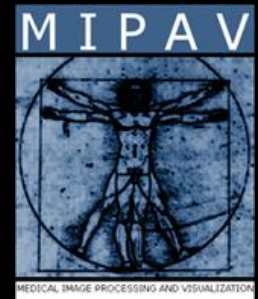


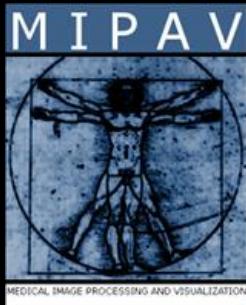


Creating a Streamlined Pipeline Utility for the Analysis of Universal Diffusion Tensor Imaging Data

Beth Tyrie

October 20, 2011

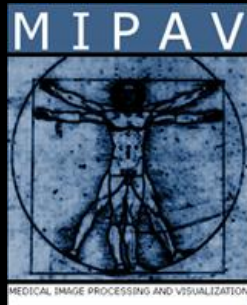




What is MIPAV?



- Medical Image Processing, Analysis, and Visualization
- Created by the Biomedical Imaging Research Services Section (BIRSS) of the Center for Information Technology (CIT)
- Enables quantitative analysis and visualization of medical images from numerous modalities (i.e. PET, MRI, CT, microscopy)



MIPAV Goals



- Develop computational methods and algorithms to analyze and quantify biomedical data
- Establish collaborations with NIH researchers and colleagues at other research centers
- Provide needed tools (in both hardware and software) to support the discovery and advancement of biomedical knowledge.

MIPAV Interface

The screenshot displays the MIPAV (Medical Image Processing and Visualization) software interface. The main window shows a 3D visualization of a brain slice, with a 2D slice index on the left. A histogram window is open, showing the intensity distribution of the slice. The interface includes a menu bar (File, VOI, LUT, Algorithms, Utilities, Systems analysis, Plugins, Scripts, Image, Toolbars, Help) and a toolbar with various tools. The histogram window shows the following data:

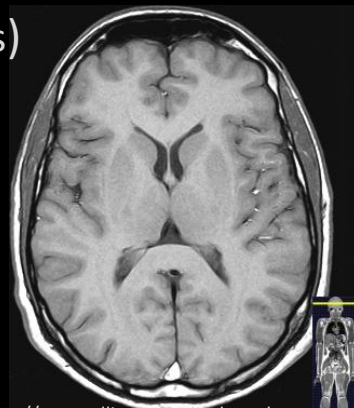
Image Intensities	Count
0.0	2
4.27	5
462.70	28
1242.0	804
6457.58	248

The histogram window also includes a 'Log scale (Histogram)' checkbox, a 'Number of colors' field set to 156, and a 'LUT' field. The X Range is 458.70 to 466.70, and the Y Range is 1 to 248. The X Scale is 458.70 to 466.70. The 2D slice index shows a total of 32 slices, with the current slice being 15. The 3D model shows a brain slice with a red box indicating the current slice position. The 2D slice window shows a brain slice with a red box indicating the current slice position. The 3D model shows a brain slice with a red box indicating the current slice position.

Conventional Magnetic Resonance Imaging vs. Diffusion Tensor Imaging

MRI

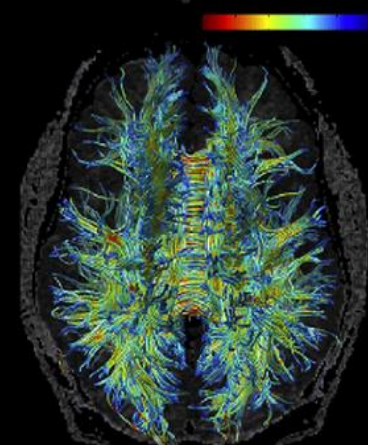
- Directly constructs an image of soft tissue in the body by using magnetic fields
- In the brain, MRI shows distinction between white and gray matter
 - Does not show fine detail of the white matter (neuronal tracts)



<http://www.alliancemedical.co.uk>

DTI

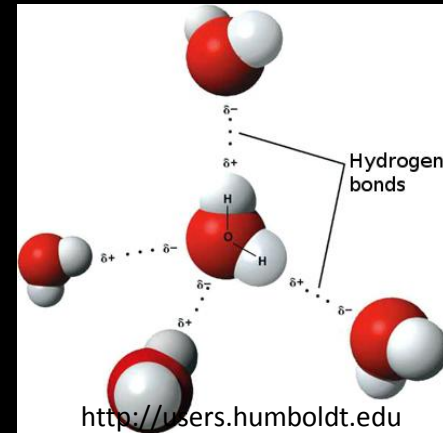
- Determines directionality and magnitude of water diffusion in the brain
 - This information is used calculate to fiber tract anatomy located in the white matter



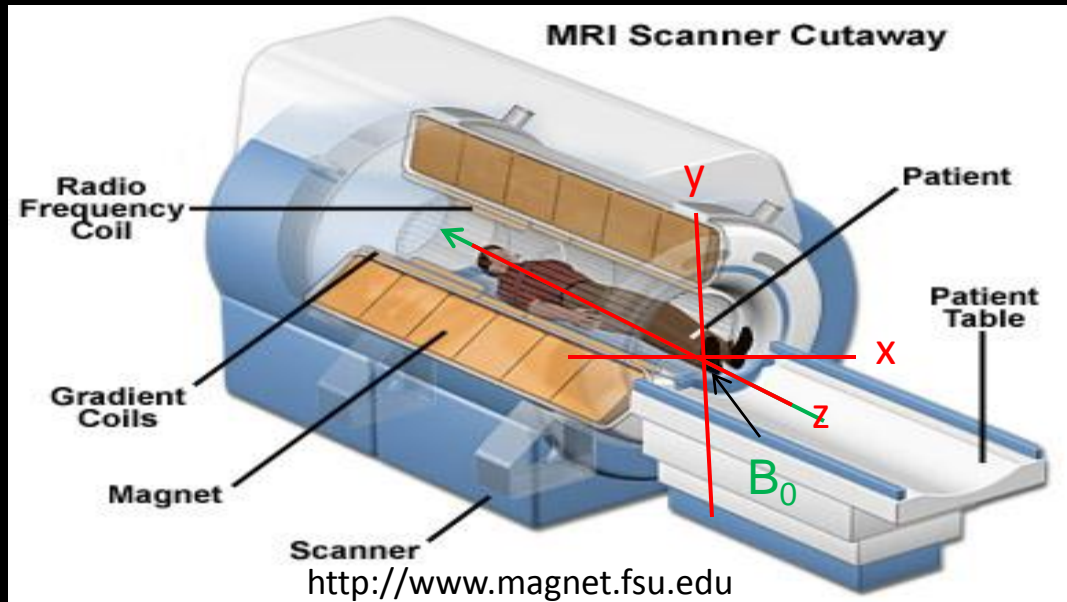
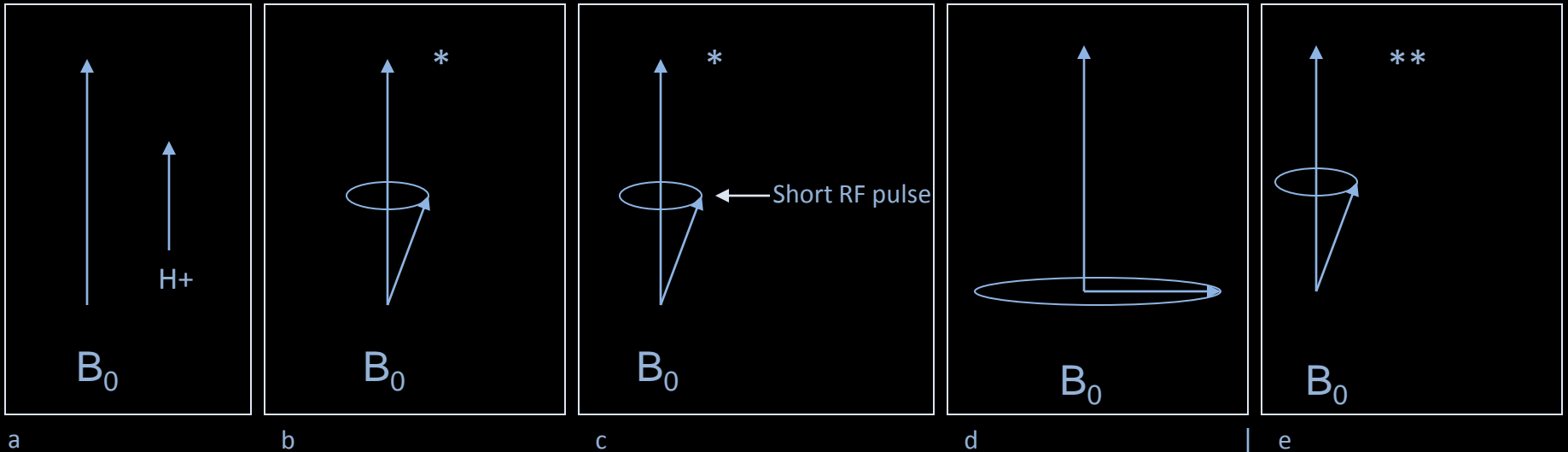
<http://lmi.bwh.harvard.edu>

Importance of Protons in MRI

- MRI images are primarily derived from the hydrogen protons (H^+) found in water
- Water is found in 70% to 90% of most body tissue and alters dramatically with disease and injury
- Protons become magnetized when immersed in an externally applied static field (B_0)



Fundamentals of MRI



Decaying Signal (FID)

X, Y, and Z gradients localize RF to allow "slices" of the body to be created

* Larmor freq = $B_0 \times$ gyromagnetic constant

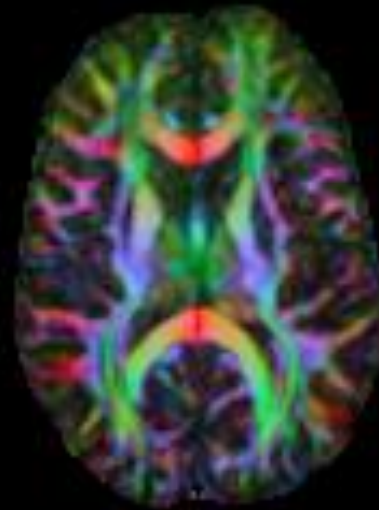
** Relaxation times are tissue dependent

Diffusion Tensor Imaging

MRI Imaging modality that indirectly images fibrous white matter brain tissue (nerve axons) by detecting water diffusion that tends to occur anisotropically along the nerve fibers.



<http://www.alliancemedical.co.uk>

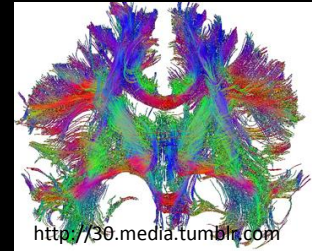


Advantages of DTI

- Non-invasive

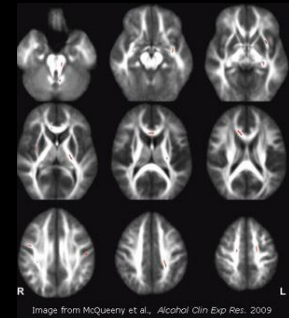


- Can measure water diffusion along any oblique angle

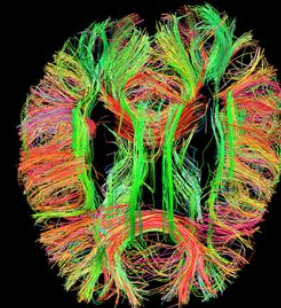


- Reveals detailed anatomy of white matter through fiber orientations

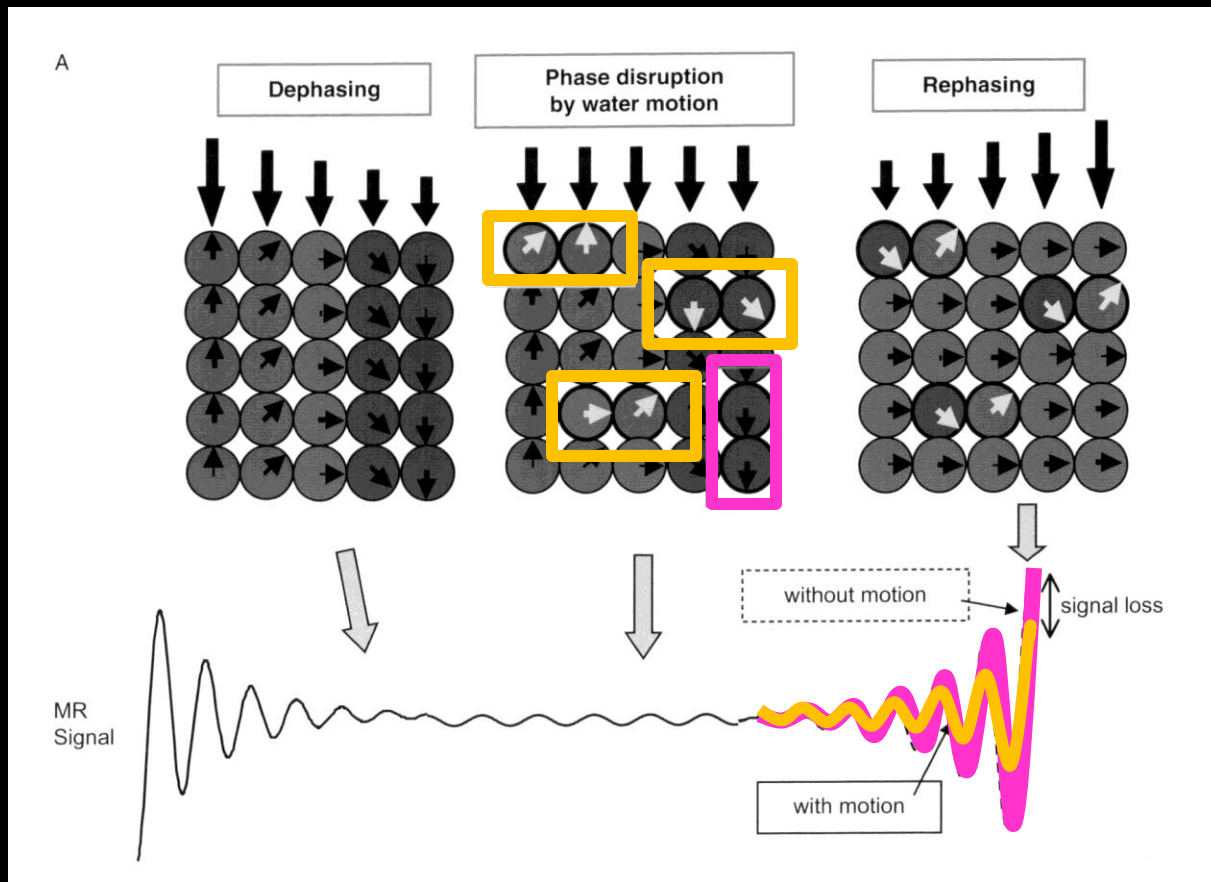
– MRI cannot view detailed anatomy due to white matter homogeneity in chemical composition



- Improves understanding of connectivity



Sensitizing the MRI Scan to Water Diffusion



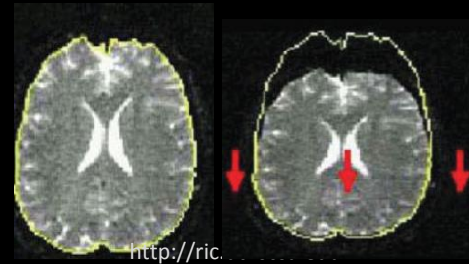
In the absence of diffusion, a rephasing gradient brings the spins back in phase. (no signal loss)

Computation of Diffusion Tensor and Applications



- Requirements for Tensor Calculation:

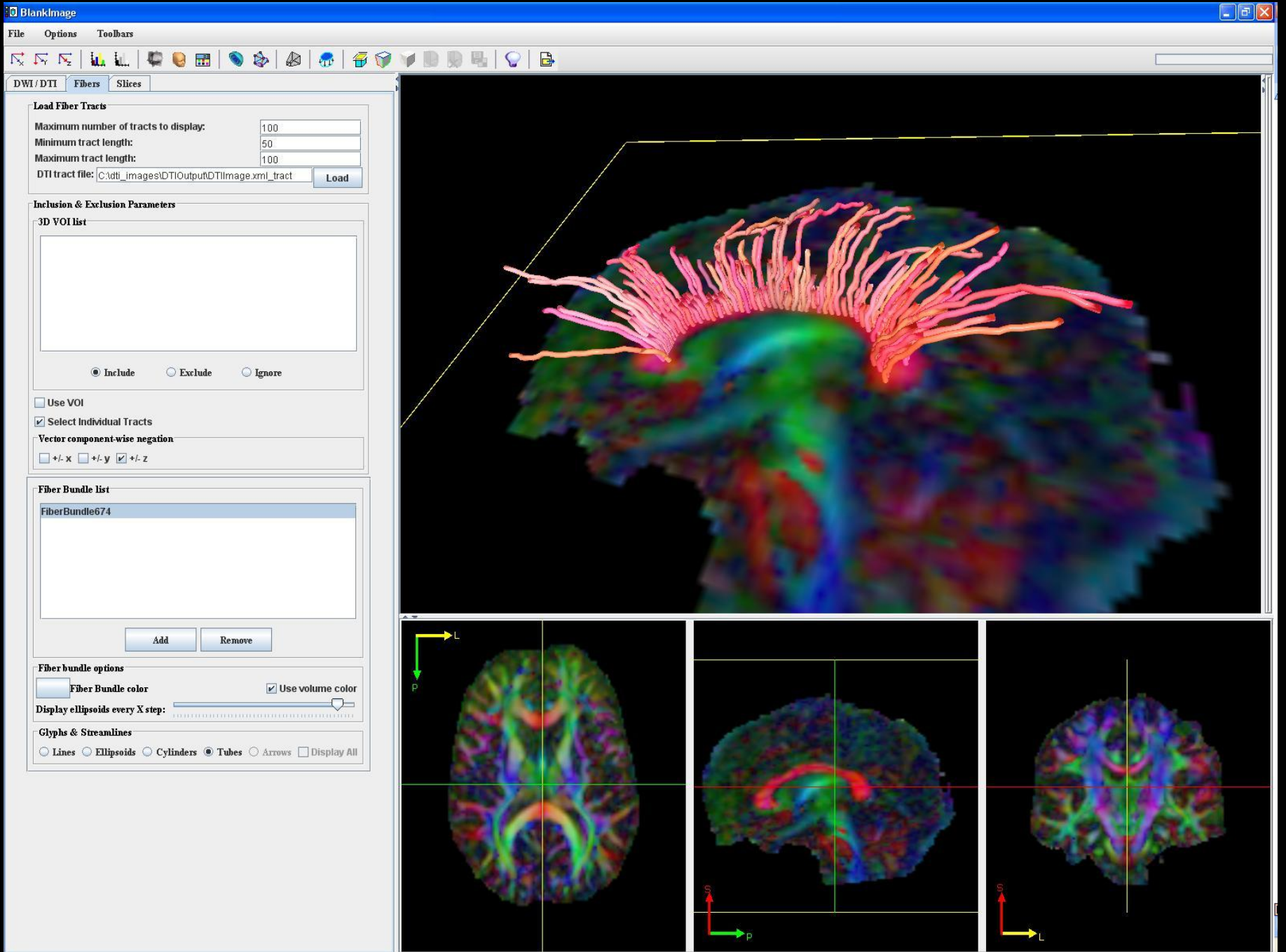
- I. A minimum of 7 scans from various directions
 - The first scan should be without the application of gradients with the B_0 set to zero
- II. Registration of the other scans in the Diffusion Weighted Imaging (DWI) data set to the first scan
- III. Distortion correction



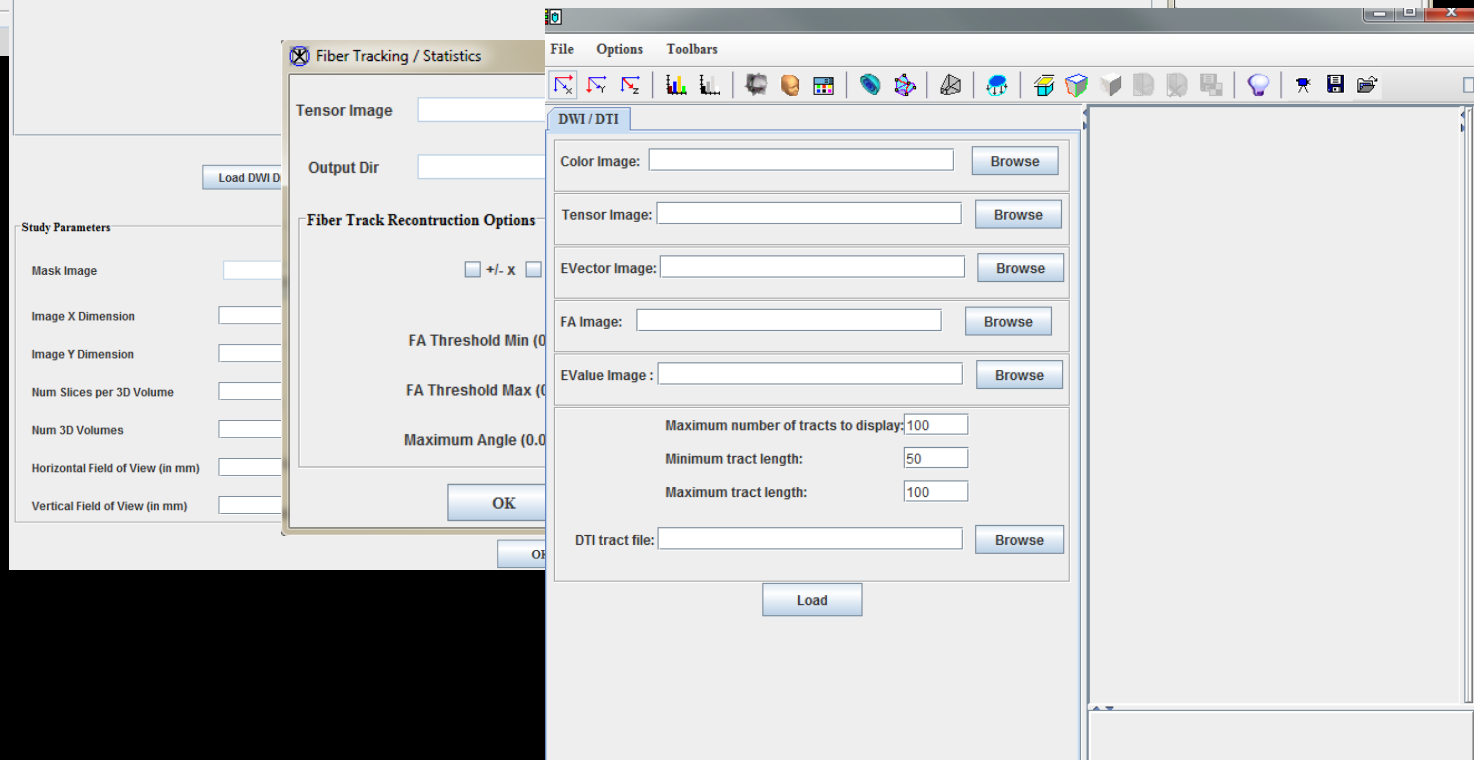
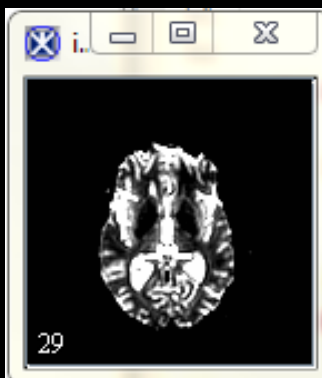
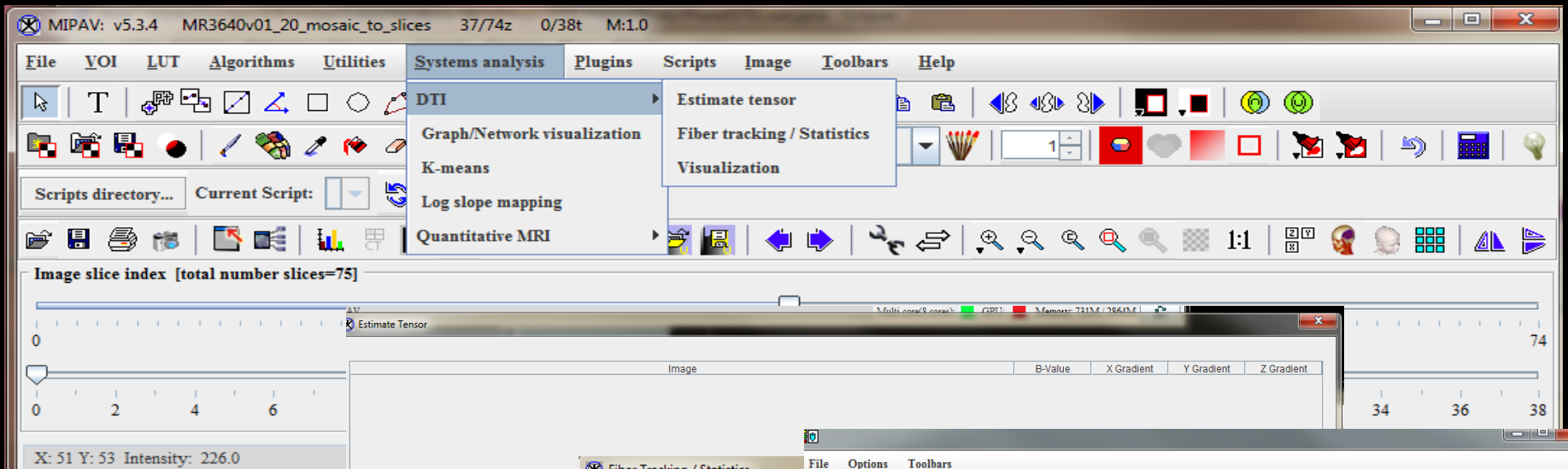
- MIPAV Tensor Calculation Applications:

- I. Fiber Tracking
- II. Visualization



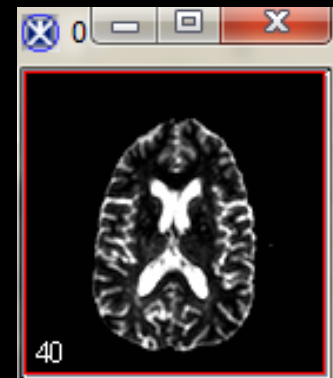
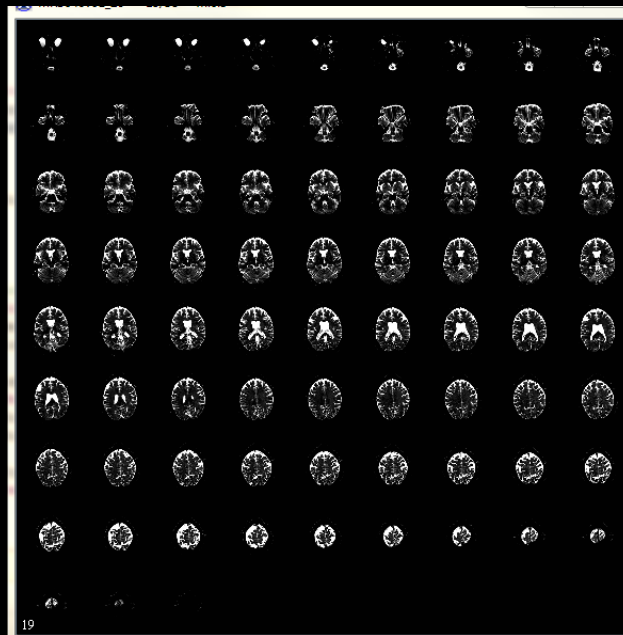
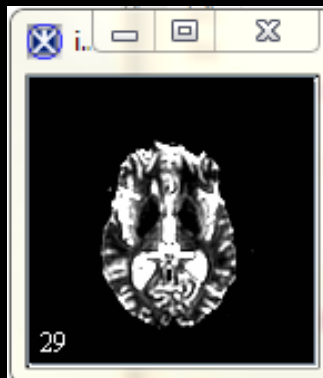


Current MIPAV DTI Pipeline



New MIPAV DTI Pipeline Project

to create a user-friendly DTI pipeline that has universal processing for all Diffusion Weighted Imaging (DWI) datasets acquired from any MRI scanner (i.e. Philips, Siemens, and GE)



New MIPAV DTI Pipeline Project

Volume	B-Value	X Gradient	Y Gradient	Z Gradient
0	0.0	0.70710677	0.0	0.70710677
1	0.0	0.70710677	0.0	0.70710677
2	300.0	-0.83203095	0.36696032	-0.38748828
3	300.0	0.8181228	-0.09182721	-0.56766444
4	1100.0	0.4652895	-0.16544881	-0.86955875
5	1105.0	0.44087344	0.76376194	-0.47148532
6	1100.0	0.27074987	-0.9347886	-0.22992377
7	1100.0	-0.84730834	0.20289089	-0.4908195
8	1100.0	0.33005223	0.9435225	0.02882328
9	1100.0	0.2650436	0.33748642	-0.9032468
10	1100.0	0.6487868	-0.6129902	-0.45090878
11	1100.0	-0.8871967	-0.15037885	-0.43619743
12	1100.0	-0.14302024	-0.4299764	-0.8914403

Siemens
(DICOM
Mosaic)

GE
(DICOM)

Philips
(DICOM,
PAR/REC)

DTI Parameters Object

Model Image

Registration

1. Motion Correction

Distortion Correction

Post-processing MIPAV
DTI Pipeline

1. Tensor Calculation
2. Fiber Tracking
3. Visualization

Completed MIPAV DTI Pipeline Steps

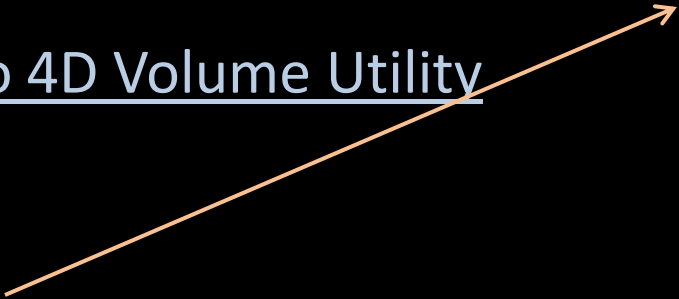
3D Mosaic to 4D Volume Utility

DTI Tab

- Automatic Population of Gradient/Bvalue Table for various image types

- Ability to save DTI parameters with image to be used for tensor calculation, fiber tracking, and visualization

- Save bvalue and gradient table to different text formats



File Type	Auto Pop of Bvals	Auto Pop of Grads
Philips P/R V3 & V4	✓	×
Philips P/R V4.1+	✓	✓
Siemens Mosaic DCM	✓	✓
GE DCM	?	?
Philips V4.1+ DCM		
fsl.txt	✓✓	✓✓
dtiStudio.txt	×	✓
mipavStandard.txt	✓	✓

The Import Data Step

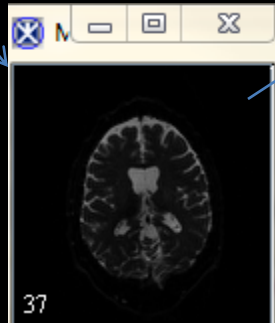
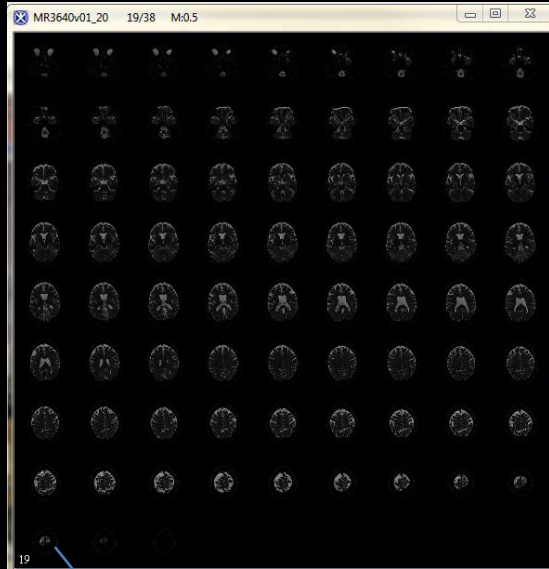
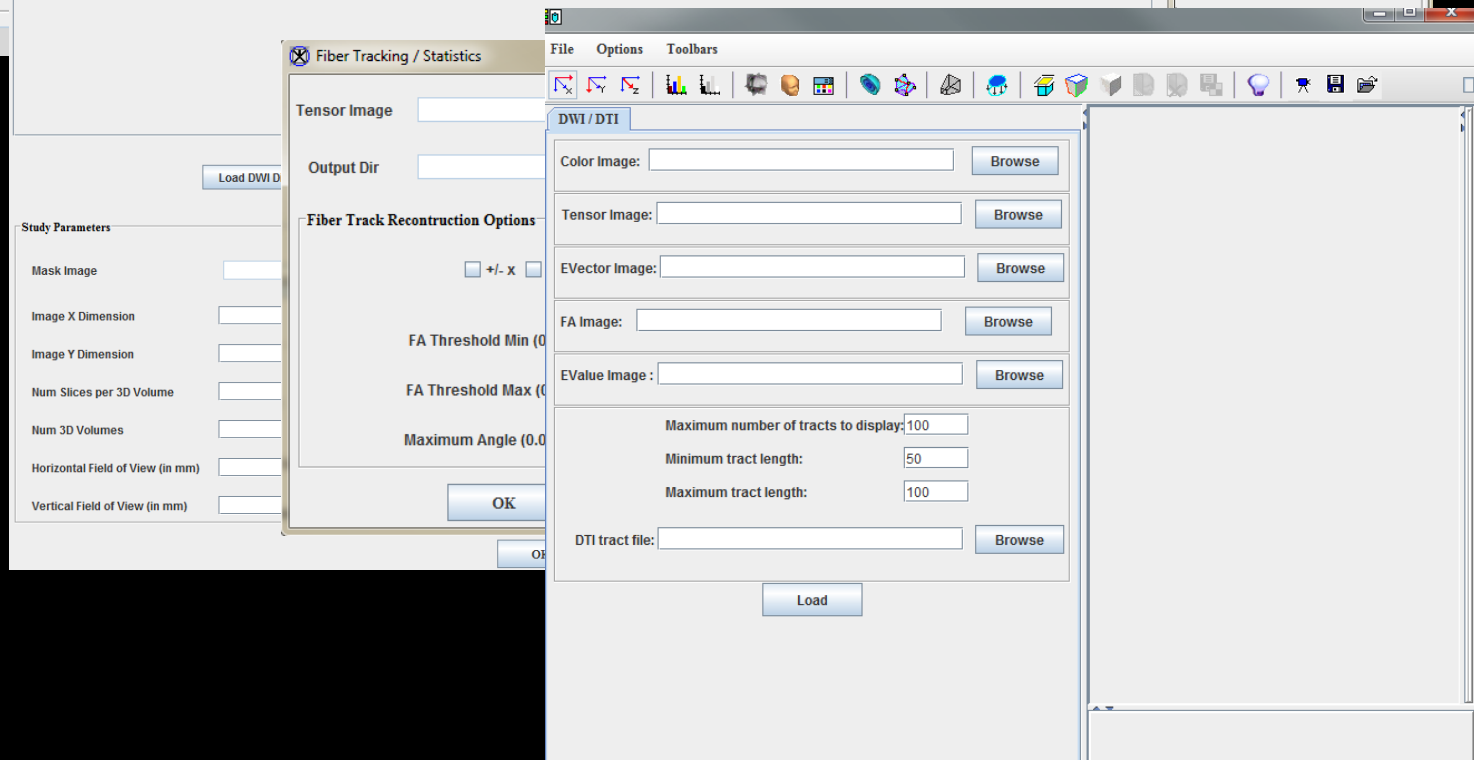
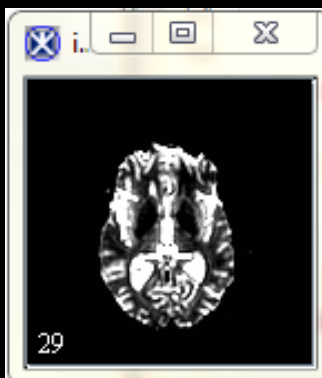
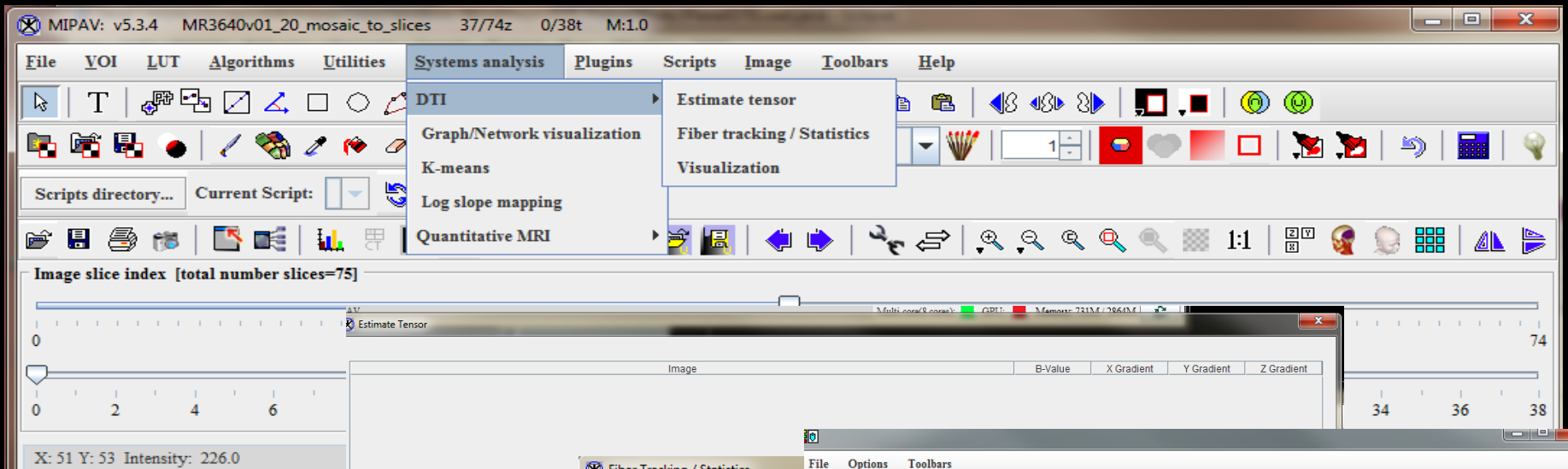


Image Attributes: MR3640v01_20_mosaic_to_slices 37 : 0

General	Resolutions	Orientations/Origin	Transform matrix	Talairach	D11
Volume	B-Value	X Gradient	Y Gradient	Z Gradient	
0	0.0	0.70710677	0.0	0.70710677	
1	0.0	0.70710677	0.0	0.70710677	
2	300.0	-0.83203095	0.39696032	-0.38748828	
3	300.0	0.8181228	-0.09182721	-0.56766444	
4	1100.0	0.4652895	-0.16544881	-0.86955875	
5	1105.0	0.44087344	0.76376194	-0.47148532	
6	1100.0	0.27074987	-0.9347886	-0.22992377	
7	1100.0	-0.84730834	0.20289089	-0.4908195	
8	1100.0	0.33005223	0.9435225	0.02882328	
9	1100.0	0.2650436	0.33748642	-0.9032468	
10	1100.0	0.6487868	-0.6129902	-0.45090878	
11	1100.0	-0.8871967	-0.15037885	-0.43619743	
12	1100.0	-0.14302024	-0.4299764	-0.8914403	

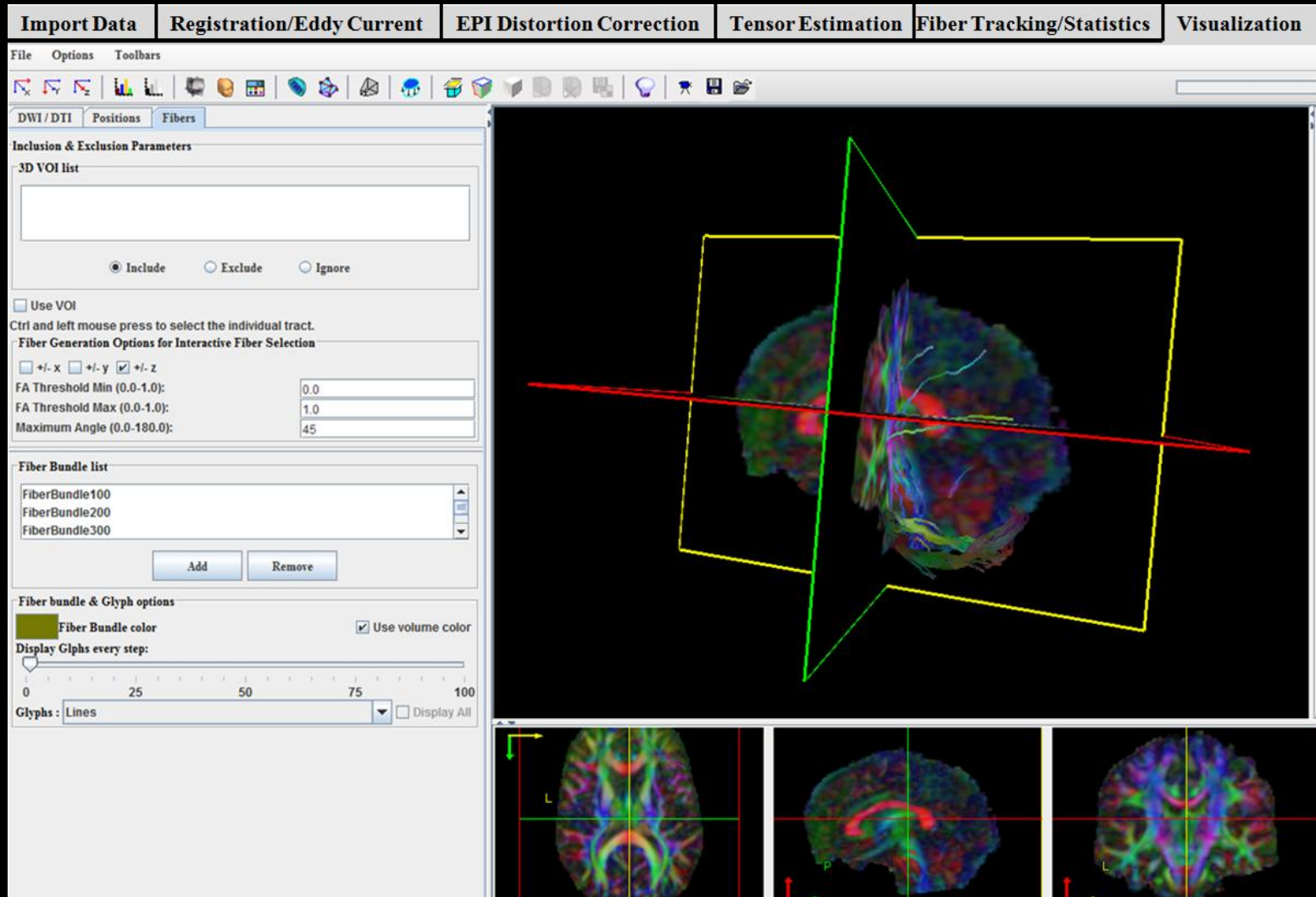
Buttons: Load B-Value/Grad File, Save Table As, Edit Table, Clear, +/- x, +/- y, +/- z, Apply, Close

Current MIPAV DTI Pipeline



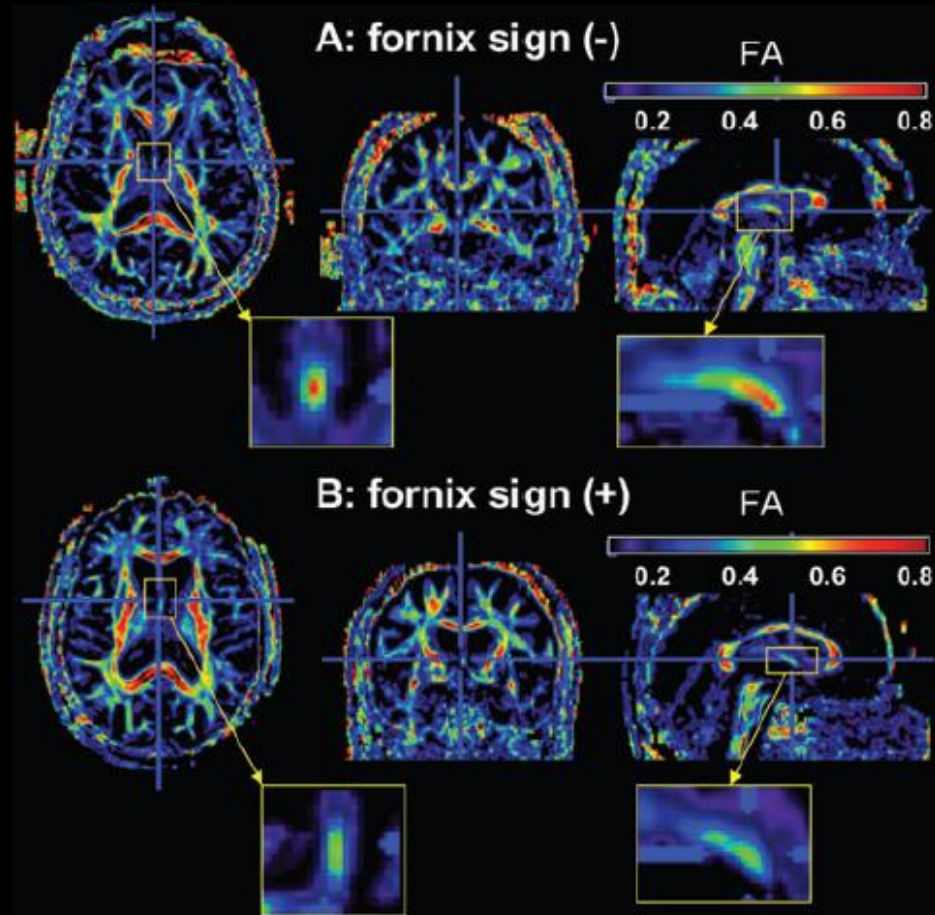
Next Step

Create an integrated DTI dialog incorporating all the steps of the pipeline including pre-processing and post-processing steps



The Fornix Sign: A Potential Sign for Alzheimer's Disease Based on Diffusion Tensor Imaging.

Oishi K, Mielke MM, Albert M, Lyetsos CG, Mori, S (2011) J Neuroimaging. 1552-6569



© 2011 by the American Society of Neuroimaging.

Fig 1. Example of the fornix sign. The axial (left), coronal (middle), and sagittal (right) slices of the color-scaled FA map are shown with the magnified view of the fornix (yellow rectangle). (A) FA map of a cognitively normal 80-year-old woman without a fornix sign. The core part of the fornix appears yellow to red (FA .5-.8). (B) FA map of an 80-year-old woman with Alzheimer's disease with the fornix sign. The fornix appears green (FA < .5). FA = fractional anisotropy.

References

<http://mipav.cit.nih.gov/>

Oishi K, Mielke MM, Albert M, Lyetsos CG, Mori S (2011) The Fornix Sign: A Potential Sign for Alzheimer's Disease Based on Diffusion Tensor Imaging. J Neuroimaging. 1552-6569

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