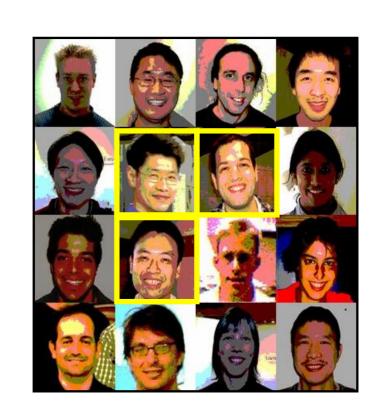


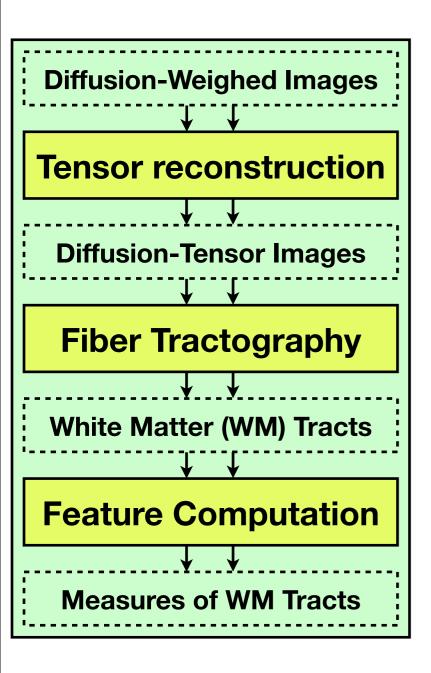
Data interoperability of DTI-TK for DTI analysis & A preview of ITK-SNAP 2.0

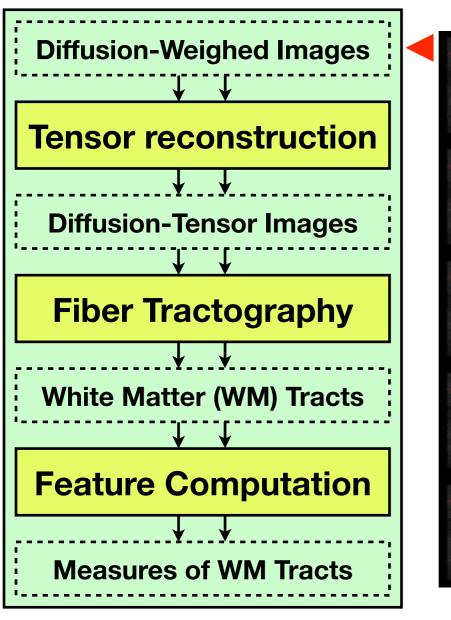
Gary Hui Zhang, Paul A Yushkevich, and James C Gee

Penn Image Computing & Science Laboratory (PICSL) Department of Radiology, University of Pennsylvania

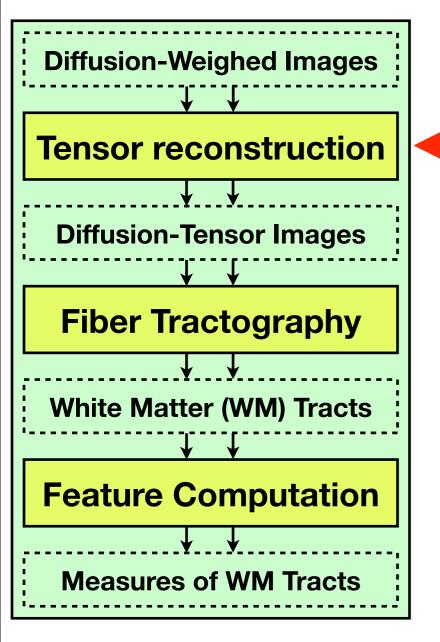


	Copyright © 2009 Gary Hui Zhang
Data interoperability of DTI-TK for DTI	analysis

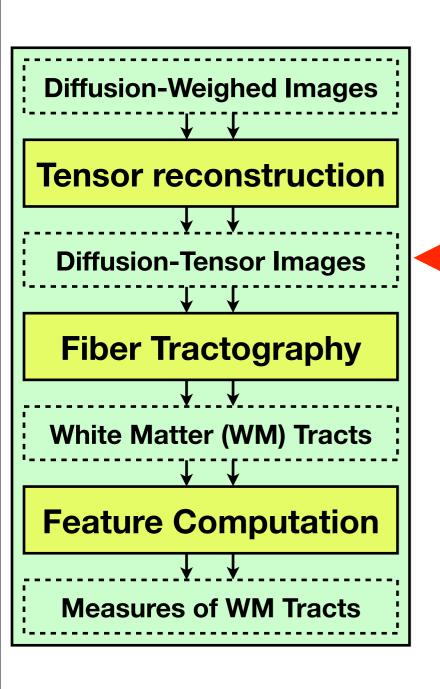


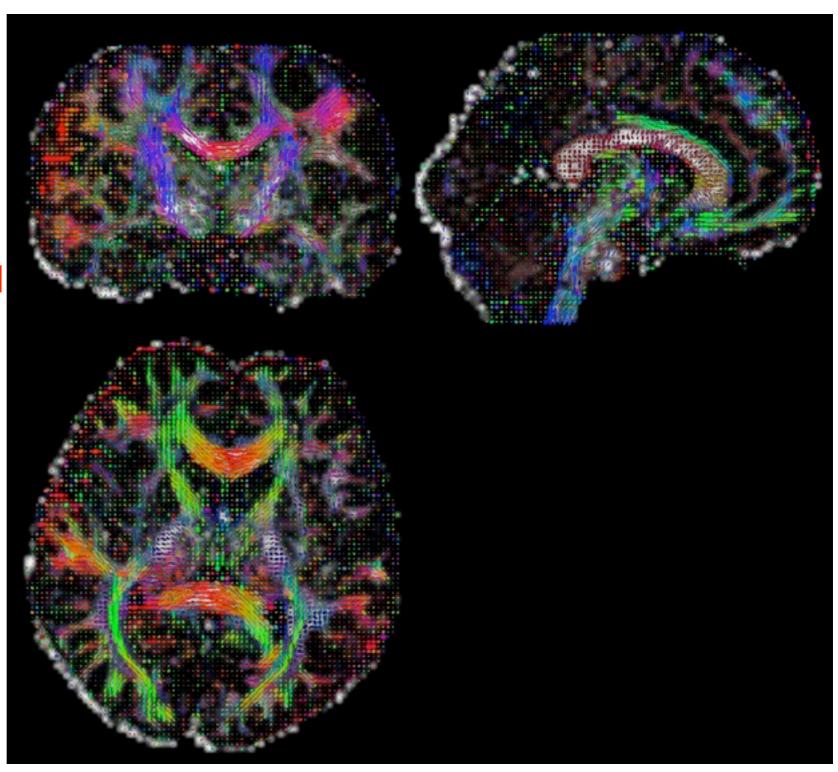


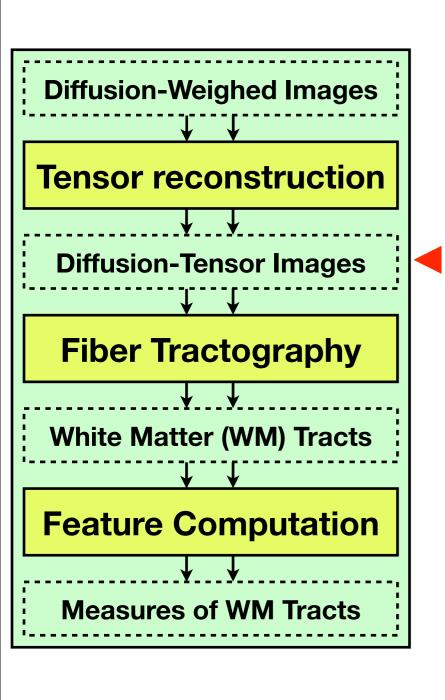


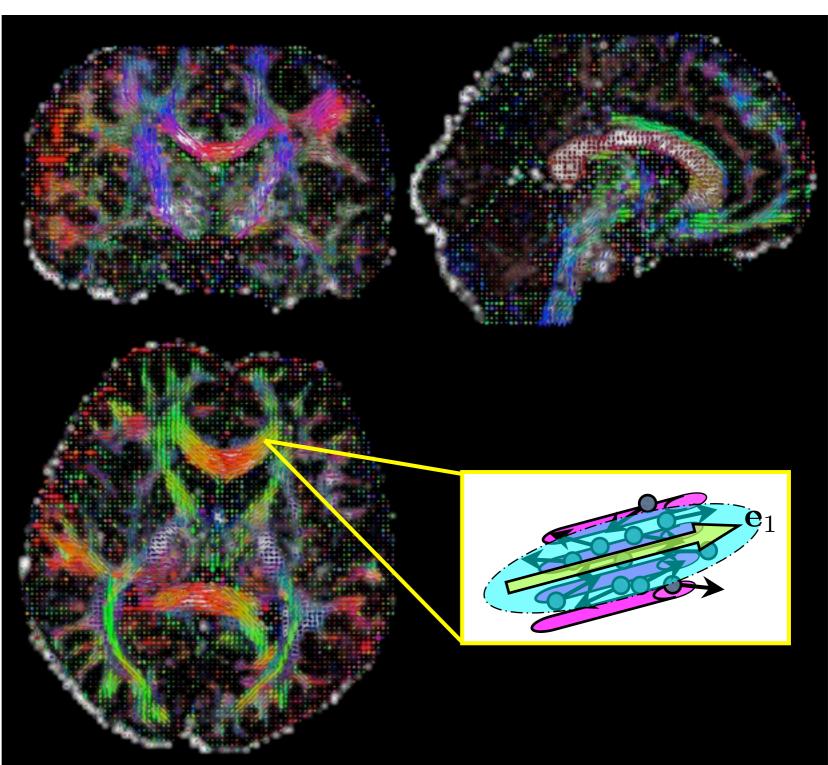


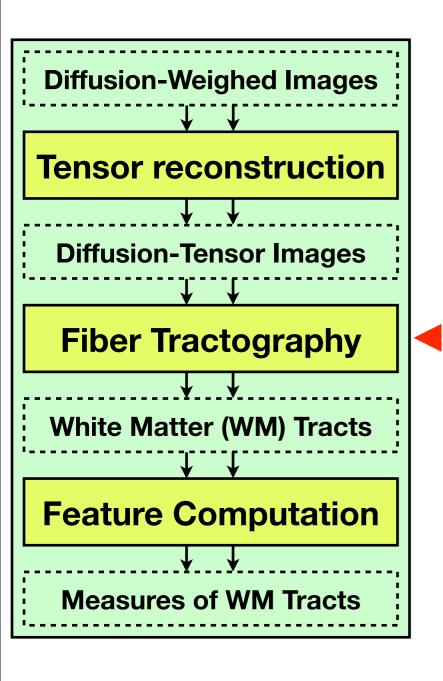


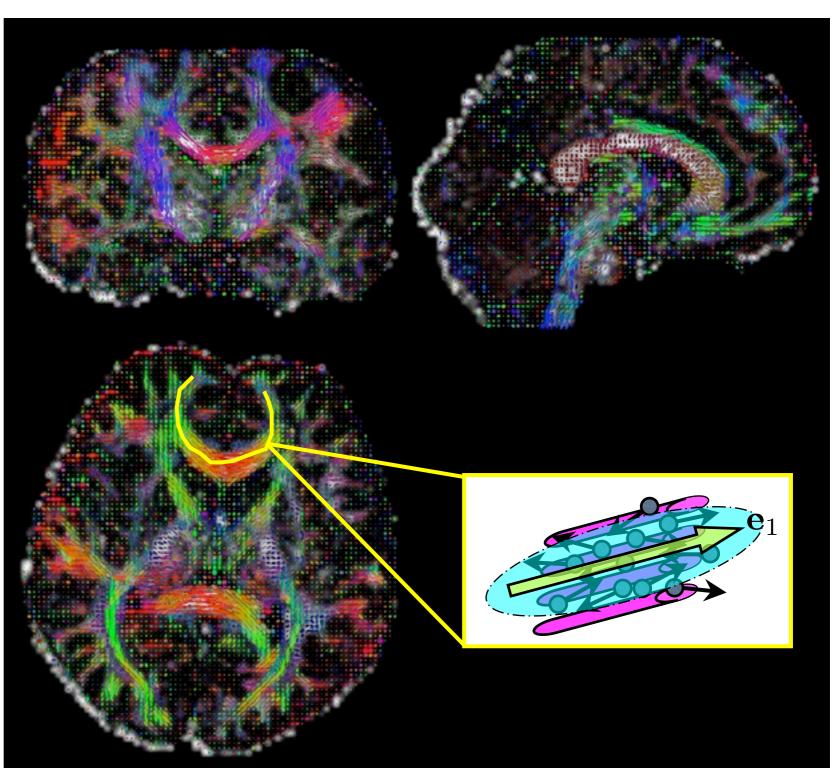


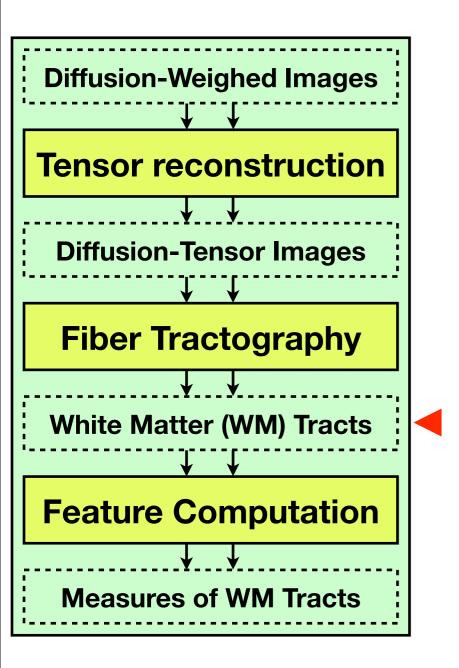


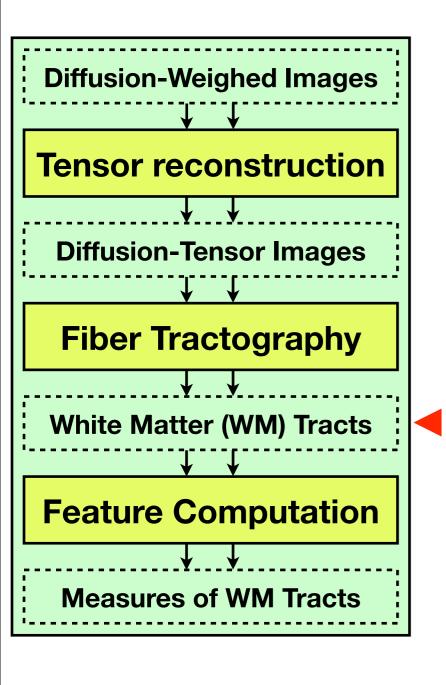


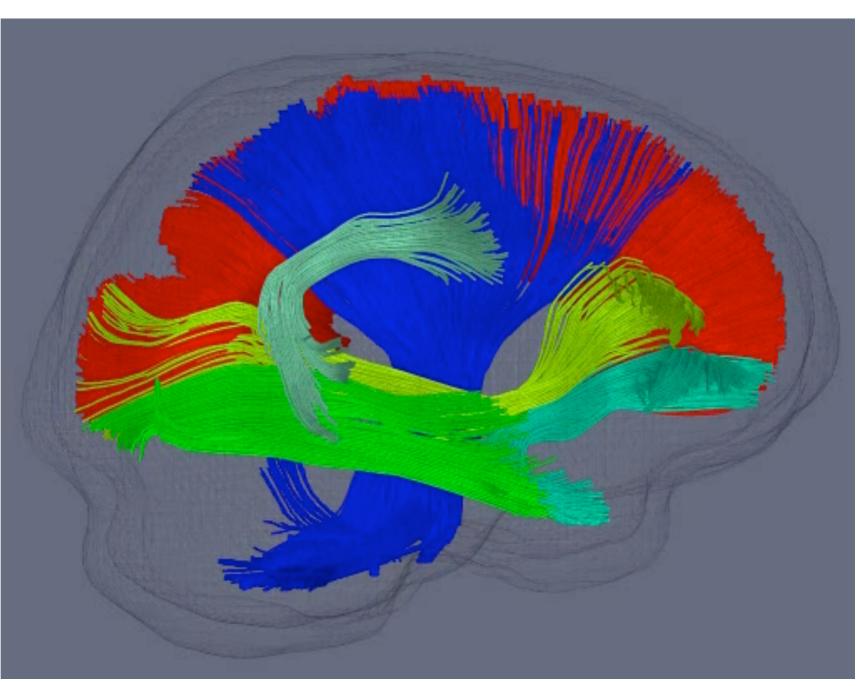


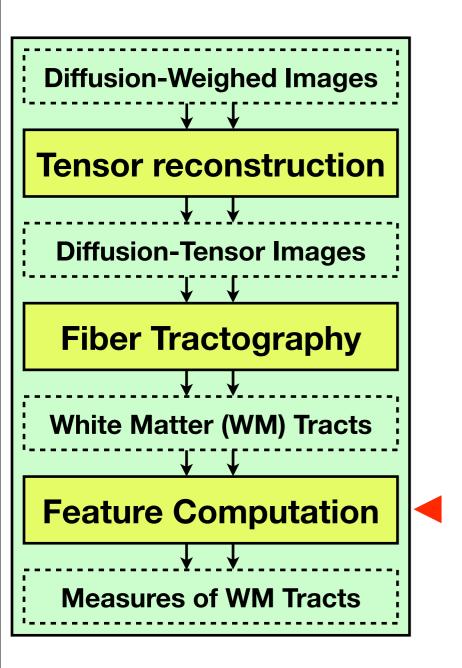


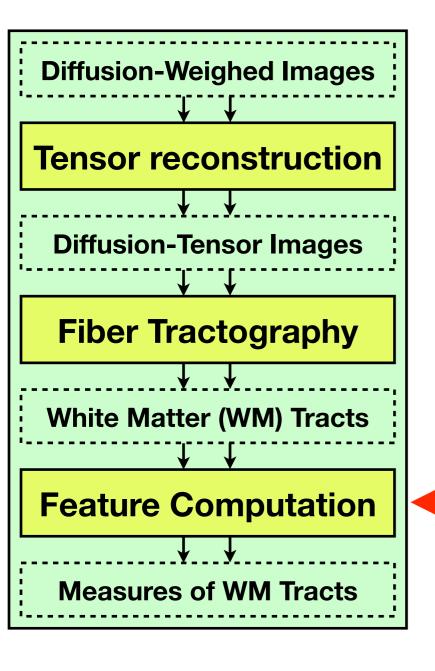


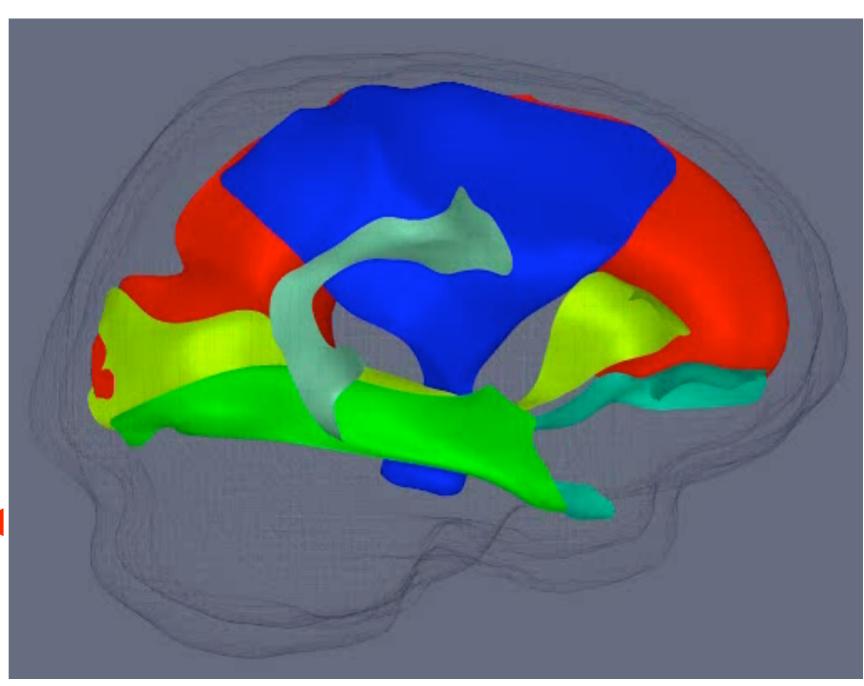


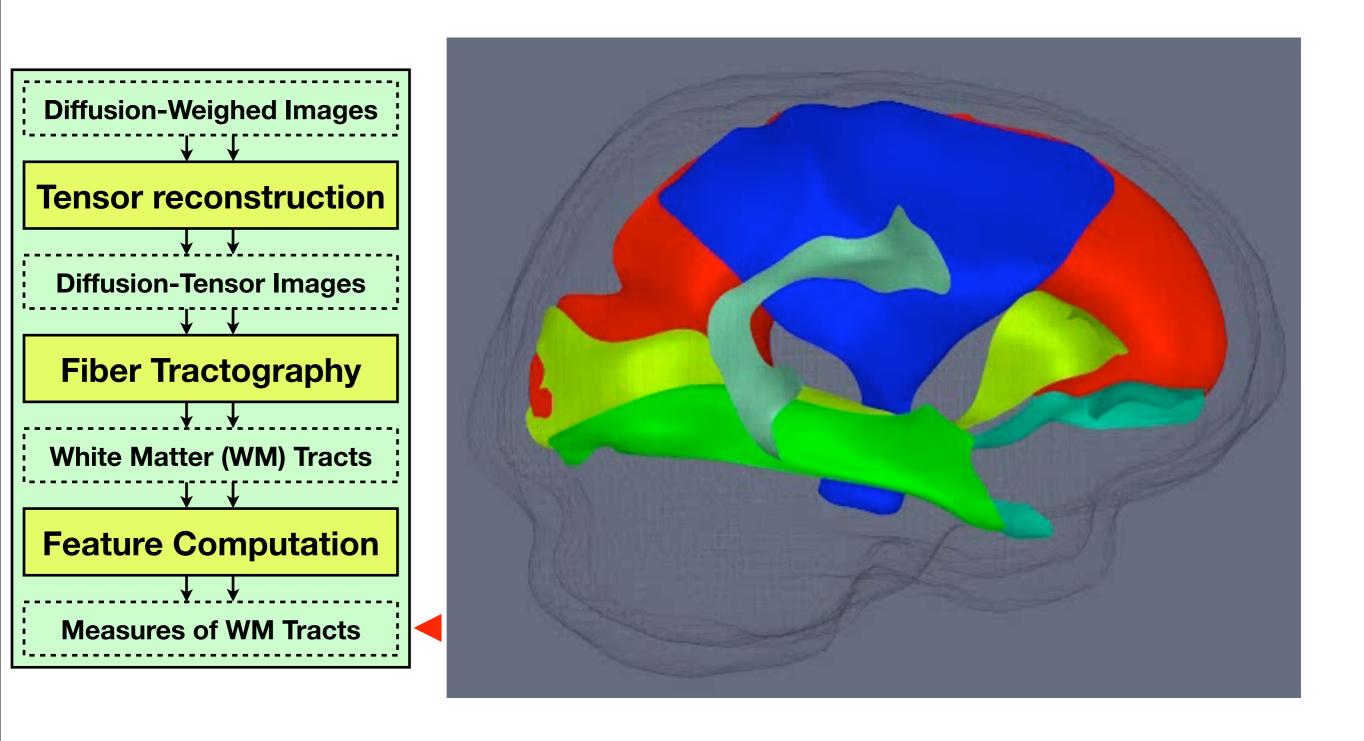




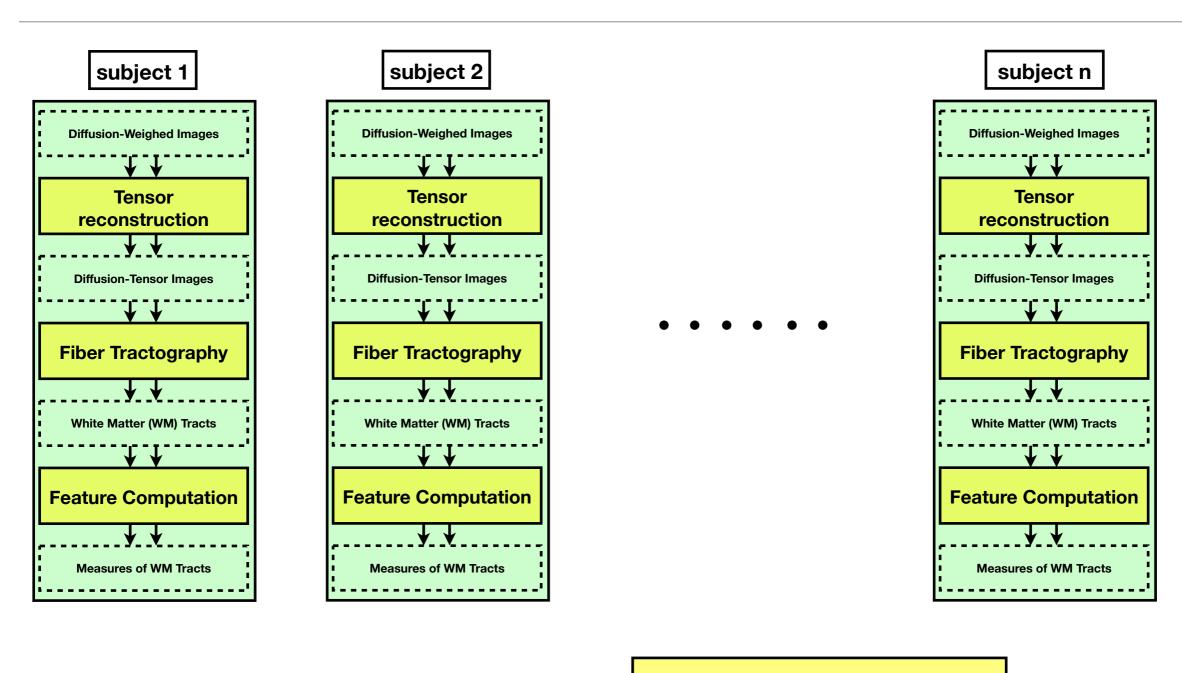








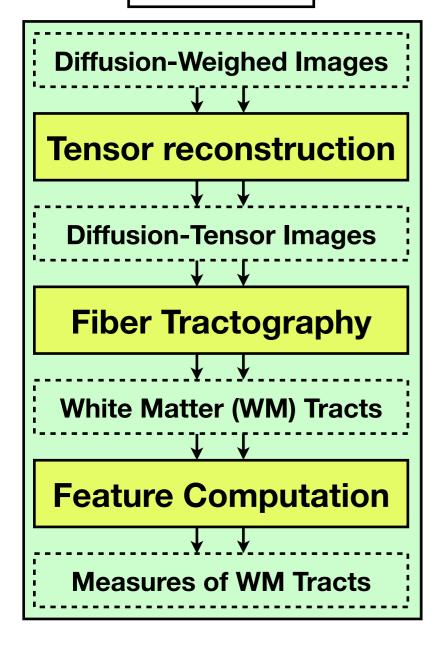
A typical DTI analysis pipeline for group studies



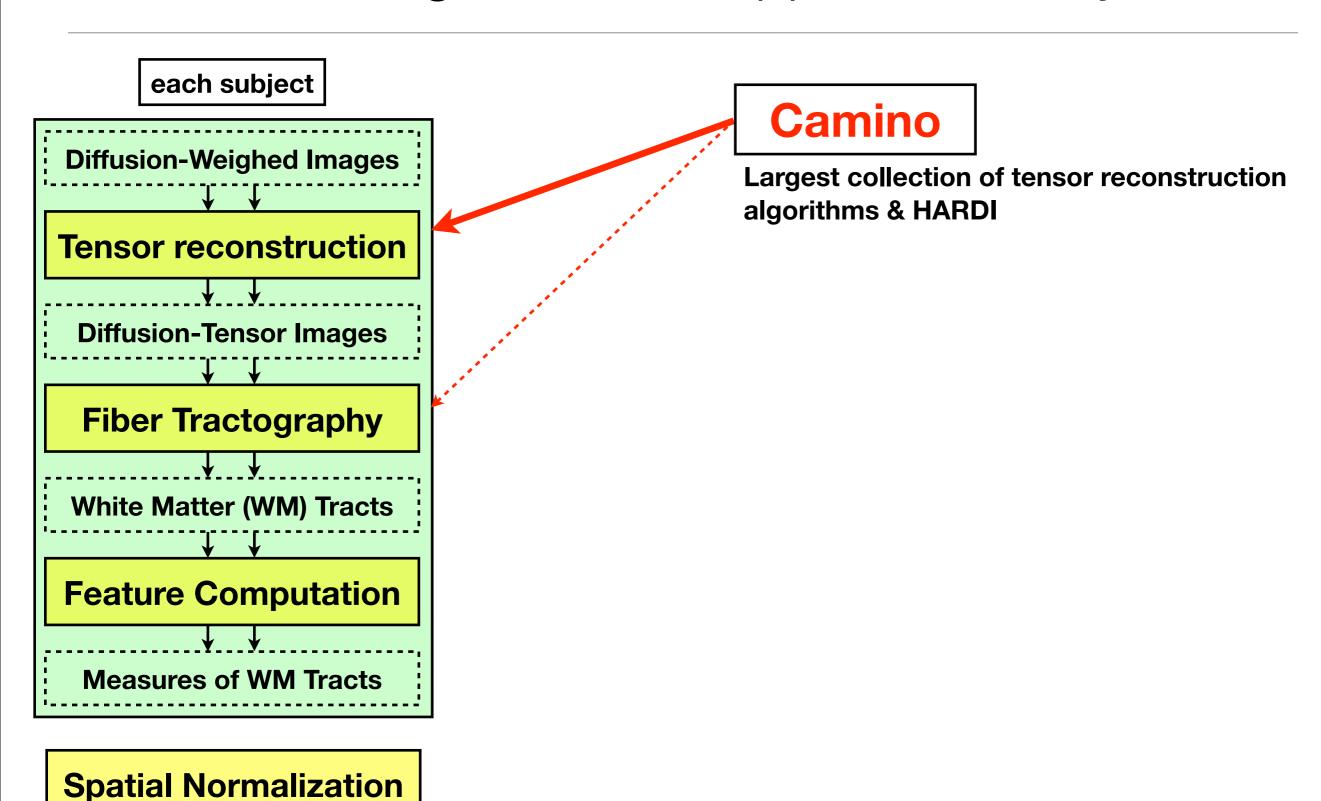
Population studies

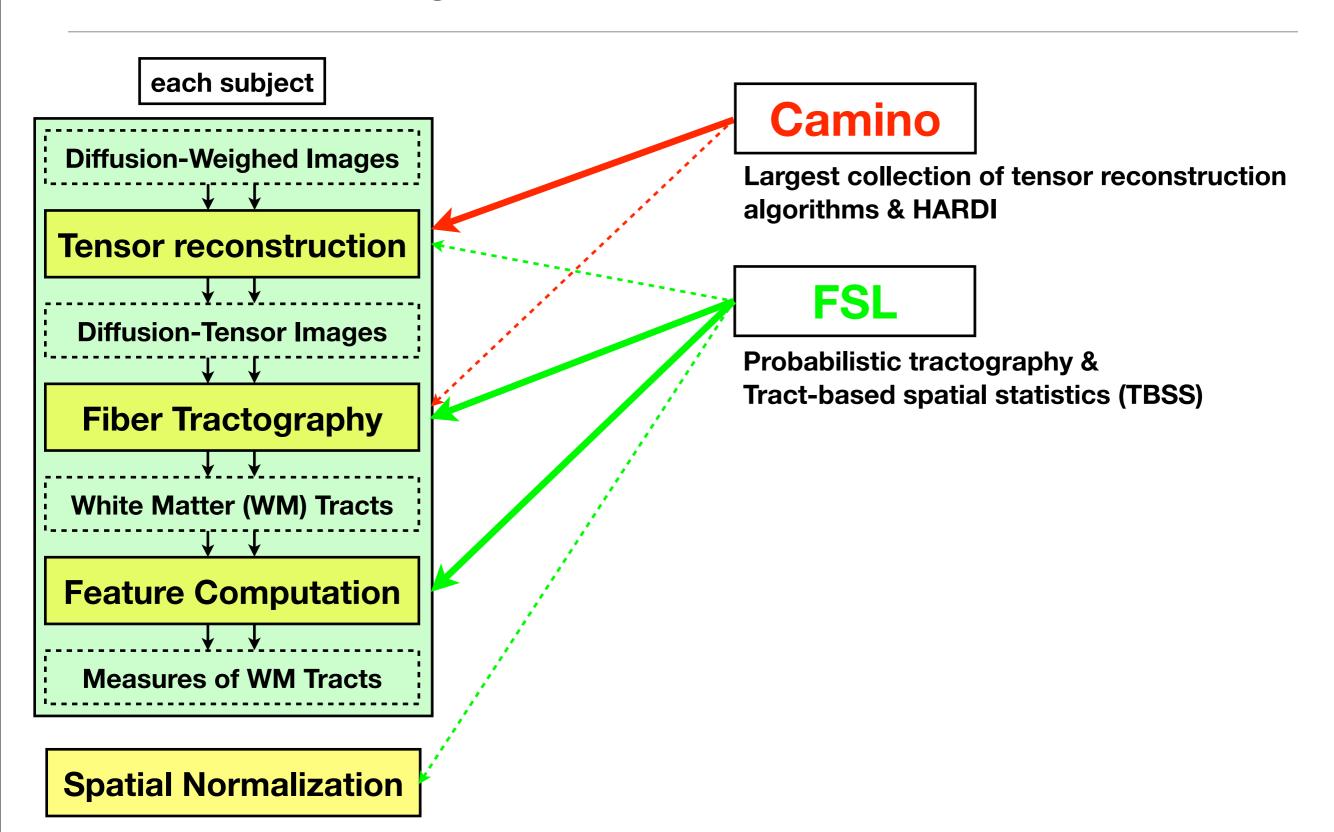
Spatial Normalization

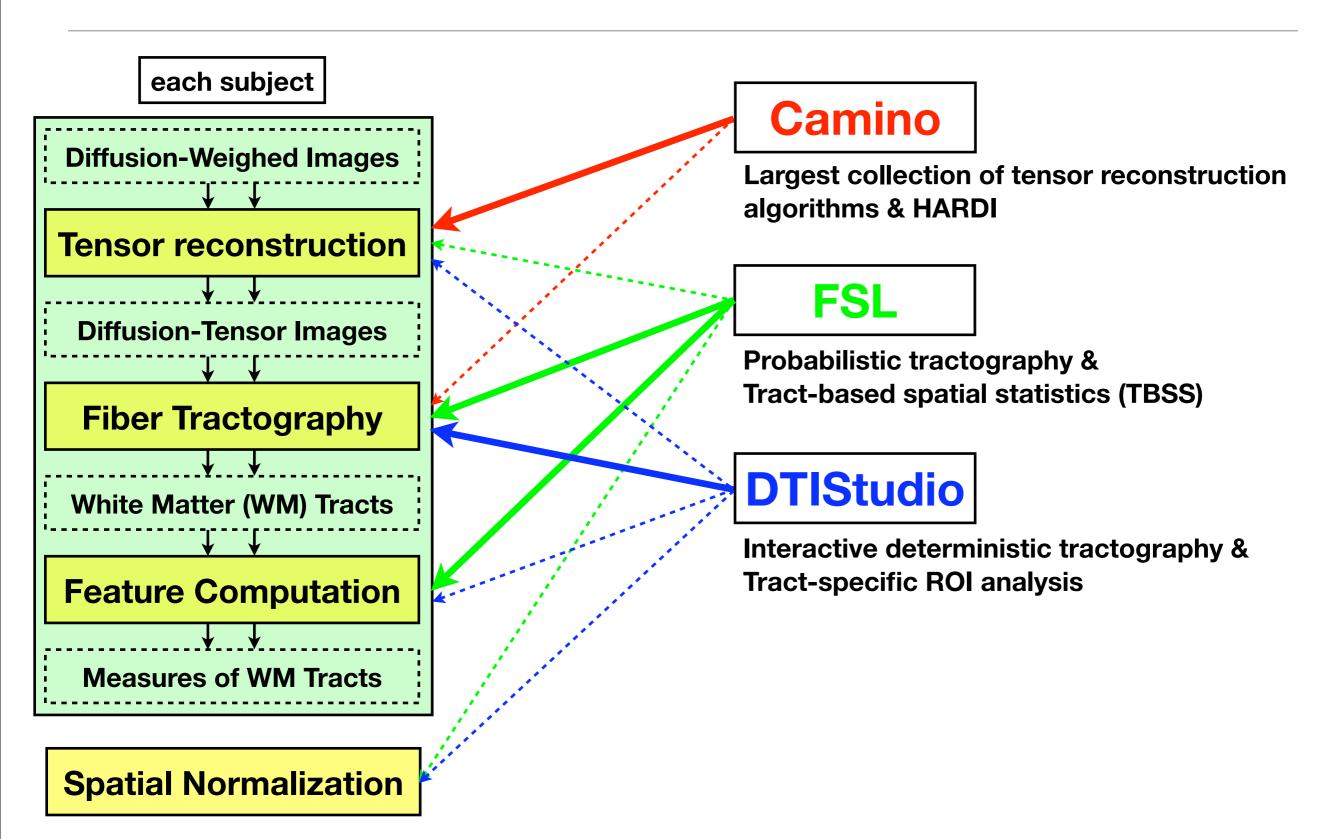
each subject

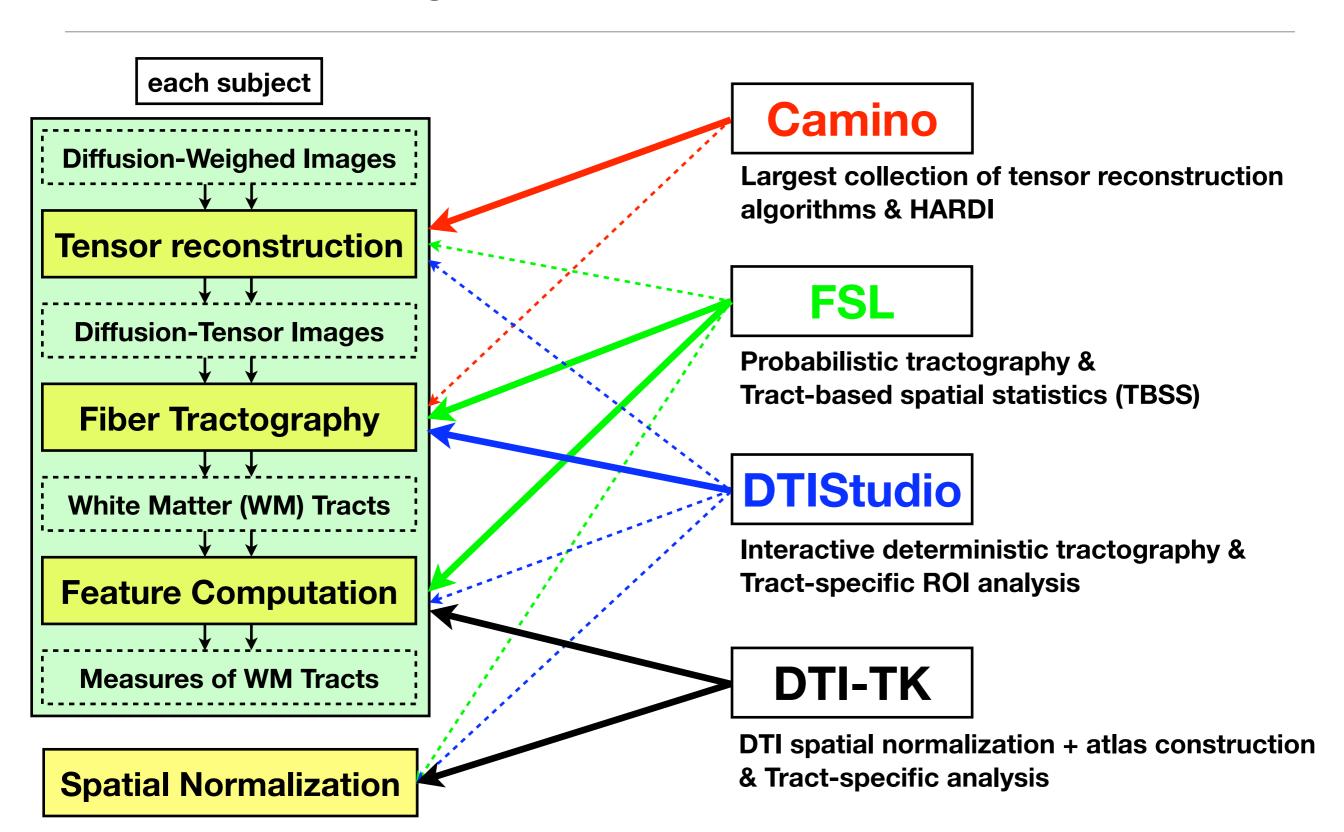


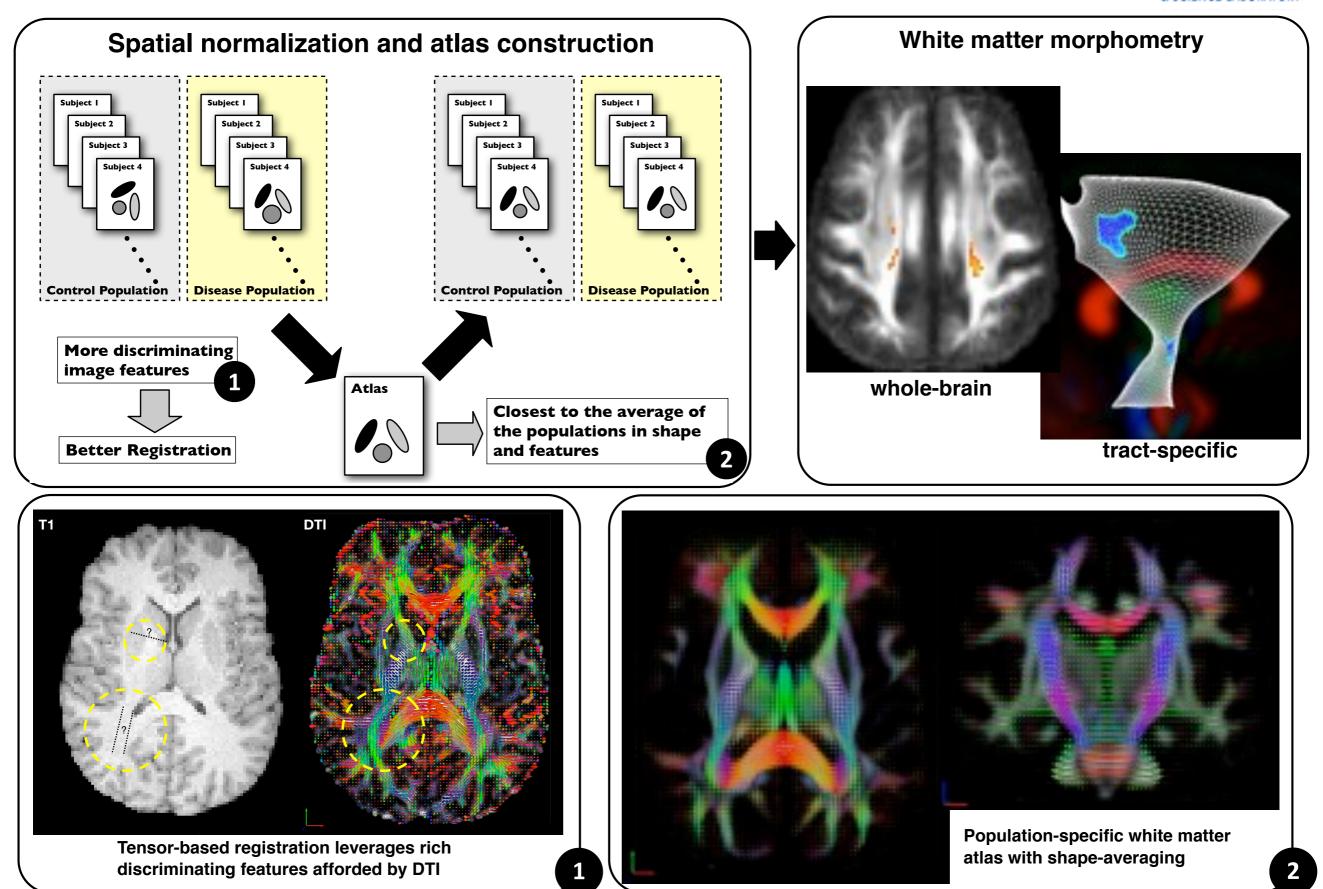
Spatial Normalization









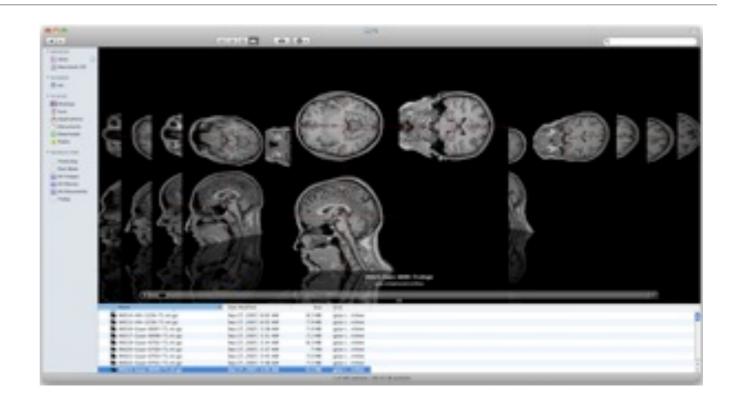


For download, visit http://www.nitrc.org/projects/dtitk

DTI-TK Quick Look Plugin for Mac OSX

About DTI-TK Quick Look Plugin

This plugin uses the Mac OS X Leopard's built-in innovative Quick Look framework to enable a quick assessment of any 3-dimensional image volume in the supported medical image formats (NIfTI / Analyze / FreeSurfer) directly from the Finder. Using the Finder's Cover Flow mode, a large collection of medical images can be browsed through and quickly inspected just as easy as flipping through your photos. Furthermore, a large number of images can be compared side-by-side by selecting them together in the Finder, then pressing the space bar to bring up the Quick Look preview mode.



System Requirements

Mac OS X 10.5 or later

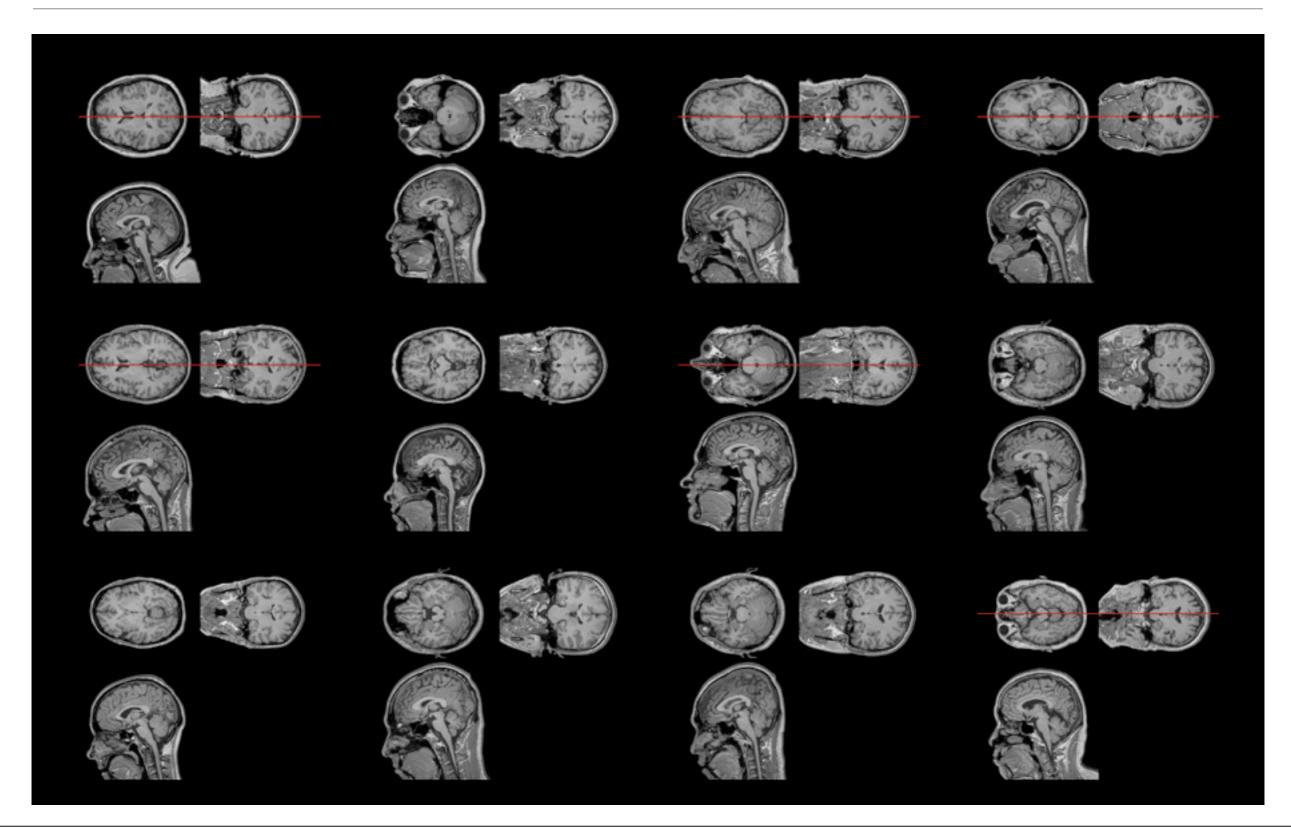
Download Details

Company:	university of pennsylvania
Version:	1.7.3
Post Date:	June 8, 2009

DTI-TK Quick Look Plugin: Cover Flow Mode



DTI-TK Quick Look Plugin: Multi-Volume Preview



$$\mathbf{D} = \begin{pmatrix} D_{xx} & D_{yx} & D_{zx} \\ D_{yx} & D_{yy} & D_{zy} \\ D_{zx} & D_{zy} & D_{zz} \end{pmatrix}$$
$$= \lambda_1 \mathbf{e}_1 \mathbf{e}_1^\mathsf{T} + \lambda_2 \mathbf{e}_2 \mathbf{e}_2^\mathsf{T} + \lambda_3 \mathbf{e}_3 \mathbf{e}_3^\mathsf{T}$$

$$\mathbf{D} = \begin{pmatrix} D_{xx} & D_{yx} & D_{zx} \\ D_{yx} & D_{yy} & D_{zy} \\ D_{zx} & D_{zy} & D_{zz} \end{pmatrix} \qquad \qquad \qquad \text{matrix representation}$$

$$= \lambda_1 \mathbf{e}_1 \mathbf{e}_1^\mathsf{T} + \lambda_2 \mathbf{e}_2 \mathbf{e}_2^\mathsf{T} + \lambda_3 \mathbf{e}_3 \mathbf{e}_3^\mathsf{T} \qquad \qquad \qquad \text{eigensystem representation}$$

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$$\begin{aligned} (D_{xx}, D_{yx}, D_{yy}, D_{zx}, D_{zy}, D_{zz}) \\ \text{lower triangular} \end{aligned}$$

This is the NIfTI Tensor standard

DTI-TK

$$(D_{xx}, D_{yx}, D_{yy}, D_{zx}, D_{zy}, D_{zz})$$
 lower triangular

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Camino

$$D_{xx}, D_{yy}, D_{zz}, D_{yx}, D_{zx}, D_{zy}$$
 individual NIfTI scalar files

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DTI-TK

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This is the NIfTI Tensor standard

Camino

$$D_{xx}, D_{yy}, D_{zz}, D_{yx}, D_{zx}, D_{zy}$$
 individual NIfTI scalar files



$$\lambda_1,\lambda_2,\lambda_3,e_1,e_2,e_3$$
 individual NIfTI scalar or vector files

$$\begin{array}{c} \mathbf{Or} & (D_{xx}, D_{yx}, D_{zx}, D_{yy}, D_{zy}, D_{zz}) \\ \mathbf{upper \ triangular} \end{array}$$

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$$D_{xx}, D_{yy}, D_{zz}, D_{yx}, D_{zx}, D_{zy}$$

individual Analyze files for export

 FA, \mathbf{e}_1

individual raw binary files for import

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 - fully implemented, tested, and documented

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 - New techniques can be more easily compared to existing tools

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A preview of ITK-SNAP 2.0

• ITK-SNAP is a software application used to <u>segment structures in 3D</u> <u>medical images</u>. It provides <u>semi-automatic segmentation</u> using active contour methods, as well as <u>manual delineation</u> and <u>image navigation</u>.

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 - Enhanced support for manual segmentation via the new adaptive brush
 - User experience enhancement, including redesigned user interface, native file chooser, and automatic check for update

ITK-SNAP 2.0 Preview Demo

For download, visit http://www.itksnap.org

Acknowledgement

- DTI-TK
 - NIH grant: R03 EB009321
 - DTI-TK user community

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 - NIH grant: R03 EB008200
 - Prof. Guido Gerig, original SNAP developers at UNC
 - ITK-SNAP user community

Thank you for your attention.