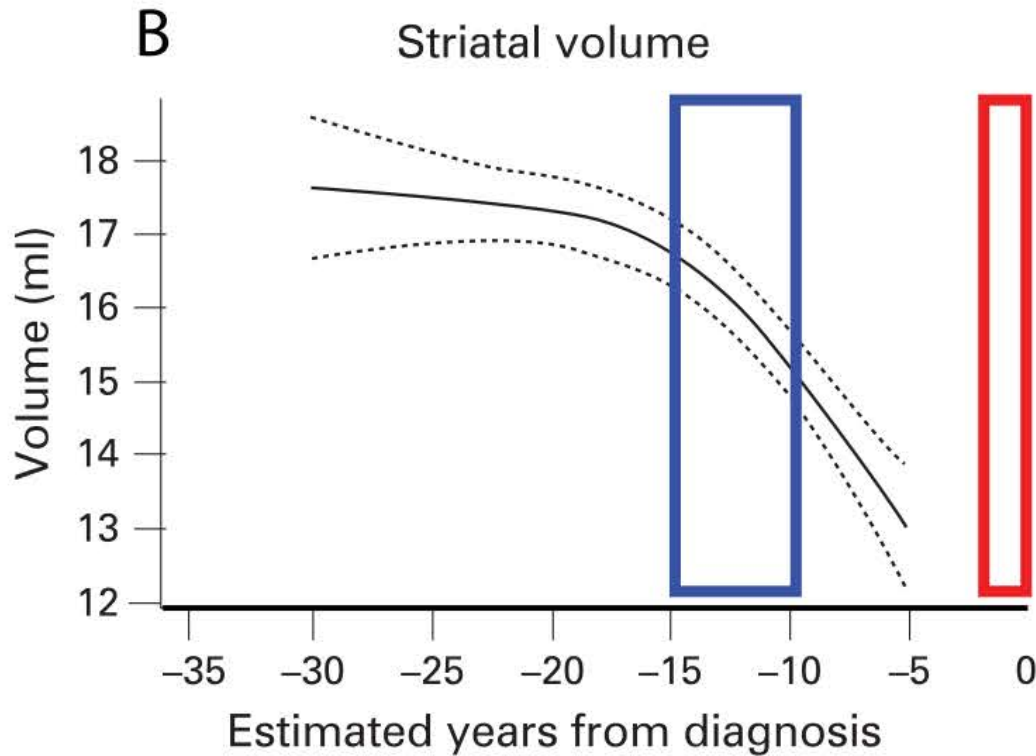
PreA



HD2



HD Background





Specific Aims

- Perform individualized longitudinal shape change quantification from multi-modal data.
- Complete full brain Diffusion Tensor Imaging tractography analysis.
- Deploy extensible tools for sharing source data, derived data, algorithms and methods to multi-site analysis teams.



Sharing HD Data

http://www.na-mic.org/Wiki/index.php/2011_Summer_Project_Week

User:

Home **New** **Upload** **Administer** **Tools**

Launch Uploader

Search
PREDICTHD currently contains **48 Projects, 1920 Subjects, and 4556 Imaging Sessions.**

Projects **Subjects** **MR** **PET** **CT**

Projects
Recent

TrackOnHD

Home **New** **Upload** **Administer**

Launch Uploader

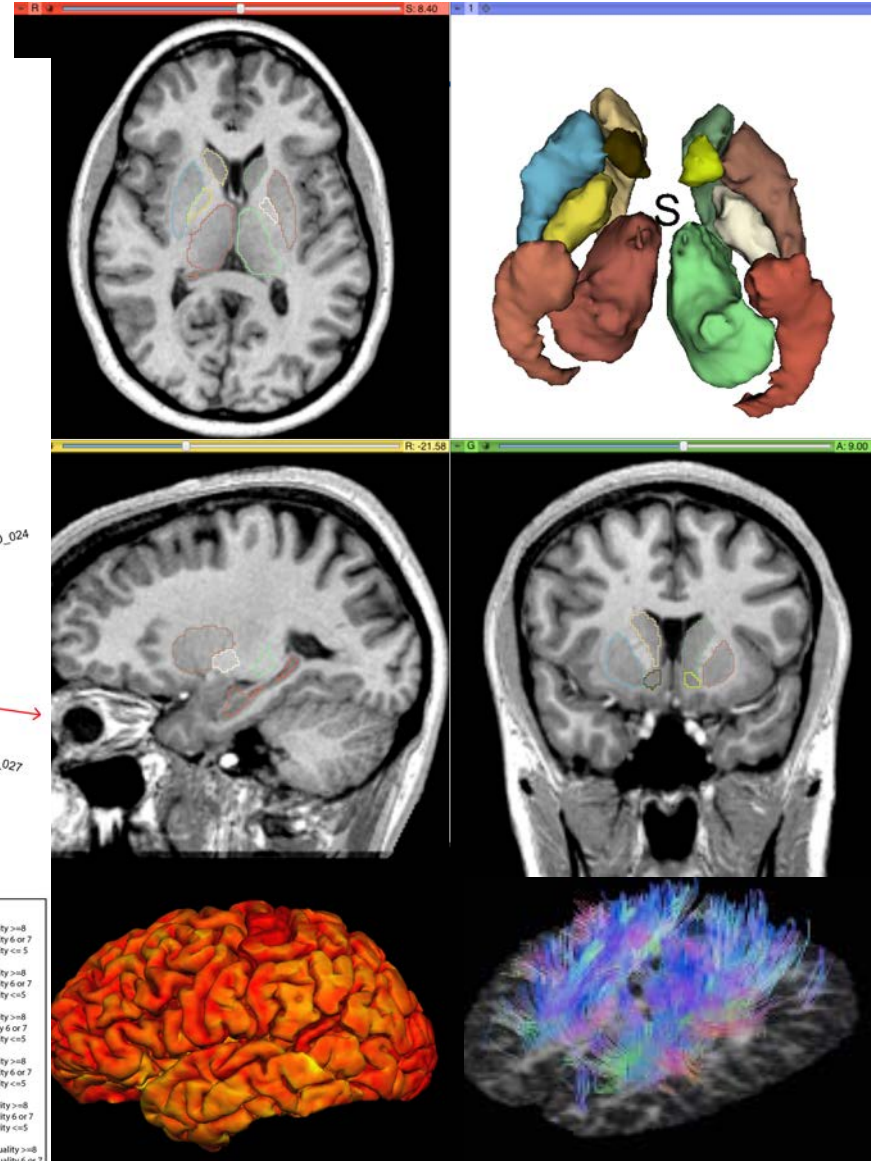
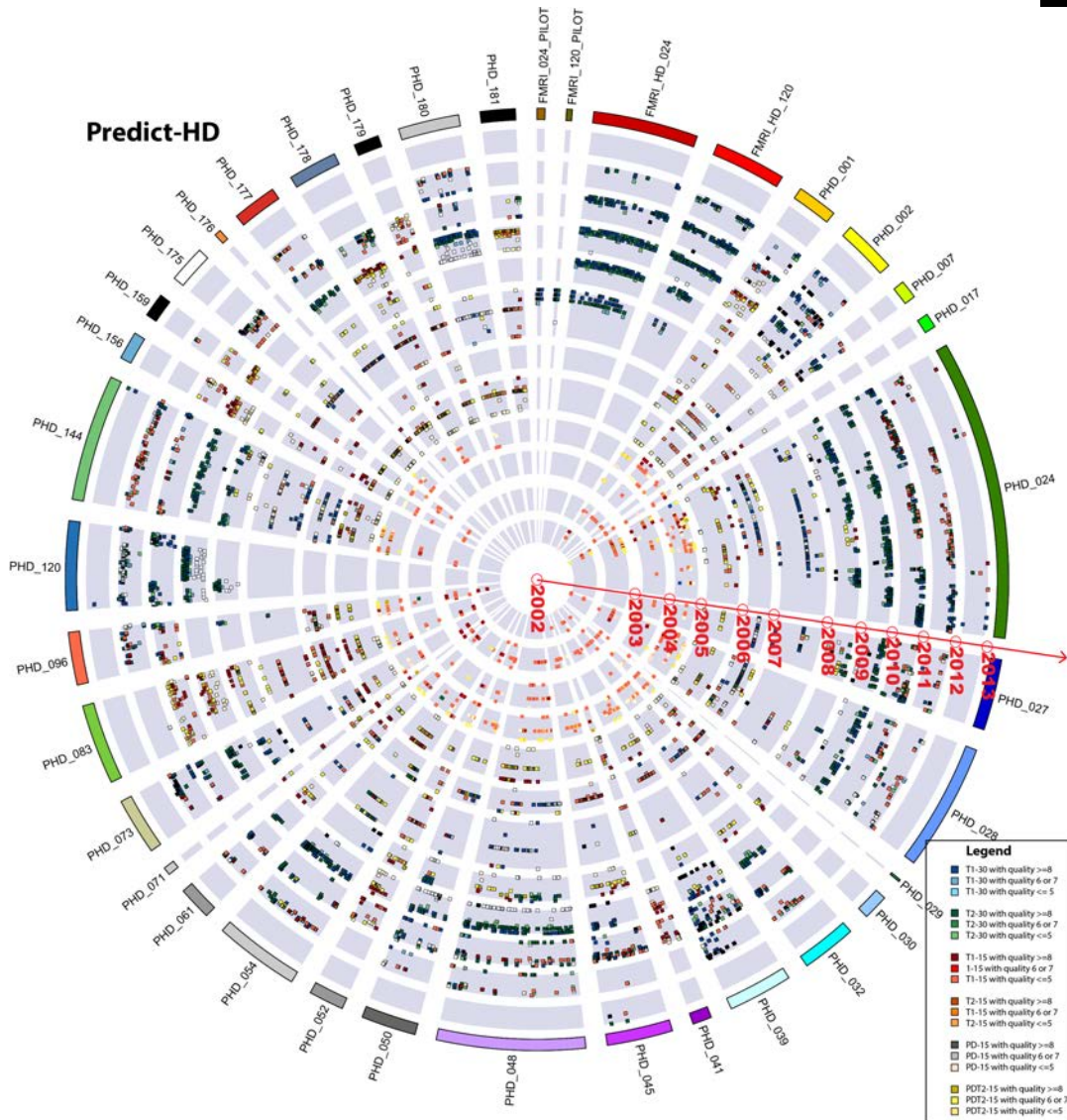
HDNI currently contains **4 projects, 454 subjects, and 1703 imaging sessions.**

Projects **Subjects** **MR** **PET** **CT**

Projects

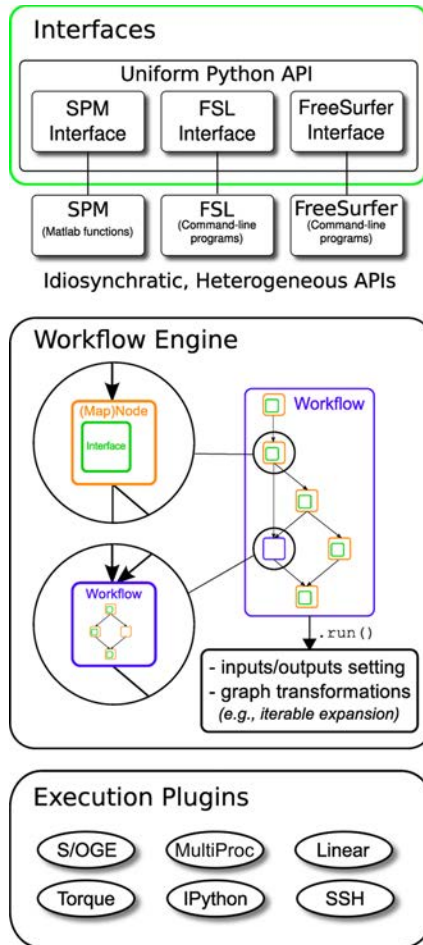


Improve Data Processing





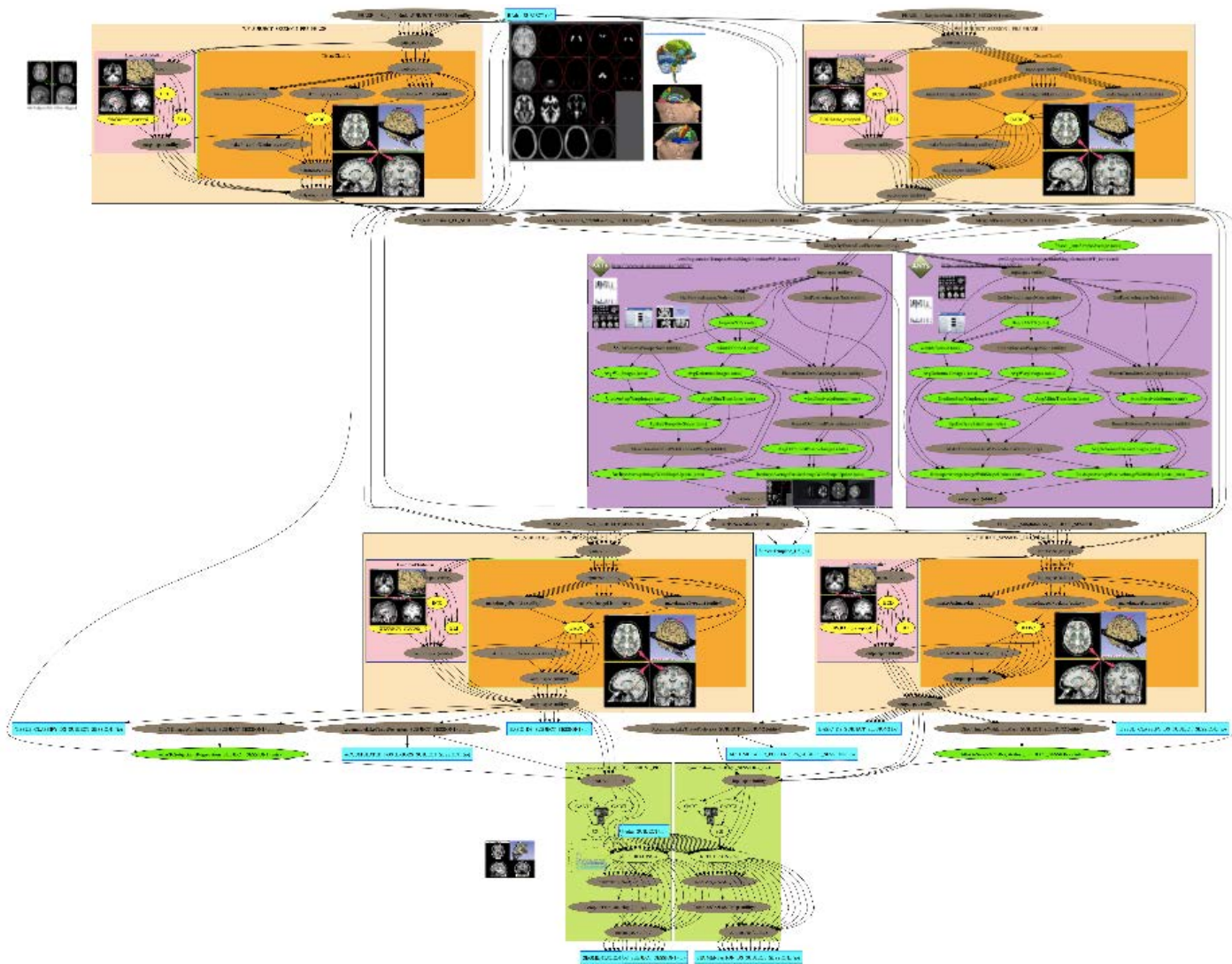
NiPype: Large catalog of tools with a uniform interface



- **Batch processing**
 - Distributed processing plugins
 - Reruns affect updated/edited node connections ONLY!
- **Uniform node creation**
 - Stable and consistent API
 - Nipype's Function node allows easy integration of CLI tools
- **Pipeline complexity**
 - Iterables, MapNodes
 - Nested workflows

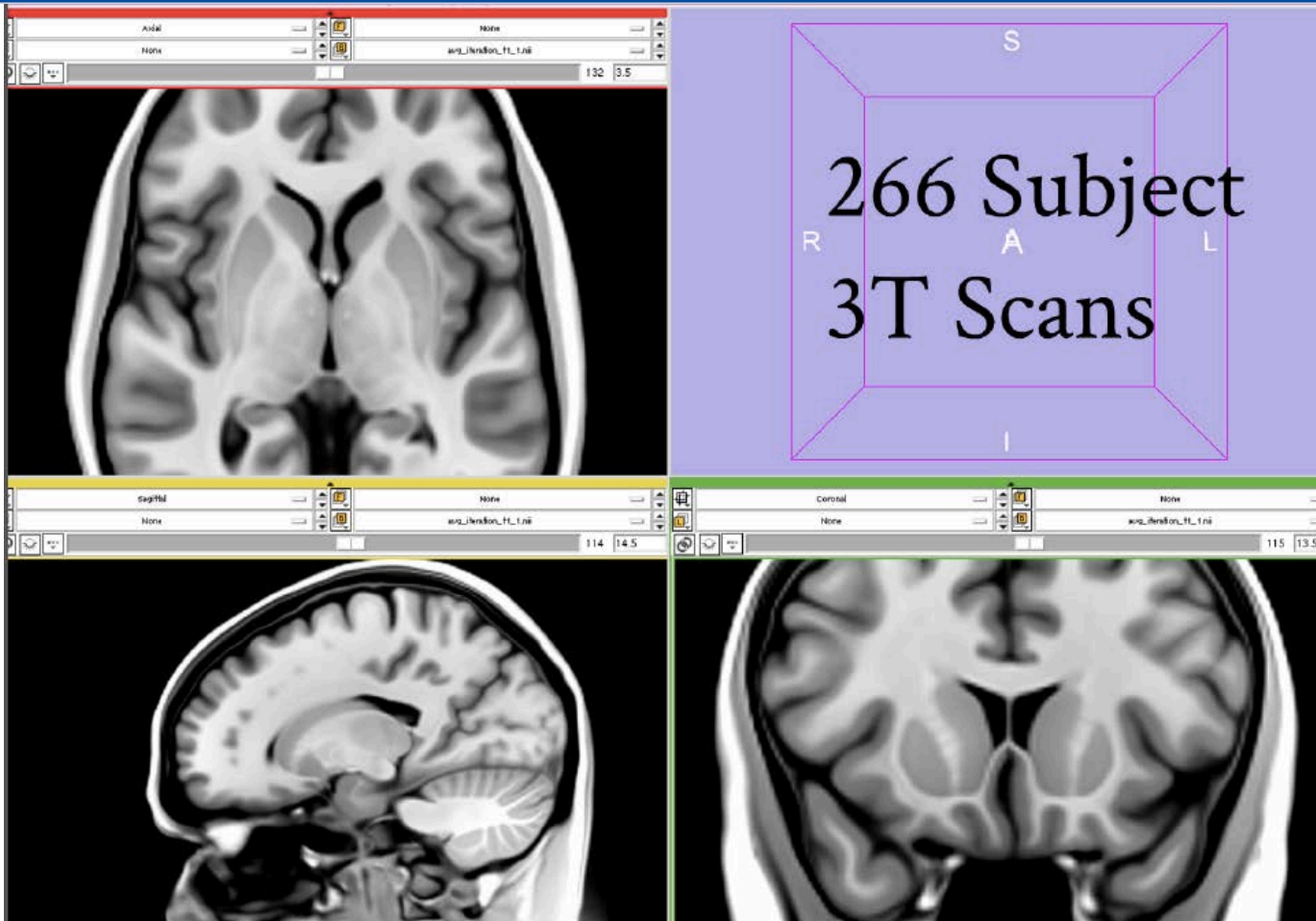


Diagram of Longitudinal Processing Pipeline





Improved Template Building with ANTS





Brain Sub-Cortical Structures: BRAINSCut (Longitudinal estimation)

Developed

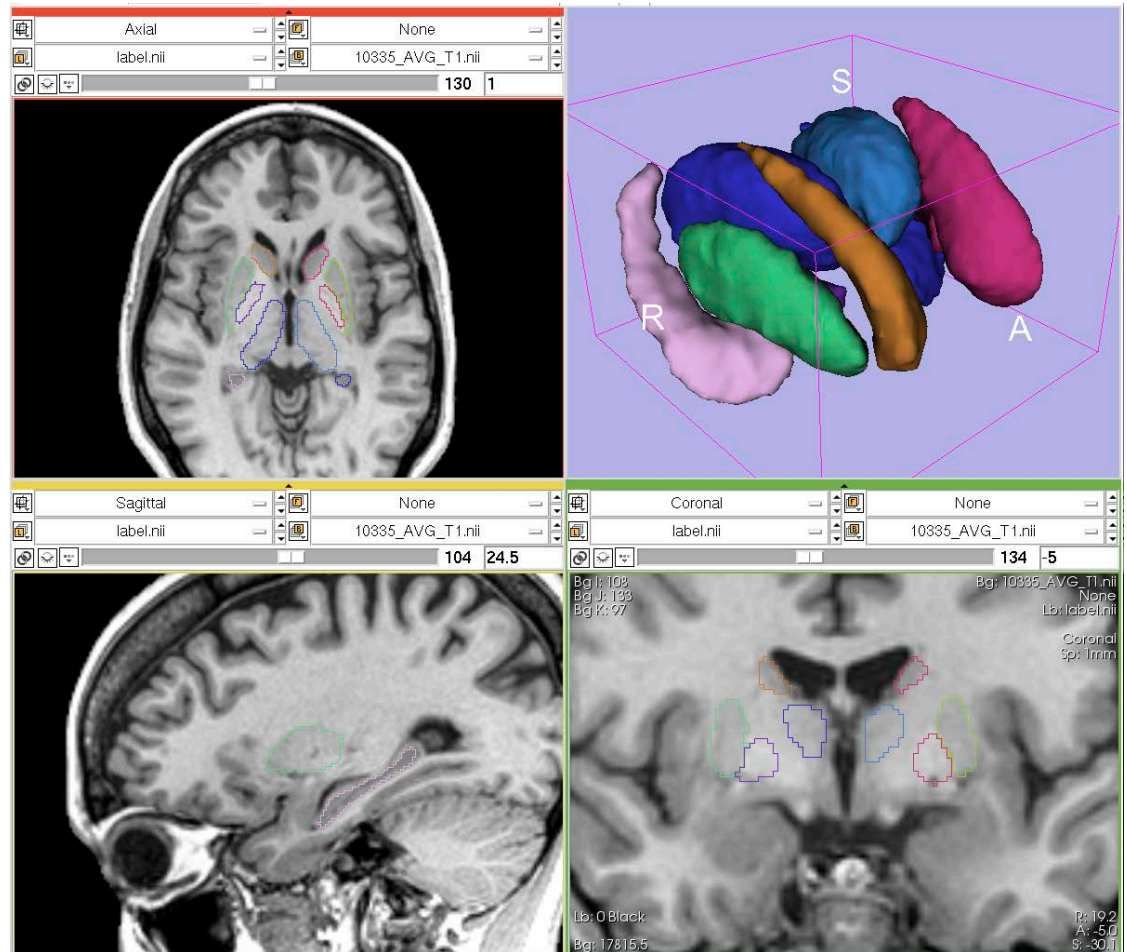
- Caudate
- Putamen
- Thalamus

New structure

- Hippocampus

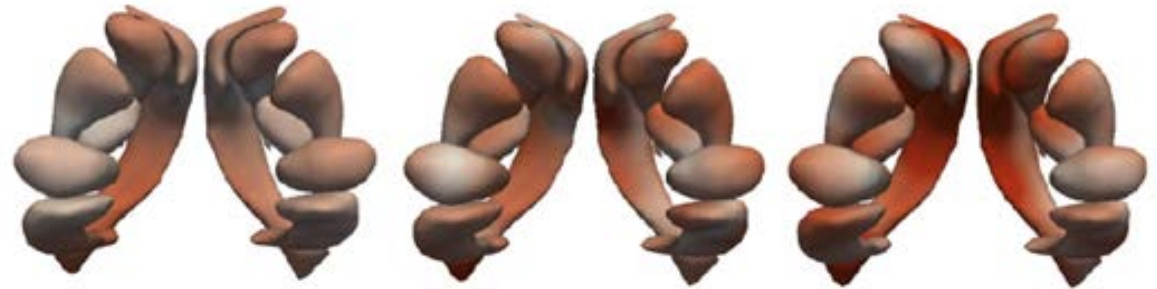
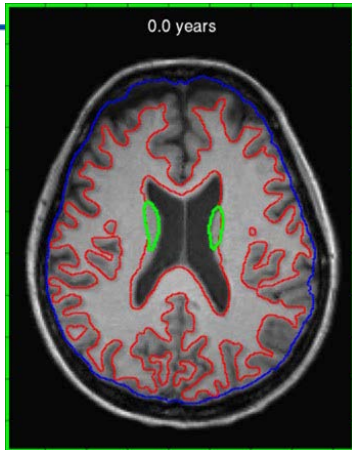
On going

- Globus
- Accumbens
- And more...





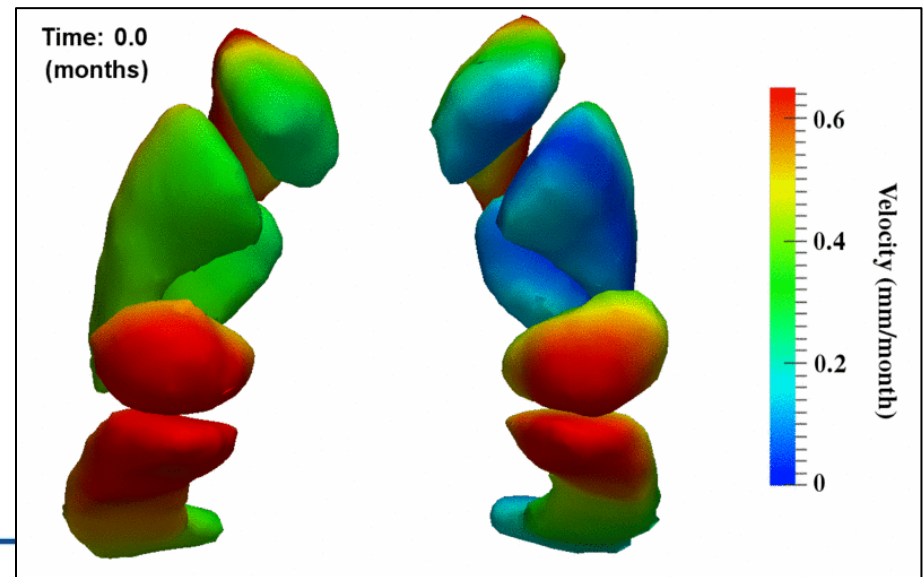
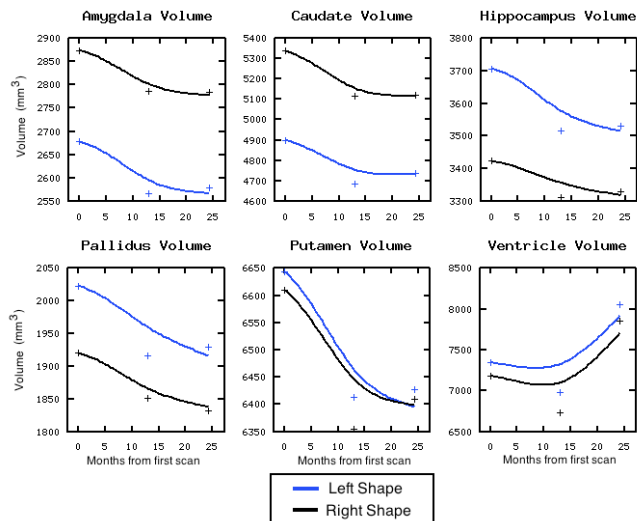
HD: Joint 4D modeling of subcortical structures from longitudinal MRI



Time point 1

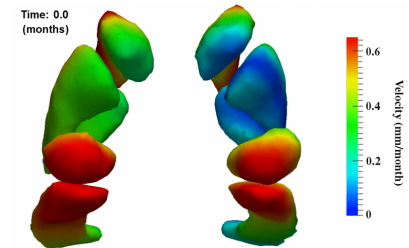
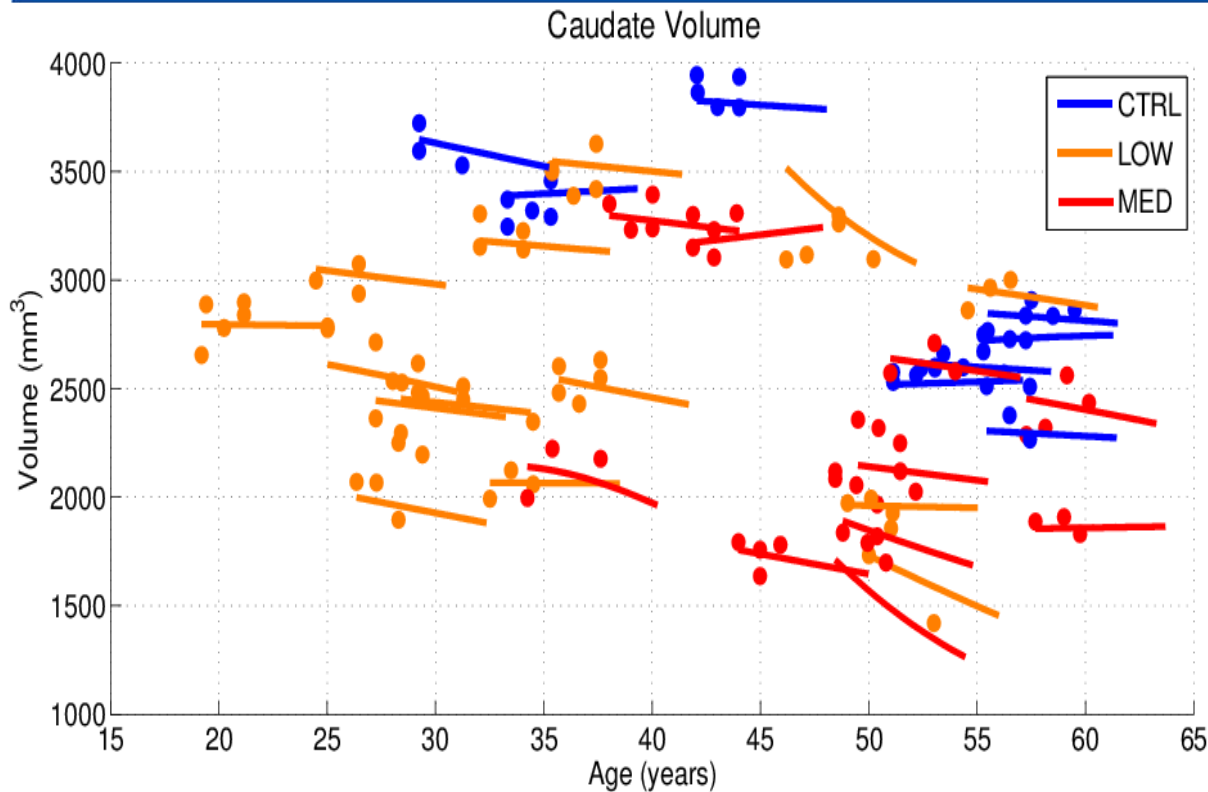
Time point 2

Time point 3





Personalized Shape Trajectory Modeling across Subjects



	PERCENT VOLUME CHANGE FROM SHAPE REGRESSION		
	CTRL	LOW	MED
CAUDATE	-0.78	-4.22	-6.25
PUTAMEN	-1.98	-6.76	-3.07
HIPPOCAMPUS	-0.65	-1.09	-2.17
THALAMUS	-0.74	-2.13	-1.18
ACUMBEN	-0.11	-2.13	-3.10
PALLIDUS	-3.11	-6.90	-7.07

Caudate volume for 32 subjects (3 time pts) extracted after shape regression. Observed volumes are shown as circles, which highlight the noise in segmentation.



Quality Assurance tools

The screenshot displays the 3DSlicer software interface with a Quality Assurance tool for DWI images. The interface is divided into several sections:

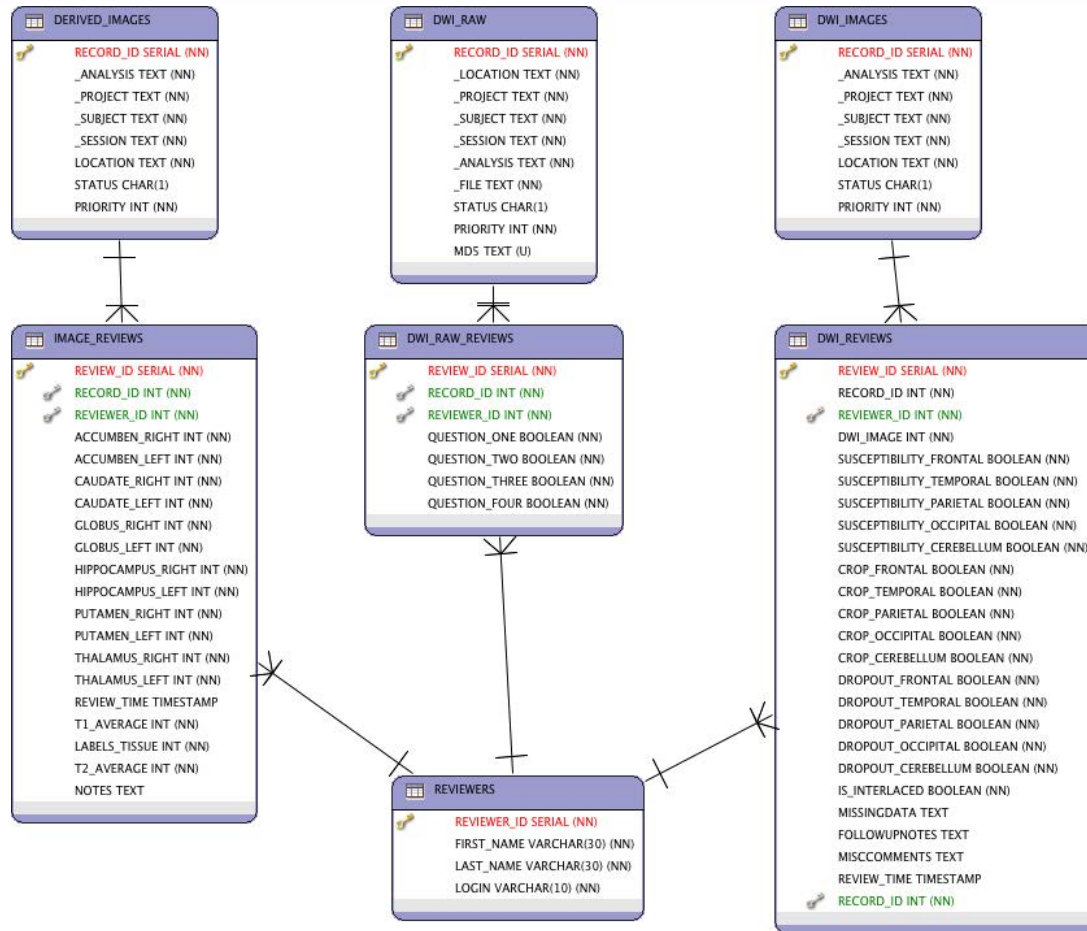
- Control Panel (Left):** Contains settings for Scalar Display, DWI Component, Lookup Table, Interpolate, and Window Level editor presets. It also includes a Histogram and an Image QA section with checkboxes for Artifacts (Susceptibility, Cropping, Dropout/Vibration), Missing Data, and a 'Get next DWI' button.
- Image Viewer (Center):** Shows a DWI image with a 3D grid overlay. The grid is labeled with 'S' (Superior), 'I' (Inferior), 'R' (Right), and 'L' (Left). A dialog box titled 'Gradient directions: 2422_18690_DWI-31_5.nrrd' is open, displaying a list of 12 gradient directions.
- Right Panel:** Shows a 3D grid overlay on a blue background, with labels 'S', 'I', 'R', and 'L' indicating the axes.

The dialog box displays the following gradient directions:

```
0000:=-0 0 0
0001:=-0.20 -0.51 -0.83
0002:=-0.19 0.51 0.83
0003:=-0.40 0.17 0.89
0004:=-0.40 -0.73 -0.54
0005:=-0.20 -0.94 -0.26
0006:=-0.85 -0.51 -0.05
0007:=-0.73 -0.51 -0.44
0008:=-0.40 -0.17 -0.89
0009:=-0.73 -0.17 -0.65
0010:=-0.65 -0.73 0.20
0011:=-0.32 -0.94 0.10
0012:=-0.32 -0.51 0.79
```




Quality Assurance tools





Data Reports

A	B	C	D	E	F
project	subject	session	ICV_2088	l_caudate	r_caudate_1
HDNI_001	015955315	015955315_20100803_30	1494658	829	
HDNI_001	015955315	015955315_20090622_30	1498021	947	1
HDNI_001	045416487	045416487_20110614_30	1413108	1714	1
HDNI_001	045416487	045416487_20080522_30	1412544	1941	2
HDNI_001	068044003	068044003_20090521_30	1770211	2580	2
HDNI_001	068044003	068044003_20100628_30	1756838	2575	2
HDNI_001	068044003	068044003_20080609_30	1746901	2729	2

dataset data_dictionary comments_scores +

A	B	C	D	E	F
"column_descriptors_id"	"column_descriptors_descripti"	"measurement_code_label"	"biological_region_label"	"experiment_label"	"tool_label"
cranial_csf_358		mm_3	cranial_csf	20130109_TrackOn_Results	BRAINSTools_20130109
r_hippocampus_162		mm_3	r_hippocampus	20130109_TrackOn_Results	BRAINSTools_20130109
ICV_20882		mm_3	ICV	20130109_TrackOn_Results	binaryBrainMask_20130401
r_caudate_134		mm_3	r_caudate	20130109_TrackOn_Results	BRAINSTools_20130109
l_thalamus_106		mm_3	l_thalamus	20130109_TrackOn_Results	BRAINSTools_20130109
l_globus_330		mm_3	l_globus	20130109_TrackOn_Results	BRAINSTools_20130109
cerebellar_white_matter_20938		mm_3	cerebellar_white_matter	20130109_TrackOn_Results	BRAINSTools_20130109

dataset data_dictionary comments_scores +

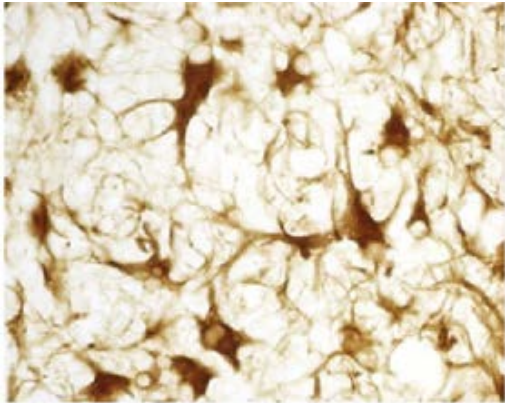
A	B	C	D	E	F
project_label	subject_label	session_label	column_descriptors_id	scores	commi
HDNI_001	560216908	560216908_20090402_30	l_hippocampus_50	75	Visual in
HDNI_003	034032275	034032275_20121001_30	r_putamen_190	95	Visual in
HDNI_002	058773254	058773254_20080609_30	l_putamen_78	95	Visual in
HDNI_003	658092030	658092030_20080204_30	r_globus_302	95	Visual in
HDNI_003	867249739	867249739_20110427_30	l_thalamus_106	95	Visual in
HDNI_003	541359481	541359481_20080611_30	r_caudate_134	95	Visual in
HDNI_004	287905458	287905458_20120801_30	r_thalamus_218	95	Visual in

dataset data_dictionary comments_scores +

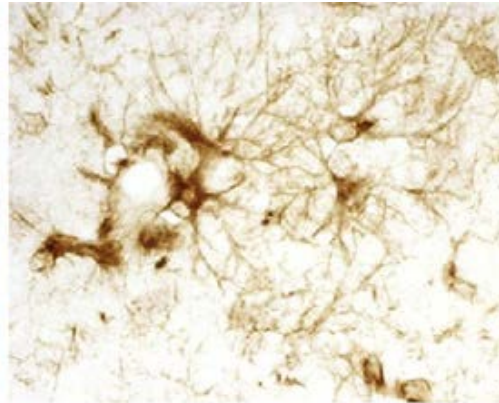


Tractography

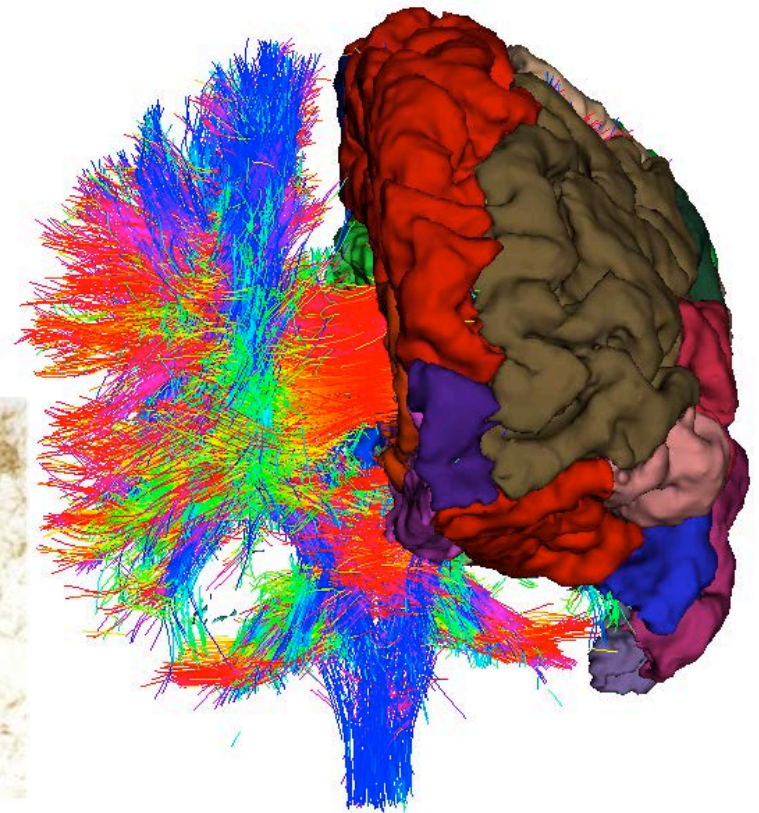
- It is clear that a more comprehensive approach is needed that could easily be extended as new questions arose. (i.e. support more exploration than direct a-priori interrogation)
- Van Camp 2012 Showed that loss of neurons can be measured by estimating “free-water” along tracks.



Normal



HD



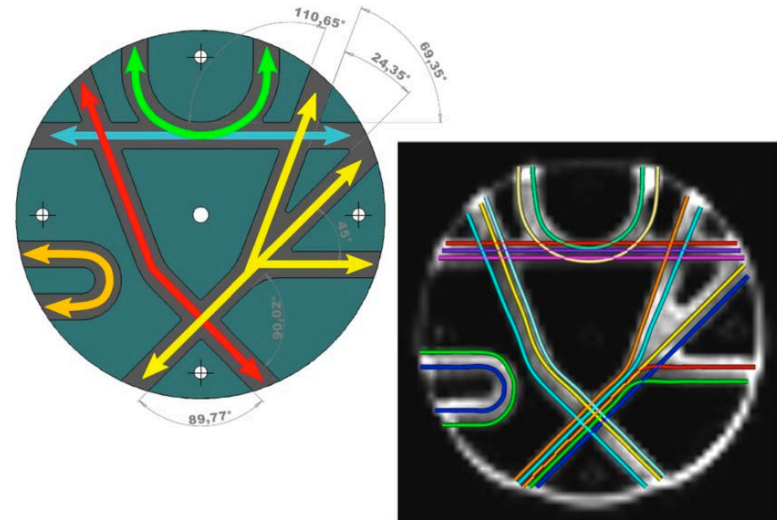


What is best tractography?

Quantitative evaluation of 10 tractography algorithms on a realistic diffusion MR phantom

Pierre Fillard ^{a,b,*}, Maxime Descoteaux ^c, Alvina Goh ^d, Sylvain Gouttard ^e, Ben Jeurissen ^f, James Malcolm ^g, Alonso Ramirez-Manzanares ^h, Marco Reisert ⁱ, Ken Sakaie ^j, Fatima Tensaouti ^k, Ting Yo ^l, Jean-François Mangin ^b, Cyril Poupon ^m

- ^a Parietal Research Team, INRIA Saclay Île-de-France, Neurospin, France
- ^b Laboratory of Computer-Assisted Neuro-Imaging, CEA Saclay, Neurospin, France
- ^c MOIVRE Center, Computer Science Department, Université de Sherbrooke, Canada
- ^d Department of Mathematics, National University of Singapore, Singapore
- ^e Scientific Computing and Imaging Institute, University of Utah, USA
- ^f IBBT-VisionLab, Department of Physics, University of Antwerp, Belgium
- ^g Psychiatry Neuroimaging Laboratory, Brigham and Womens Hospital, Harvard Medical School, USA
- ^h Mathematics Department, University of Guanajuato, Mexico
- ⁱ Department of Radiology, Medical Physics, University Hospital Freiburg, Germany
- ^j Imaging Institute, The Cleveland Clinic, Cleveland, USA
- ^k National Institute for Medical Research, INSERM, U825, France
- ^l Max Planck Institute for Human Cognitive and Brain Sciences, Germany
- ^m Imaging and Spectroscopy Laboratory, CEA Saclay, Neurospin, France



5 2-DT

Streamline tractography with filtered estimation of propagation direction

$3 \times 3 \times 3$,
 $b = 1500$

The diffusion model estimation is guided by the previous propagation direction using unscented Kalman filtering.

(Malcolm et al., 2009; Malcolm et al., 2010)

3D Slicer

James Malcolm



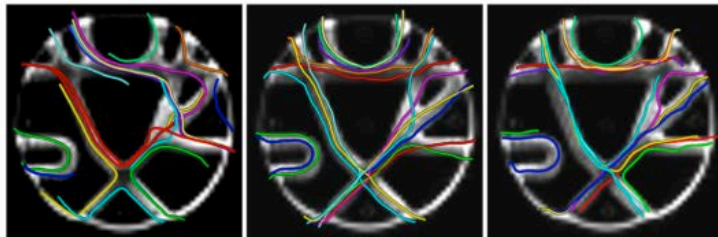
From Publication To Clinical Application



(1)

(2)

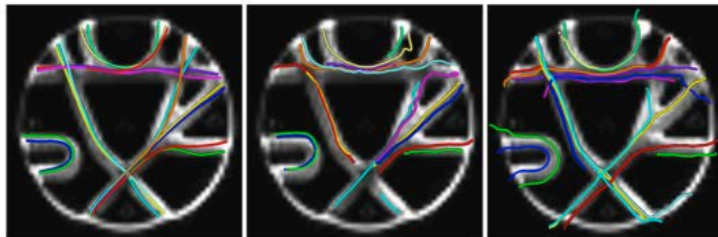
(3)



(4)

(5)

(6)



(7)

(8)

(9)



(10)

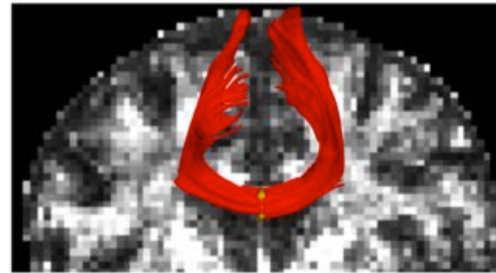


(11) - Ground truth

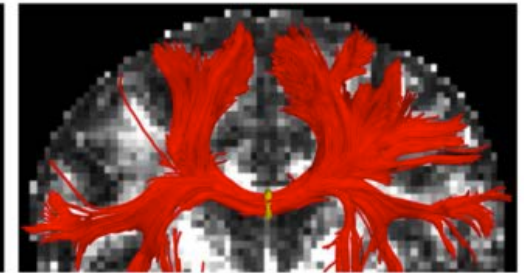
Filtered multi-tensor tractography

James G. Malcolm, Martha E. Shenton, and Yogesh Rathi

Psychiatry Neuroimaging Laboratory, Brigham and Womens Hospital, Harvard Medical School

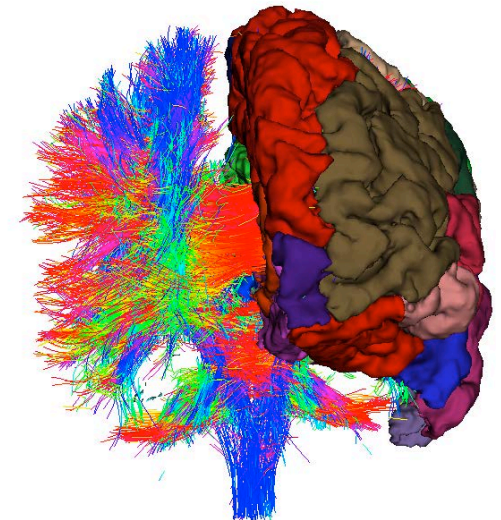


(a) Single-tensor streamline



(b) Filtered two-tensor

- * We Choose the Unscented Kahlman Filter algorithm “UKFTractography”
- * Improved speed and integration with Slicer



Computing



How to select tracts of interest?

- WMQL – White Matter Query Language

Table 1: Association Tract Definitions in WMQL: Cingulum bundle (CG); Extreme Capsule (EmC); and the fasciculus: Superior Longitudinal (SLF) sections I to III; Arcuate (AF); Inferior occipito frontal (IOFF); Middle Longitudinal (MdLF); Uncinate (UF)

CB.side = **only**((cingular.side or cingular.cortex.side) and (middle.frontal.side or cuneus.side or entorhinal.side or superior.frontal.side or inferior.parietal.side or fusiform.side or medial.orbitofrontal.side or lateral.orbitofrontal.side or parahippocampal.side or precuneus.side or lingual.side or centrum.semiovale.side))

EmC.side = **endpoints.in**(inferior.frontal.side or middle.frontal.side) and **endpoints.in**(inferior.parietal.lobule.side) and **temporal.side** and **insula.side** not in hemisphere.opposite

SLF_I.side = (superior.parietal.side and precuneus.side and superior.frontal.side or (superior.parietal.side and precuneus.side and superior.frontal.side and lateral.occipital.side) not in cingular.side not in temporal.side not in subcortical.side not in hemisphere.opposite

SLF_II.side = (inferior.parietal.side or supramarginal.side or lateral.occipital.side) and **endpoints.in**(middle.frontal.side) not in temporal.side not in subcortical.side not in hemisphere.opposite

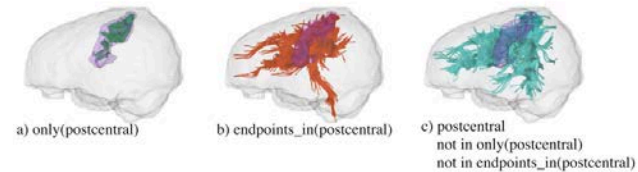
SLF_III.side = ((inferior.parietal.side or supramarginal.side or lateral.occipital.side) and **endpoints.in**(inferior.frontal.side) not in temporal.side not in subcortical.side not in hemisphere.opposite

AF.side = (inferior.frontal.side or middle.frontal.side or precentral.side) and (superior.temporal.side or middle.temporal.side) not in medial.of(supramarginal.side) not in subcortical.side not in hemisphere.opposite

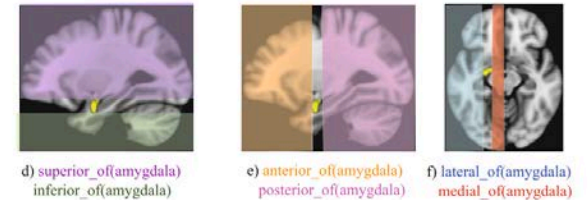
IOFF.side = (lateral.orbitofrontal.side and occipital.side) and **temporal.side** not in subcortical.side not in cingular.side not in superior.parietal.lobule.side not in hemisphere.opposite

ILF.side = **only**(temporal.side and occipital.side) and **anterior.of**(hippocampus.side) not in parahippocampal.side

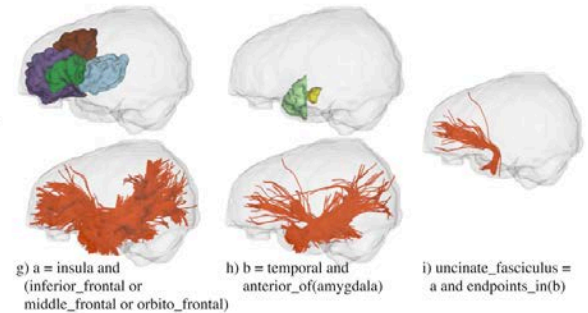
Anatomical Terms



Relative Position Terms



Logical Operations

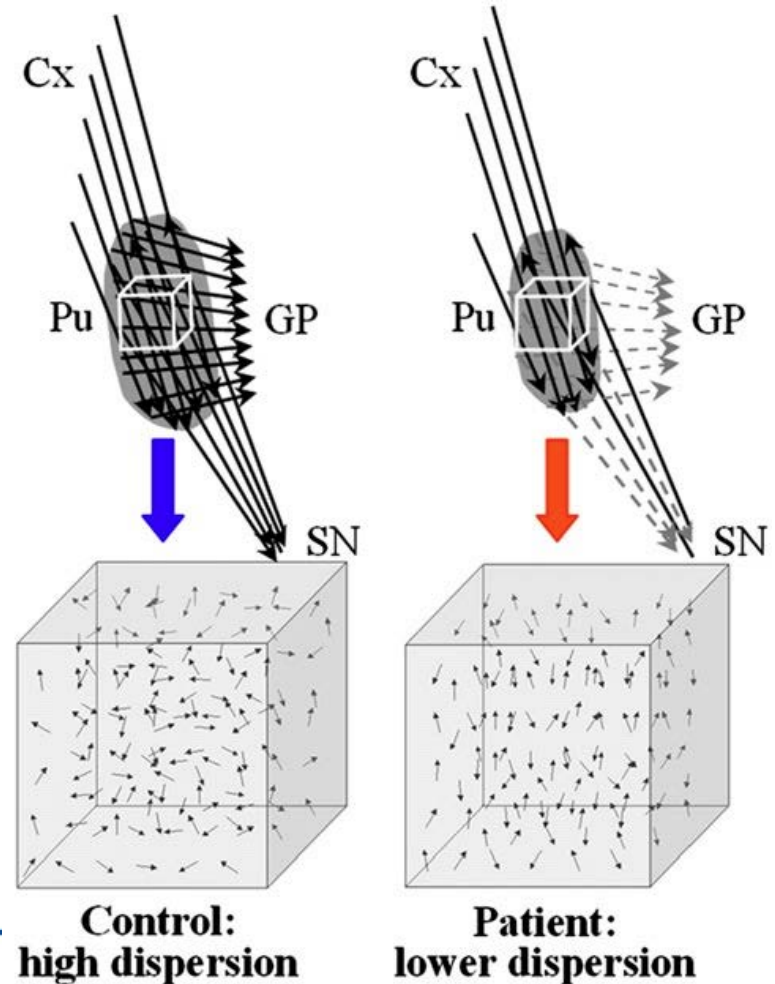
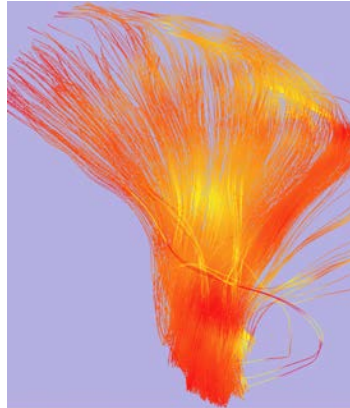




Dispersion is an important measure

- [Douaud09] describes the selective loss in fibers leading to lower dispersion in HD subjects (left).
- Need to translate methods that enable measurement of **dispersion**: A fiber tract connecting the substantia nigra in the brainstem to the striatum is colored by fiber dispersion [Savadjiev12].

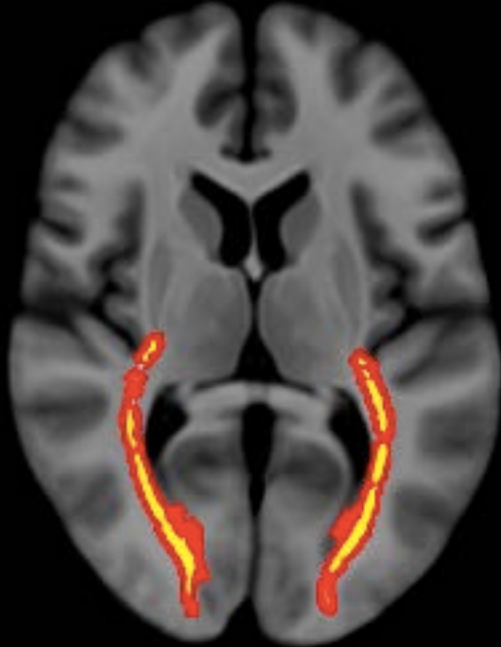
Working with Peter Savadjiev & Yogesh Rathi to transfer this methodology from prototype environment to production environment



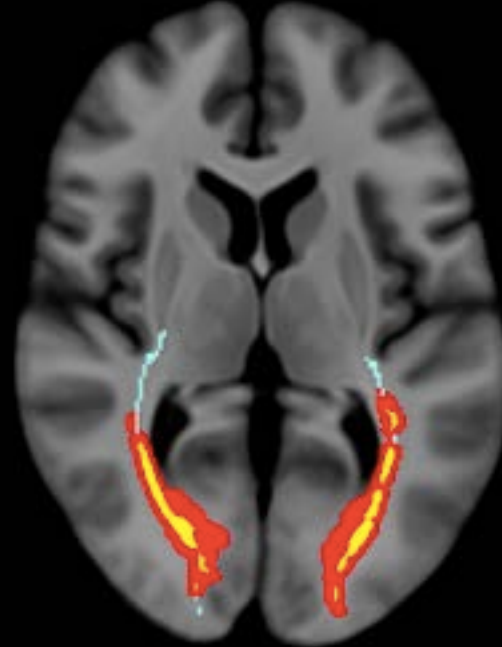


Preliminary Longitudinal DWI Processing Results

IFO, MD, C < H



IFO, RD, C < H





Share the Knowledge!



UNC-Utah NA-MIC framework for DTI fiber tract analysis

Audrey R. Verde^{1*}, Francois Budin¹, Jean-Baptiste Berger¹, Aditya Gupta^{1,2}, Mahshid Farzinfar¹, Adrien Kaiser¹, Mihye Ahn³, Hans Johnson⁴, Joy Matsui⁴, Heather C. Hazlett¹, Anuja Shama⁵, Casey Goodlett⁶, Yundi Shi¹, Sylvain Gouttard⁵, Clement Vachet^{1,5}, Joseph Piven¹, Hongtu Zhu³, Guido Gerig⁵ and Martin Styner^{1,7}

Psychiatry Research: Neuroimaging
Volume 214, Issue 3, 30 December 2013, Pages 389–394

Characterizing white matter health and organization in atherosclerotic vascular disease: A diffusion tensor imaging study

Kelly Rowe Bijanki^a, Stephan Arndt^{a,1}, Vincent A. Magnotta^{a, b, e}, David J. Moser^a, Peg Nopoulos^{a, c, d}, Sergio Paradiso^{a, g}, Joy T. Matsui^{a, e}, Hans J. Johnson^{a, e}

This repository Search or type a command Explore

INSTTools / BRAINTools

Processing focused on brain analysis

6 branches 1 releases

E.Y. Regina Kim^{1,2}, Hans J. Johnson^{1,2} and Norman K. Williams¹

Derivation of fiber tracts representing the corticospinal tract using anatomical landmarks

Joy Matsui^{1,2}, Eun Young Kim¹, Vincent Magnotta^{3,1}, and Hans Johnson¹

¹ University of Iowa, Department of Psychiatry, Iowa City, IA, USA
² University of Hawaii, John A. Burns School of Medicine, Honolulu, HI, USA
³ University of Iowa, Department of Radiology, Iowa City, IA, USA

Predictors of phenotypic progression in premanifest and early-stage Huntington's disease: TRACK-HD study: analysis of 36-month observational data

Rachael I. Scahill^a, Nicola Z. Hobbs^a, Miranda J. Say^a, Natalie Bechtel^a, Susie M.D. Henley^a, Harpreet Hyare^a, Douglas R. Langbehn^a, Rebecca Jones^a, Blair R. Leavitt^a, Raymond A.C. Roos^a, Alexandra Durr^a, Hans Johnson^a, Stéphane Léhericy^a, David Craufurd^a, Christopher Kennard^a, Stephen L. Hicks^a, Julie C. Stout^a, Ralf Reilmann^a, Sarah J. Tabrizi^a and the TRACK-HD investigators

Frontiers in Neurosciences

Robust Multi-site MR Data Processing: Iterative Optimization of Bias Correction, Tissue Classification, and Registration

Eun Young Kim¹ and Hans J. Johnson^{1,2*}

¹ University of Iowa, Biomedical Engineering Dept., Iowa City, IA, USA
² University of Iowa Hospital and Clinic, Psychiatry, Iowa City, IA, USA

Correspondence*:
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Hans J. Johnson, hans-johnson@uiowa.edu

Robust Multi-site MR Data Processing: Iterative Optimization of Bias Correction, Tissue Classification, and Registration

Eun Young Kim¹ and Hans J. Johnson^{1,2*}

¹ University of Iowa, Biomedical Engineering Dept., Iowa City, IA, USA
² University of Iowa Hospital and Clinic, Psychiatry, Iowa City, IA, USA

Correspondence*:
Eun Young Kim, eunyoungkim@uiowa.edu
Hans J. Johnson, hans-johnson@uiowa.edu

Striatal and white matter predictors of estimated diagnosis for Huntington disease

Jane S. Paulsen^{a,d,*}, Peggy C. Nopoulos^{a,b,1}, Elizabeth Aylward^{f,2}, Christopher A. Ross^{g,h,i,3}, Vincent A. Magnotta^{c,5}, Andrew Juhl^{a,6}, Ronald K. Pierson^{a,7}, James Mills^{a,8}, Douglas R. Langbehn^{a,9}, and the PREDICT-HD Investigators and Coordinators

Clinical Impairment in Premanifest and Early Huntington's Disease is Associated with Regionally Specific Atrophy

Rachael I. Scahill^{1a}, Nicola Z. Hobbs¹, Miranda J. Say¹, Natalie Bechtel², Susie M.D. Henley³, Harpreet Hyare⁴, Douglas R. Langbehn⁵, Rebecca Jones⁶, Blair R. Leavitt⁶, Raymond A.C. Roos⁷, Alexandra Durr⁸, Hans Johnson⁹, Stéphane Léhericy¹⁰, David Craufurd¹¹, Christopher Kennard¹², Stephen L. Hicks¹³, Julie C. Stout¹⁴, Ralf Reilmann¹⁵, Sarah J. Tabrizi¹⁶ and the TRACK-HD investigators

Longitudinal change in regional brain volumes in prodromal Huntington disease

Elizabeth H Aylward¹, Peggy C Nopoulos², Christopher A Ross³, Douglas R Langbehn², Ronald K Pierson², James A Mills², Hans J Johnson², Vincent A Magnotta², Andrew R Juhl², Jane S Paulsen², the PREDICT-HD Investigators and Coordinators

Biological and clinical changes in premanifest and early stage Huntington's disease in the TRACK-HD study: A longitudinal analysis

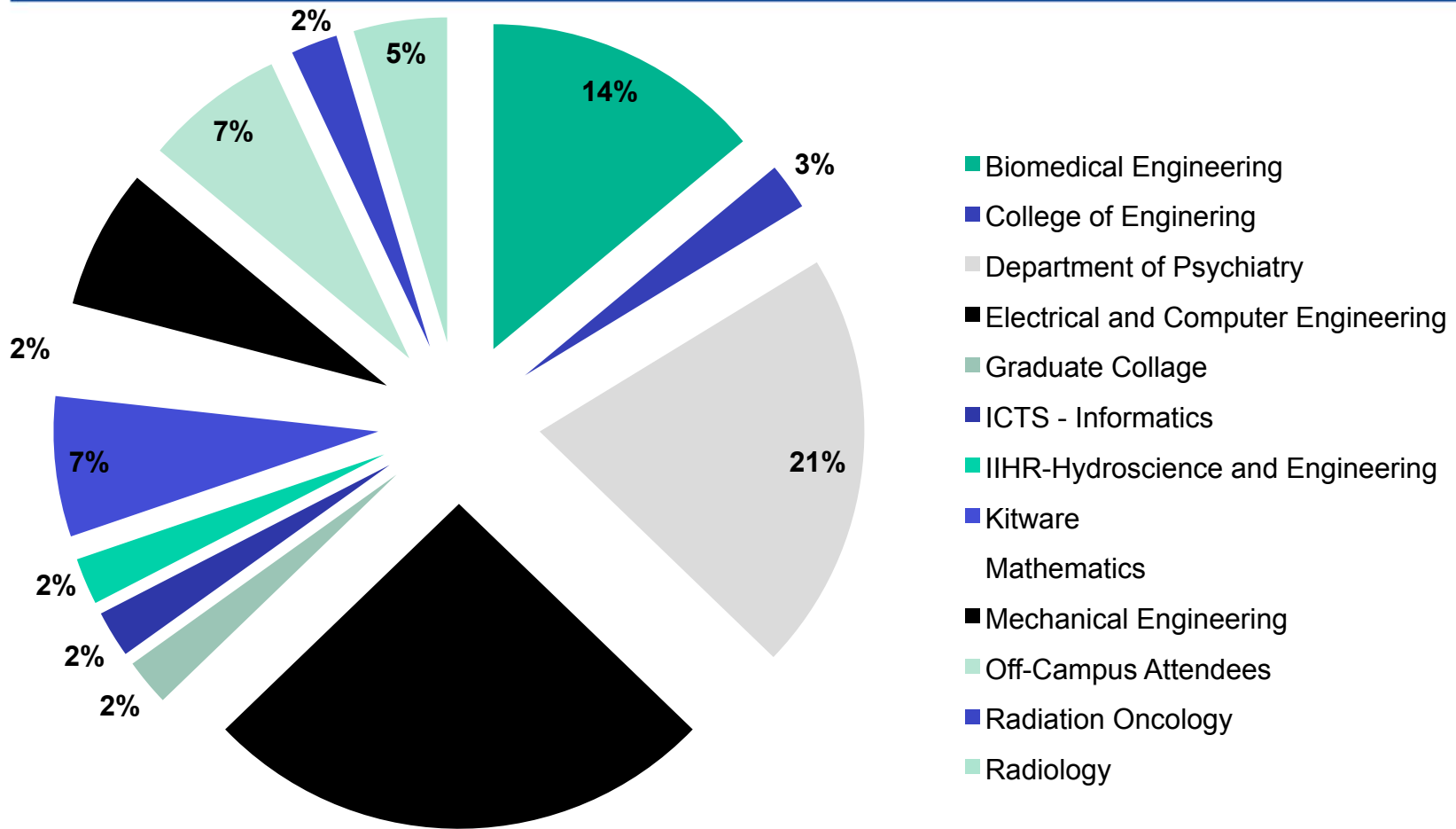
Rachael I. Scahill^{1a}, Nicola Z. Hobbs¹, Miranda J. Say¹, Natalie Bechtel², Susie M.D. Henley³, Harpreet Hyare⁴, Douglas R. Langbehn⁵, Rebecca Jones⁶, Blair R. Leavitt⁶, Raymond A.C. Roos⁷, Alexandra Durr⁸, Hans Johnson⁹, Stéphane Léhericy¹⁰, David Craufurd¹¹, Christopher Kennard¹², Stephen L. Hicks¹³, Julie C. Stout¹⁴, Ralf Reilmann¹⁵, Sarah J. Tabrizi¹⁶ and the TRACK-HD investigators

And Many Under



ITK/Simple ITK

ITK/Simple ITK Departmental Attendance





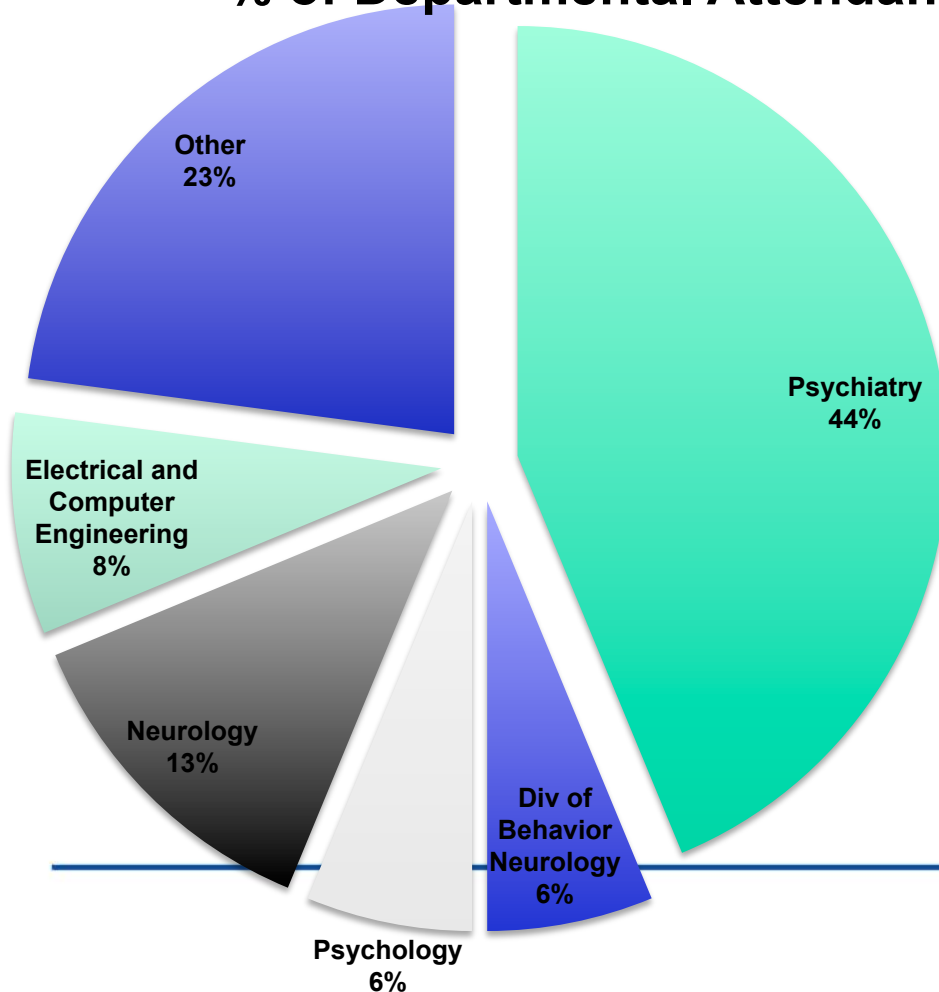
BRAINSCamp 2013

NA-MIC

National Alliance for Medical Image Computing

<http://www.na-mic.org>

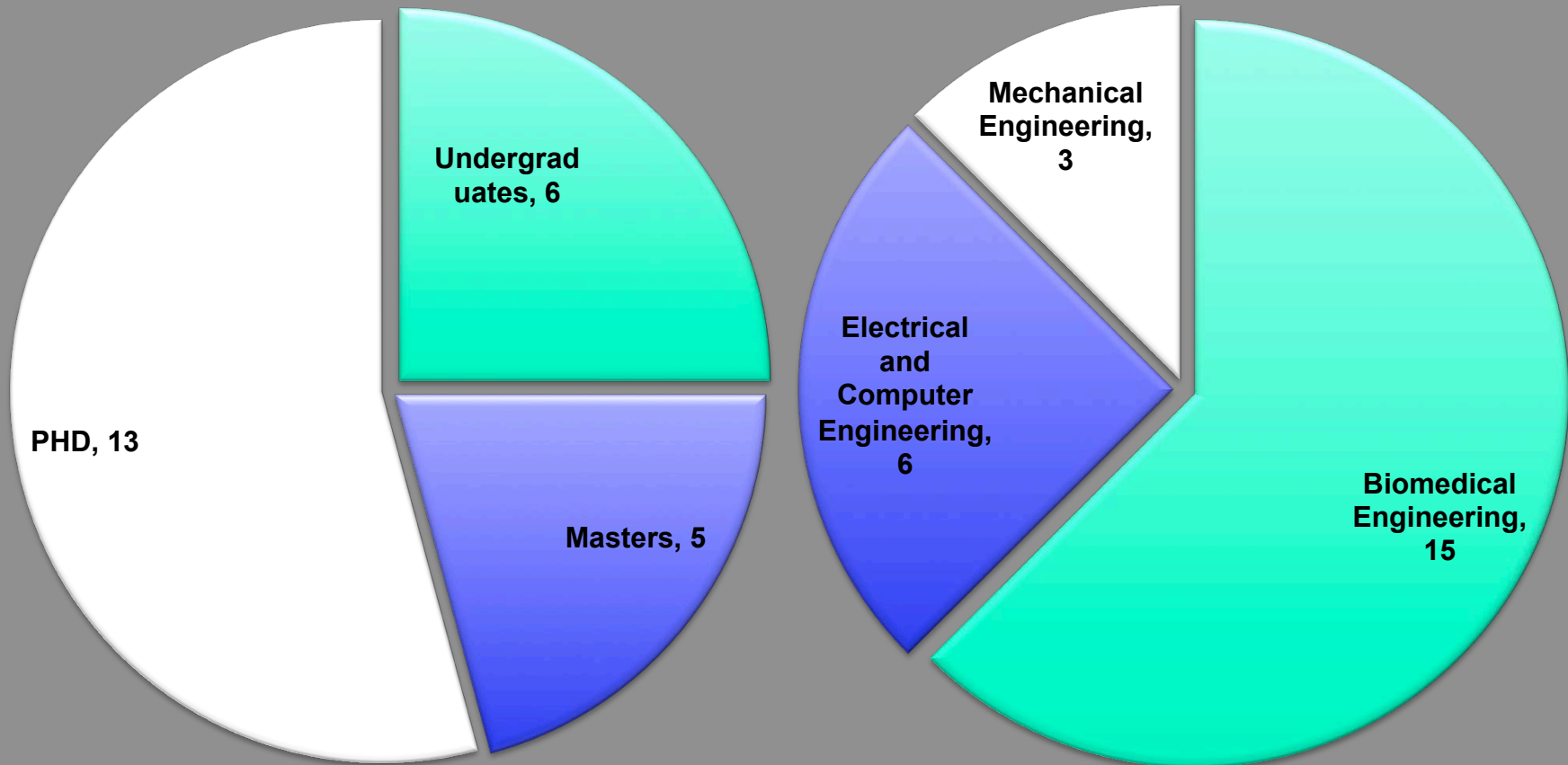
% of Departmental Attendance 41 Attendees



#	Answer	Bar
1	Very Unlikely	
2	Unlikely	
3	Undecided	
4	Likely	
5	Very Likely	
	Total	



Medical Image Processing Course 24 New “NA-MIC Certified” Engineers





Specific Aims

- Perform individualized longitudinal shape change quantification from multi-modal data. **(DONE!)**
- Complete full brain Diffusion Tensor Imaging tractography analysis. **(DONE!)**
- Deploy extensible tools for sharing source data, derived data, algorithms and methods to multi-site analysis teams. **(DONE!)**



Funded HD-Projects Using NAMIC DWI Processing Tools

- (CHDI) TRACK-HD/TRACK-ON (Private Foundation)
 - (R01 NS040068 12S1) PREDICT-HD
 - (U01 NS082083) *Functional Connectivity in Pre-manifest HD* (Stephen Rao, Cleveland Clinic)
 - (U01 NS082085) *Study of Basal Ganglia Shape Analysis and Circuitry* (Michael Miller & Chris Ross, JHU)
 - (U01 NS082086) *4D Shape Analysis for Modeling Spatiotemporal Change Trajectories in Huntington's* (Guido Gehrig, Utah)
 - (U01 NS083223) *Characterization of White Matter in HD Using Diffusion MRI* (CF Westin, Harvard)
 - (U01 NS082074) *Imaging and Genetics in Huntington's Disease* (Jessica Turner & Vincent Calhoun, MIND Institute)
 - (U54 EB005149) DBP-National Alliance For Medical Image Computing
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Thank You!

- Questions?