Open-source diffusion MRI for cancer research

Lauren J. O'Donnell, Ph.D.

Assistant Professor of Radiology

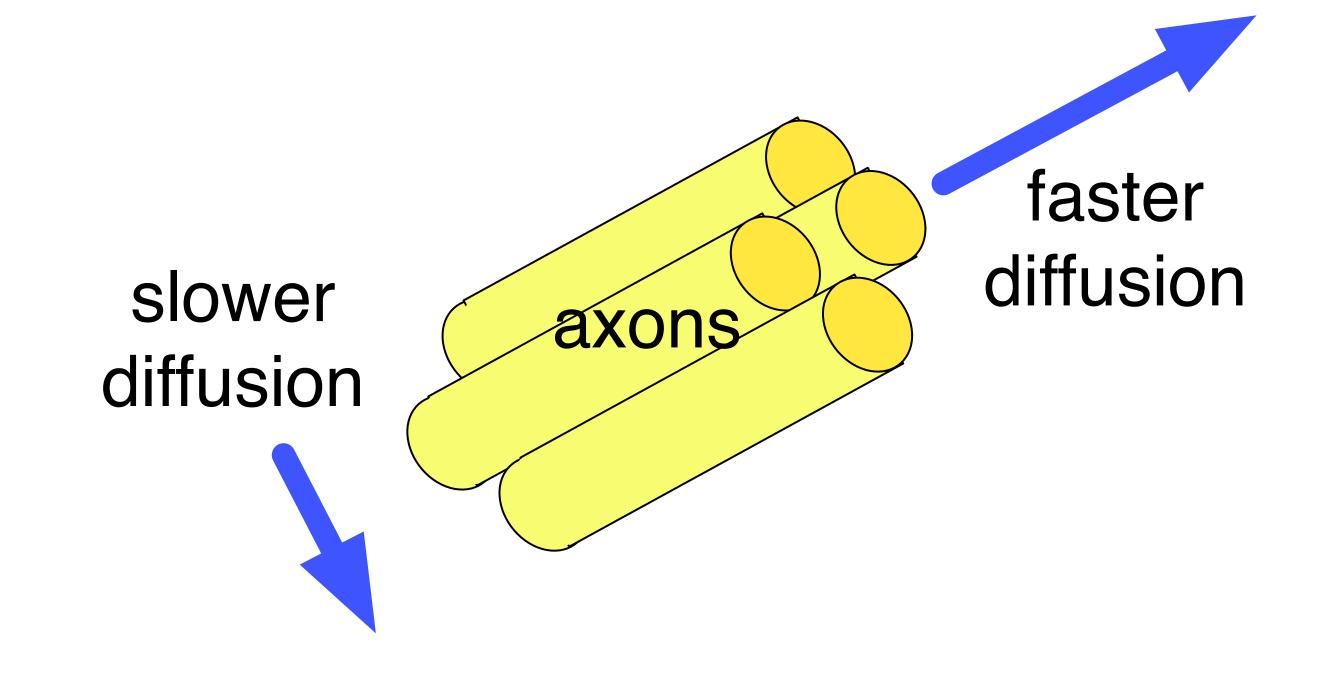
Harvard Medical School and Brigham and Women's Hospital

Informatics Technology for Cancer Research (ITCR) PI Meeting

June 2016

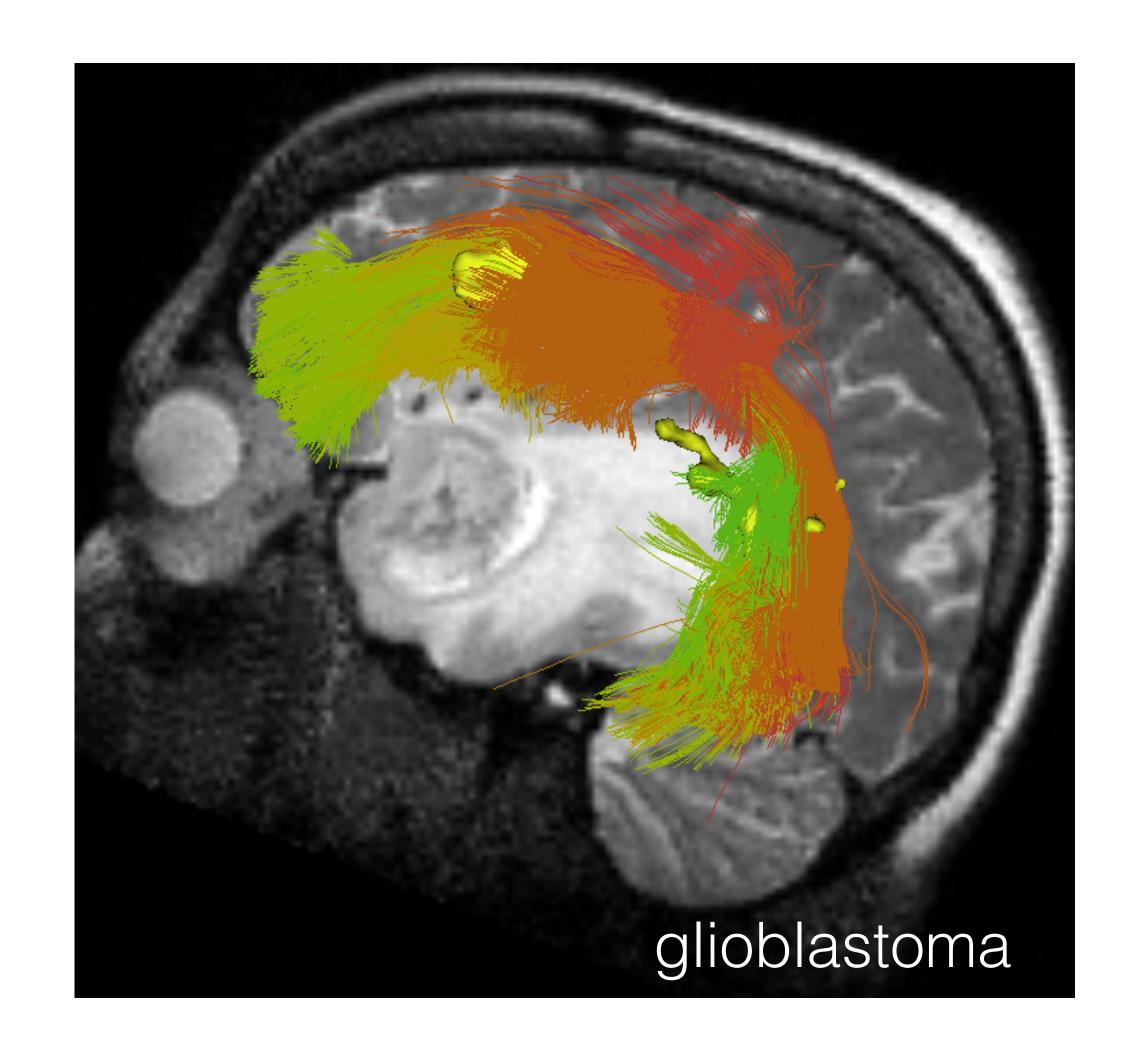
Diffusion MRI (dMRI)

- Only non-invasive probe of brain's microstructure
- Measure water diffusion in many orientations using MRI
- Map brain connections
- Tractography



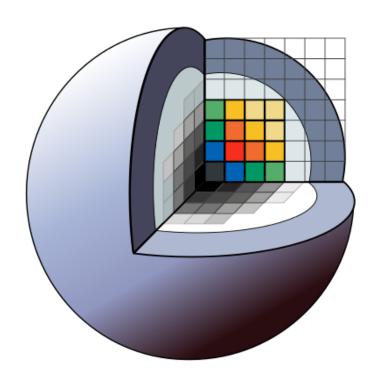
Neurosurgery for Brain Tumors

- Maximal tumor resection improves patient outcome
- dMRI: Map white matter tracts
- dMRI: Map brain microstructure
- dMRI is used for surgical planning and neuronavigation during surgery.



SlicerDMRI U01

- Create state-of-the-art dMRI workflows for brain cancer research.
- 3D Slicer: Open-source platform for visualization, surgical planning, surgical navigation, and medical image computing
- Latest improvements are available:
 - www.slicer.org
 - https://github.com/SlicerDMRI



Year 1: DICOM Tractography

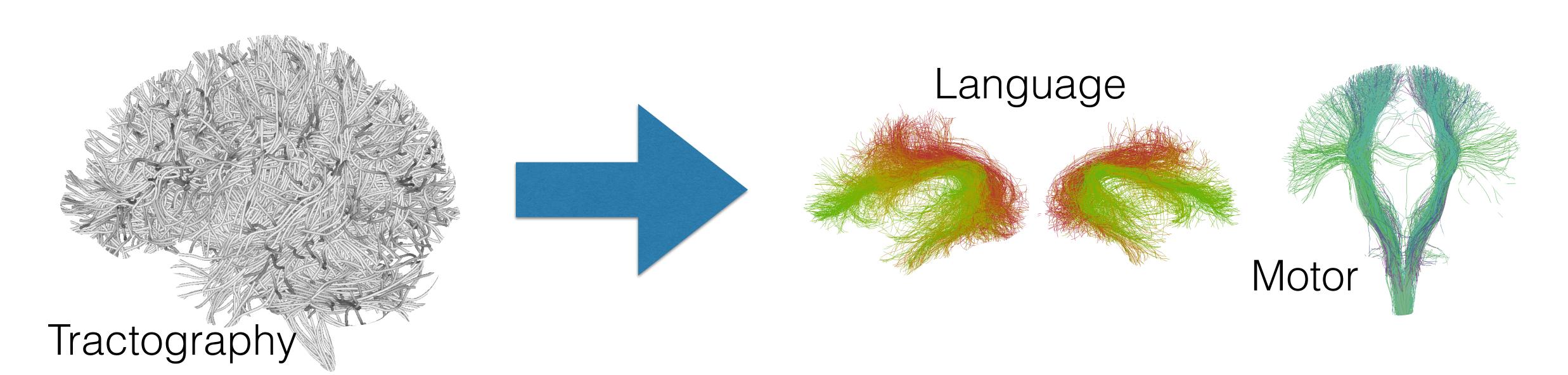
- First implementation of the standard
- DCMTK: Open-source DICOM library
- Slicer DICOM I/O module
- DICOM <-> major research formats
- Future interoperability with commercial neurosurgical neuronavigation!

Table C.8.X-2 TRACTOGRAPHY RESULTS MODULE ATTRIBUTES

Attribute Name	Tag	Туре	Attribute Description
Include Table 10-12 "Content Identification	n Macro Attributes"		
Content Date	(0008,0023)	1/	The date the content creation started.
Content Time	(0008,0033)	1	The time the content creation started.
Track Set Sequence	(0066,0101)	1	Describes the track sets that are contained within the data.
			One or more Items shall be included in this sequence.
>Track Set Number	(0066,0105)	1	Identification number of the Track Set. Uniquely identifies a track set within this SOP instance. Shall start at a value of 1, and increase monotonically by 1.

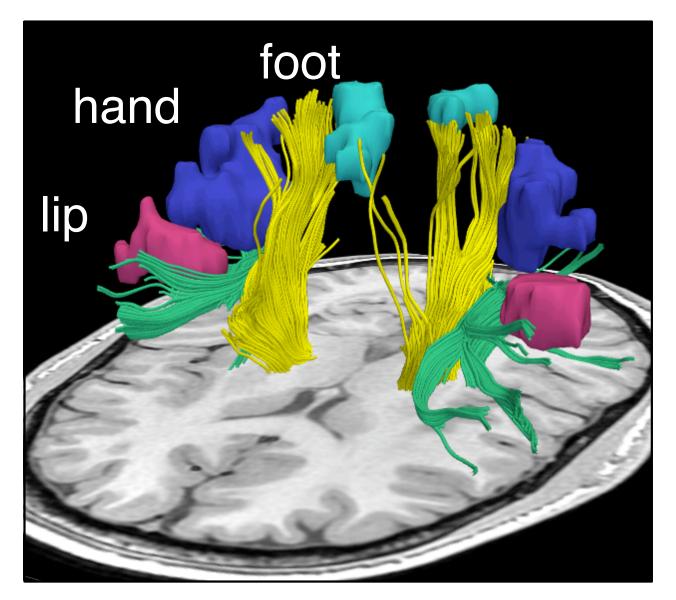
Research: Tractography for Neurosurgery

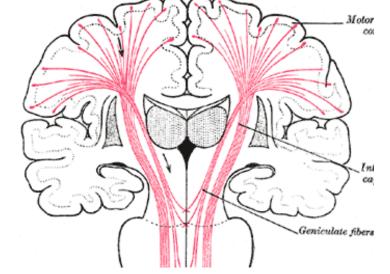
- Tractography challenges: crossing fibers, edema, displacement
- Analysis challenge: identify key tracts from 100,000's of fibers



Challenge: Crossing Fibers

- Clinical method: Diffusion tensor (DTI)
- Models one tract per voxel
- Multi-fiber models needed [1] [2]



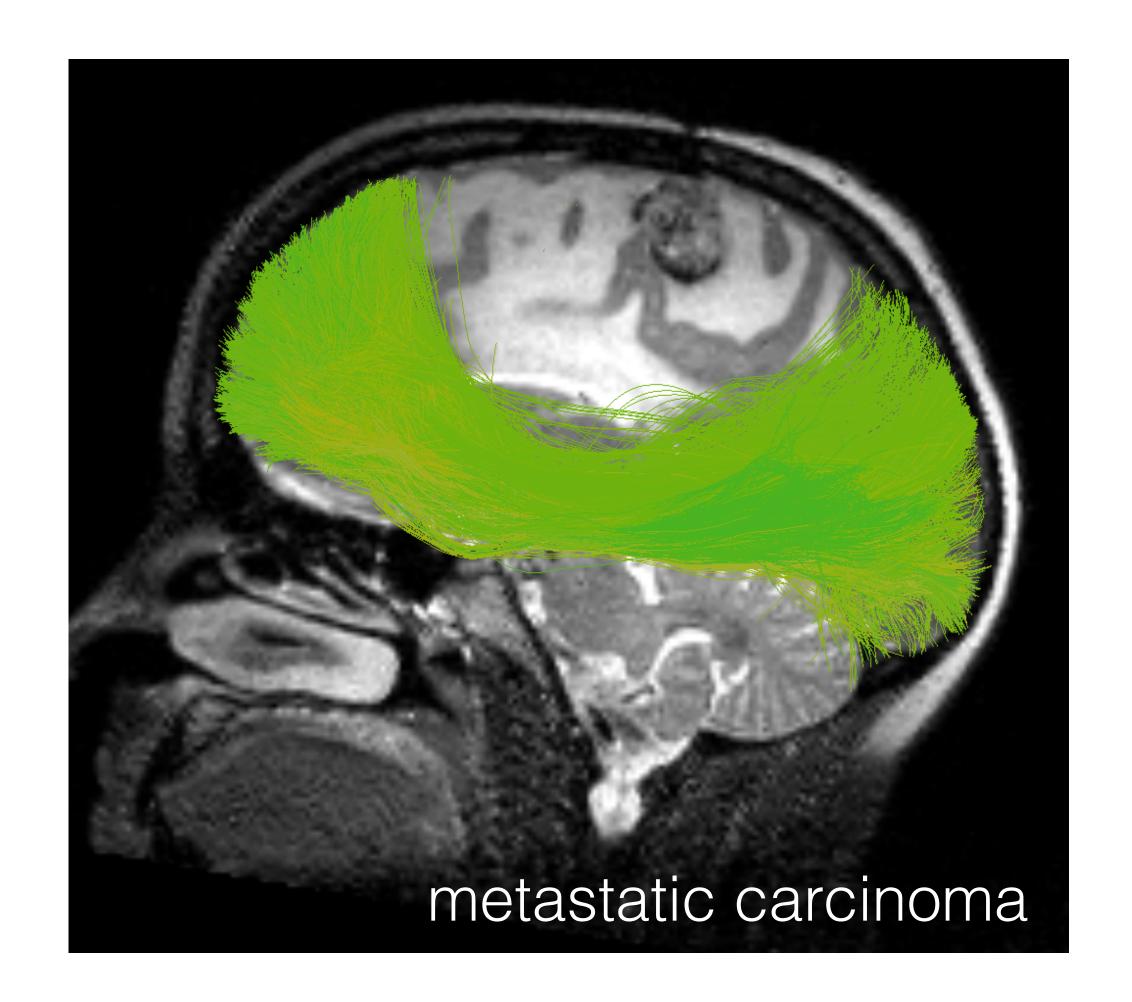


false negative

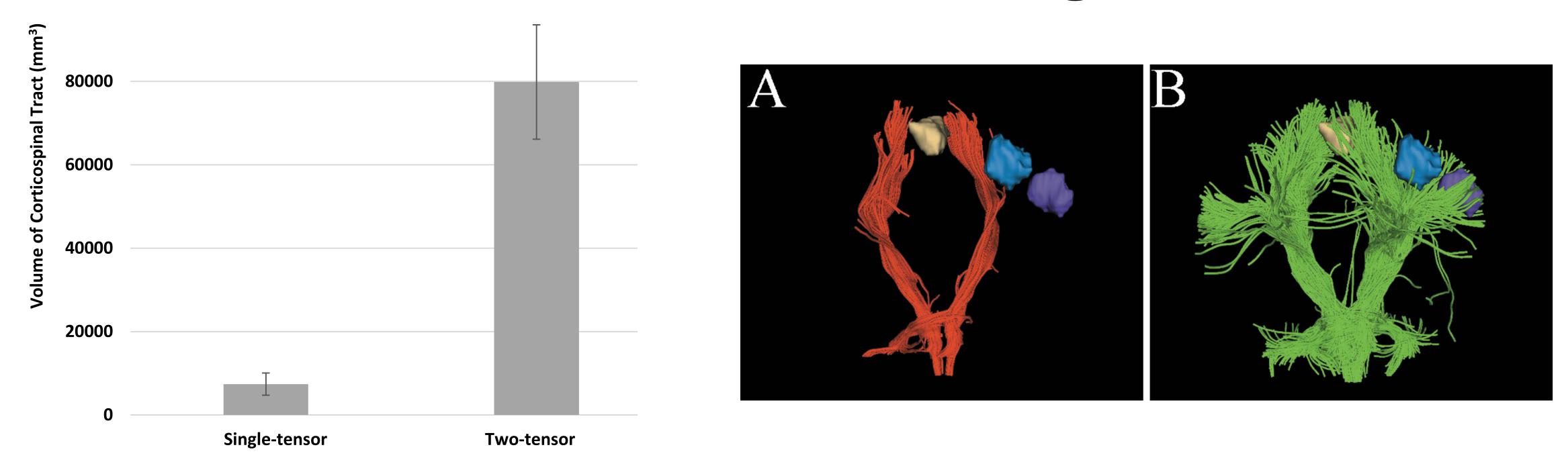
- [1] A. A. Qazi, A. Radmanesh, L. O'Donnell, G. Kindlmann, S. Peled, S. Whalen, C.-F. Westin, A.J. Golby, Resolving crossings in the corticospinal tract by two-tensor streamline tractography: method and clinical assessment using fMRI, Neurolmage 47 (2009) T98–T106
- [2] C. Nimsky, Fiber tracking we should move beyond diffusion tensor imaging, World neurosurgery 2014

Challenge: Edema

- Peritumoral edema increases water around tumor
- Changes dMRI signal
- Increases difficulty of tractography



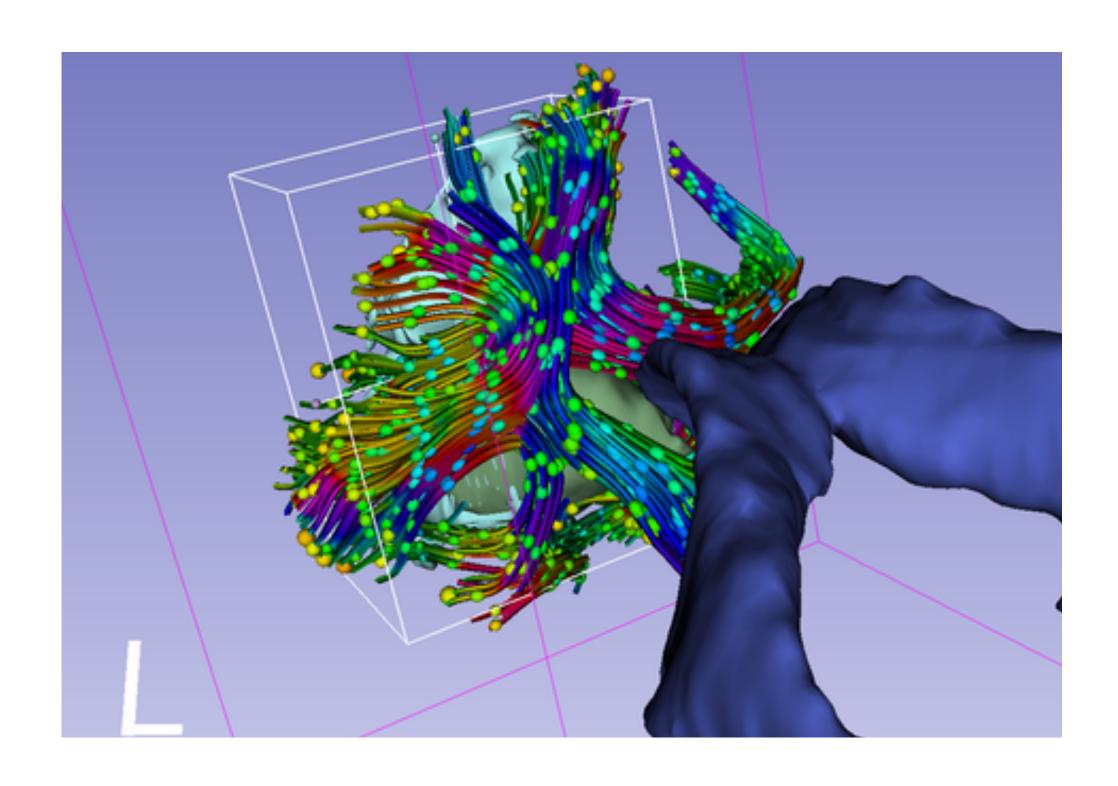
Robustness to crossings, edema



[3] Chen, Zhenrui, et al. "Corticospinal tract modeling for neurosurgical planning by tracking through regions of peritumoral edema and crossing fibers using two-tensor unscented Kalman filter tractography." International journal of computer assisted radiology and surgery 2016
[4] Chen, Zhenrui, et al. "Reconstruction of the arcuate fasciculus for surgical planning in the setting of peritumoral edema using two-tensor unscented Kalman filter tractography." NeuroImage: Clinical 2015

Challenge: Identification of Important Tracts

- Brain is highly connected: 100,000's of "fibers"
- Current clinical method: interactive selection of fibers
- Time consuming, operatordependent, and variable [5] [6]



- [5] U. Burgel et al, Fiber tracking with distinct software tools results in a clear diversity in anatomical fiber tract portrayal., Central European Neurosurgery 2009
- [6] A. Radmanesh et al. Comparison of seeding methods for visualization of the corticospinal tracts using single tensor tractography, Clinical neurology and neurosurgery 2015.

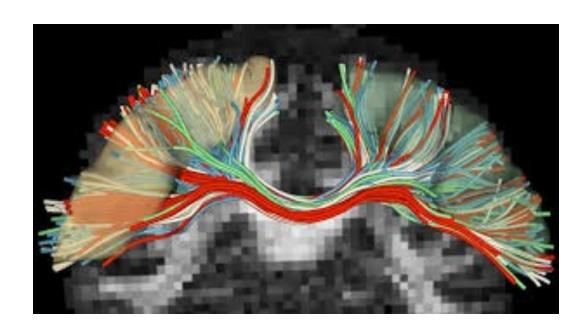
Open-source tools for dMRI







tractography visualization, dMRI quantification, anatomical hierarchies



github.com/pnlbwh/ukftractography

multi-fiber tractography



github.com/SlicerDMRI/whitematteranalysis

tract clustering and registration

Thank You

- Alexandra J. Golby, NCIGT
- Isaiah Norton, software engineer U01
- Yannick Suter, visiting masters student
- Fan Zhang, PhD student
- Laura Rigolo, Golby Lab
- Pegah Kahaliardabili, MD postdoc
- Yogesh Rathi, UKF tractography
- Carl-Fredrik Westin, NAC
- Ron Kikinis, NAC
- NIH NCI ITCR U01 CA199459



