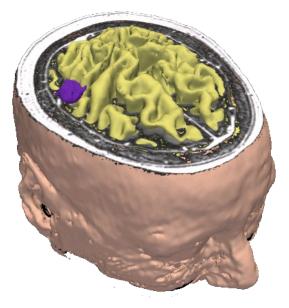


Slicer3 Training Compendium

Measuring Volume Change in Tumors



Kilian M. Pohl, Ph.D.

Ender Konugolu, Ph.D.

Andriy Fedorov, Ph.D.



Material

This course requires the following installation:

3DSlicer version 3.6 Software, which can be installed from:

http://www.slicer.org/pages/Special:SlicerDownloads

A reliable internet connection for downloading the data

Disclaimer

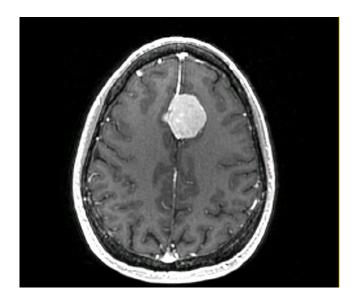
It is the responsibility of the user of 3DSlicer to comply with both the terms of the license and with the applicable laws, regulations and rules.

The module described in this tutorial was tested on Axial 3D SPGR T1 post Gadolinium scans (Voxel dimension: 0.94mm x 0.94mm x 1.20mm, FOV: 240mm, Matrix: 256 x 256)

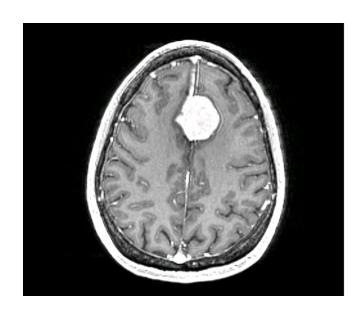




This course is built upon two scans of a patient with meningioma:



MR Scan 1

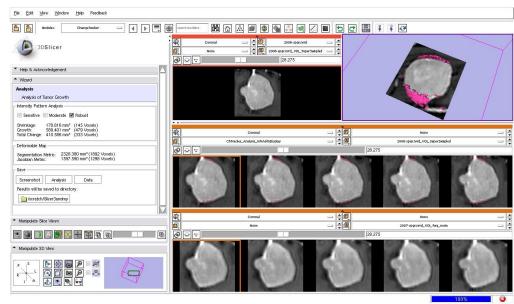


MR Scan 2



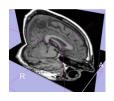
Learning objective

Following this tutorial, you'll be able to load two scans into Slicer3, and measure and visualize the change in volume of the meningioma between the two scans.

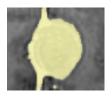




Overview

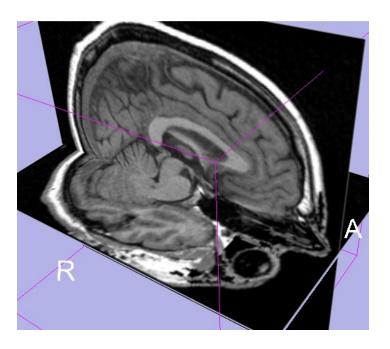


Loading tutorial data



Measuring volume change in tumors





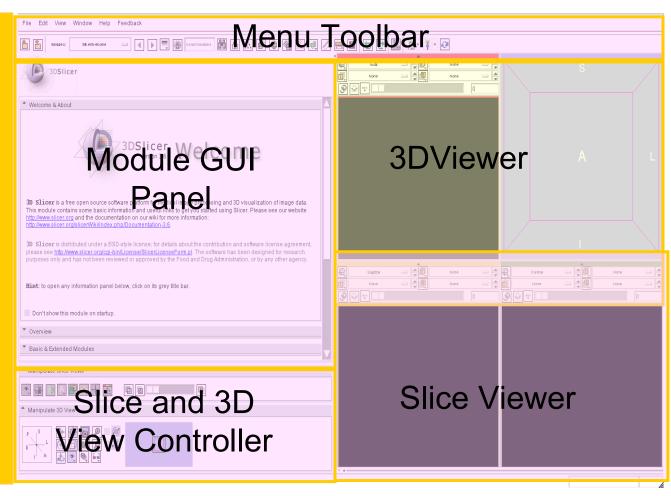
Part 1: Loading the tutorial data



Slicer3 GUI

The Graphical User Interface (GUI) of Slicer3 integrates five components:

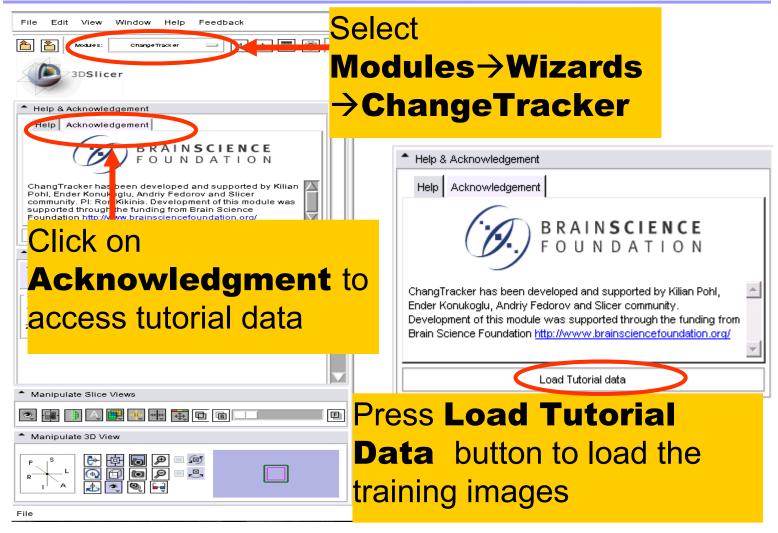
- the Menu Toolbar
- •the Module GUI Panel
- the 3D Viewer
- the Slice Viewer
- •the Slice and 3D View Controller



Slide courtesy of Sonia Pujol, Ph.D.

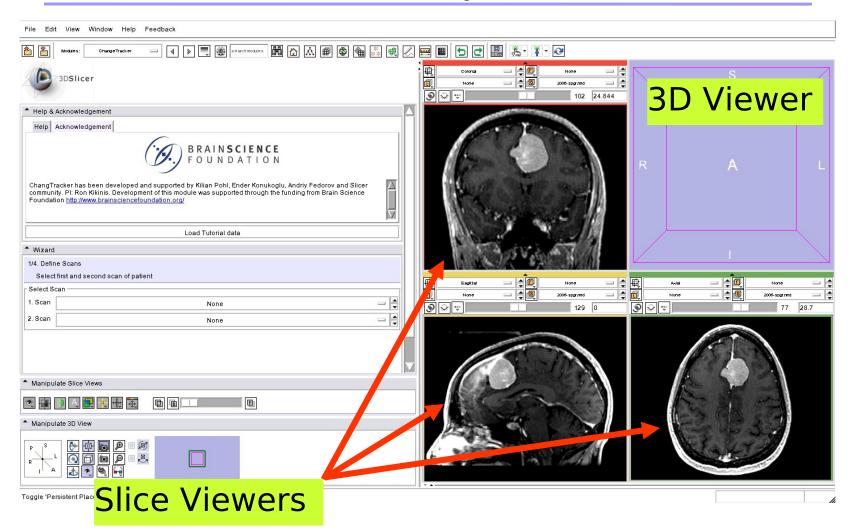


Loading training dataset

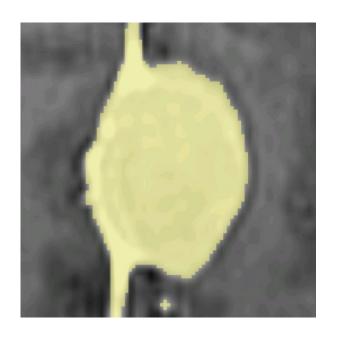




New Layout of Viewer







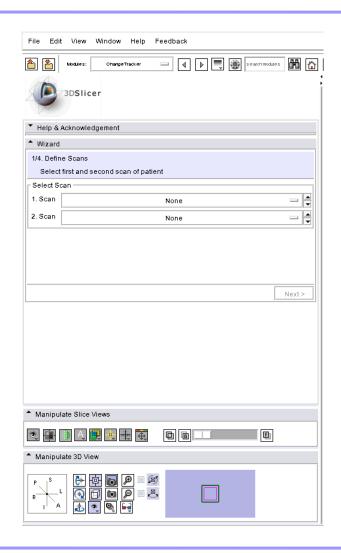
Part 2: Measuring Volume Change



Workflow Wizard

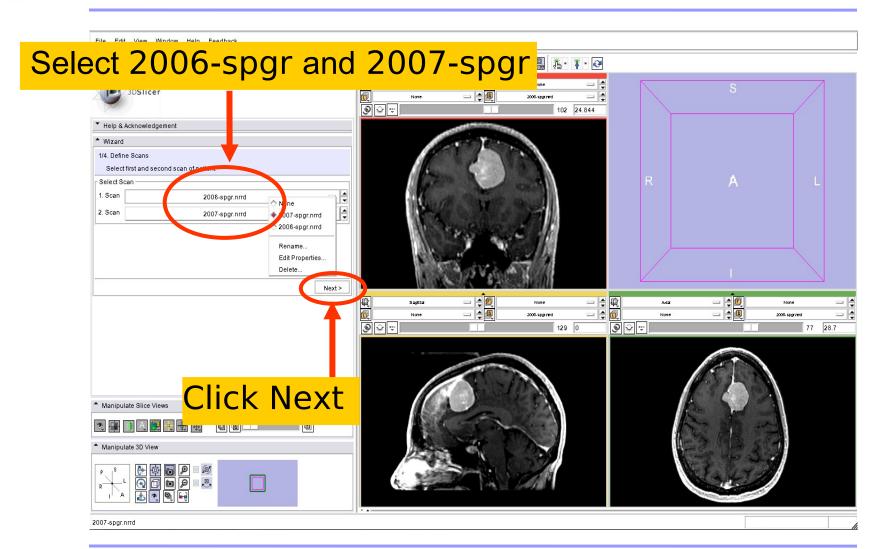
The Workflow Wizard of Slicer3 guides the user through a sequence of steps and is defined by the following components:

- •the Step Panel
- •the User Panel
- the Navigation Panel





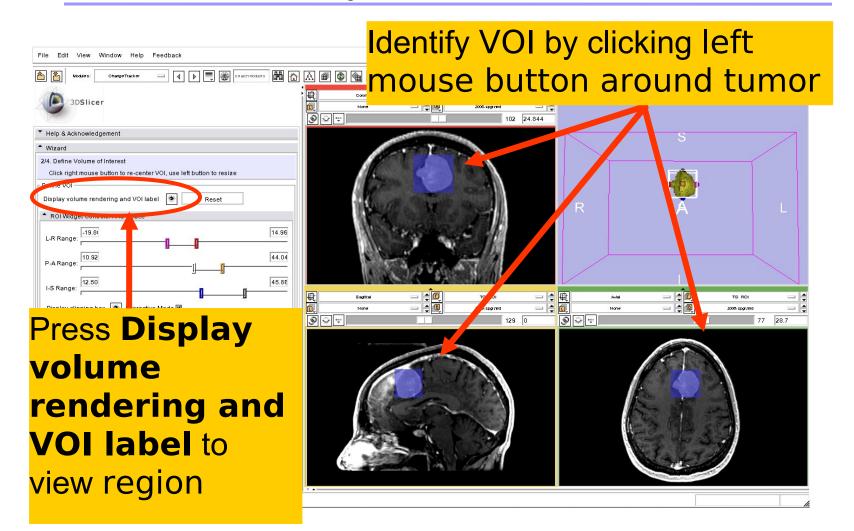
Select Scans



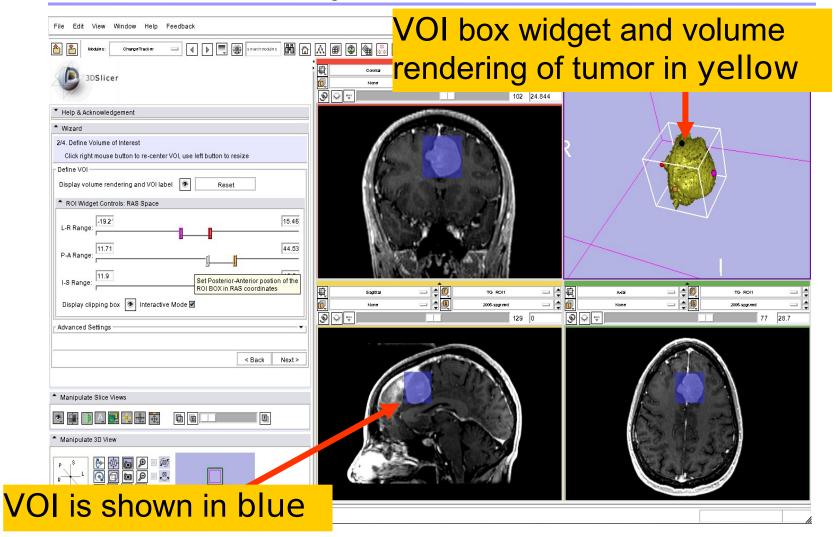






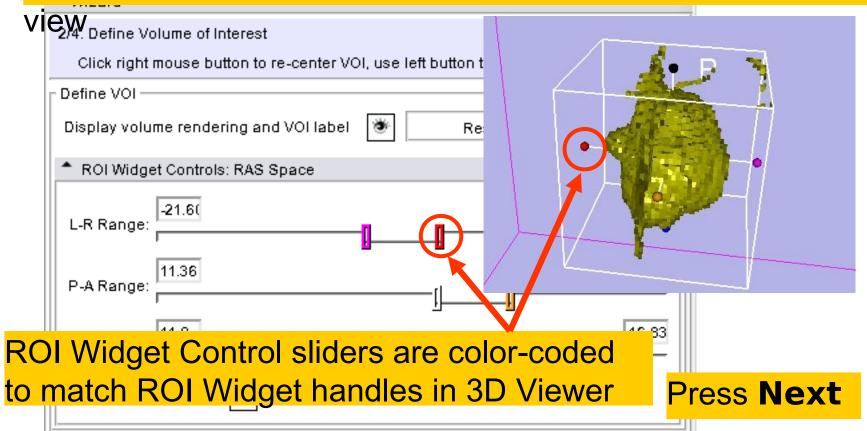






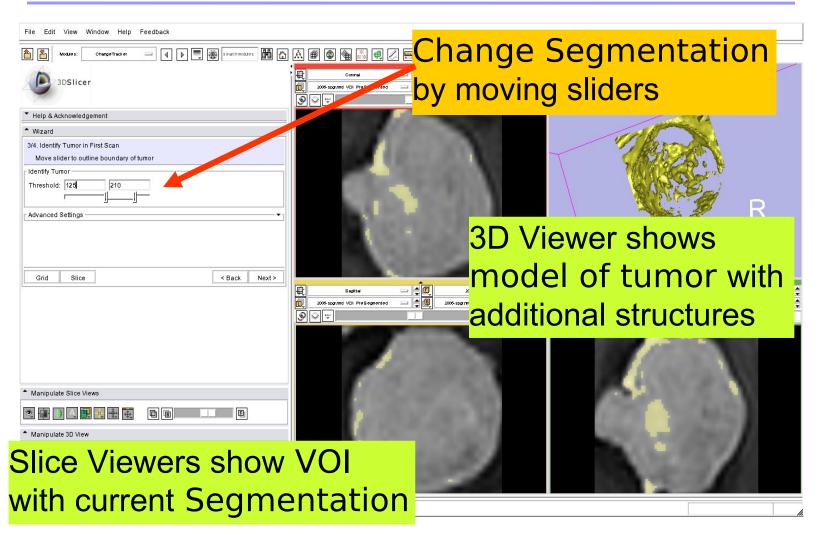


Adjust ROI by moving the **ROI Widget Control sliders** in the Step panel, or by moving the ROI Widget handles in 3D



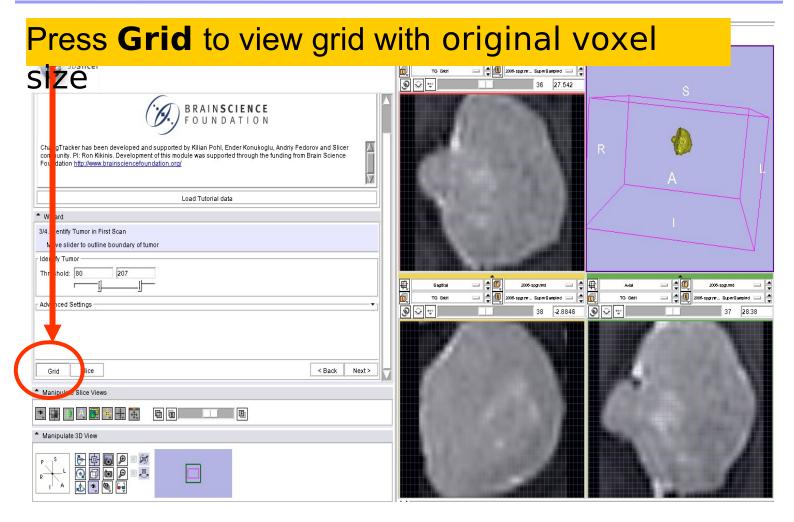


Segment Tumor





Segment Tumor



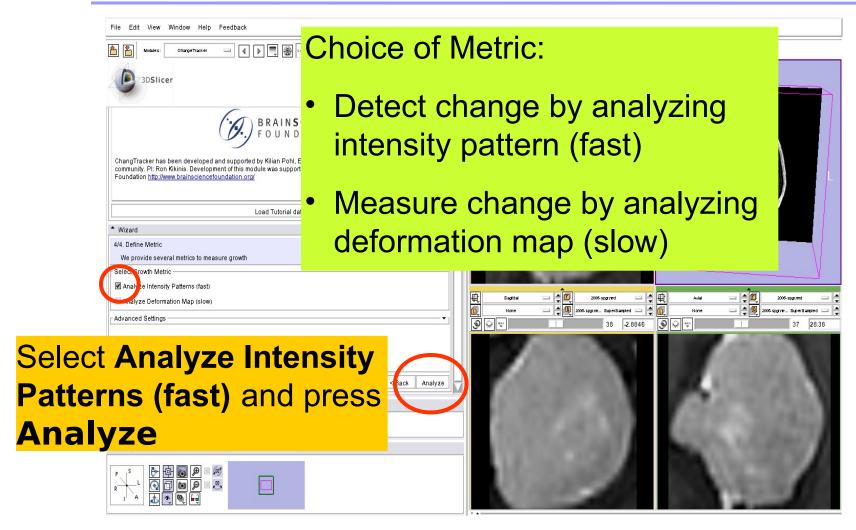


Segment Tumor

Press **Slice** to see the Coror
Non first scan in the 3D Viewer, then press **Next** Foundation http://www.brainsciencefoundation.org/ Load Tutorial data Wizard 3/4. Identify Tumo n First Scan Move slider to tline boundary of tumor Identify Tumor Threshold: 80 Advanced Setting 38 -2.8846 🔊 🔯 🙂 < Back Next > Manipulate 3D View

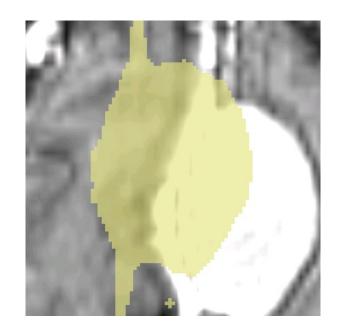


Choose Metric Type

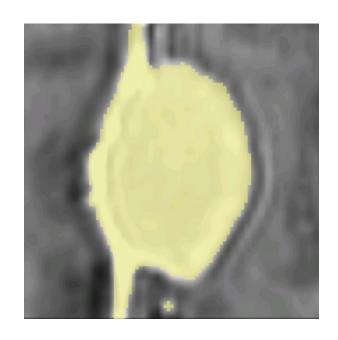




Analysis I: Volume Preserving Registration



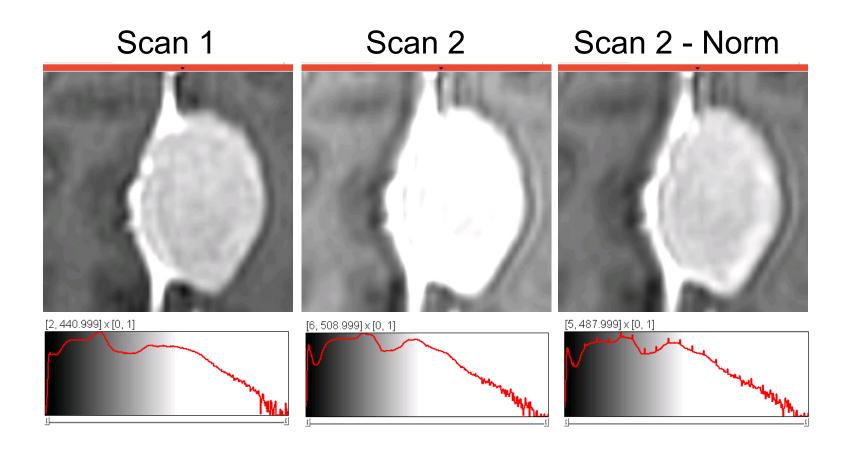
Before



After

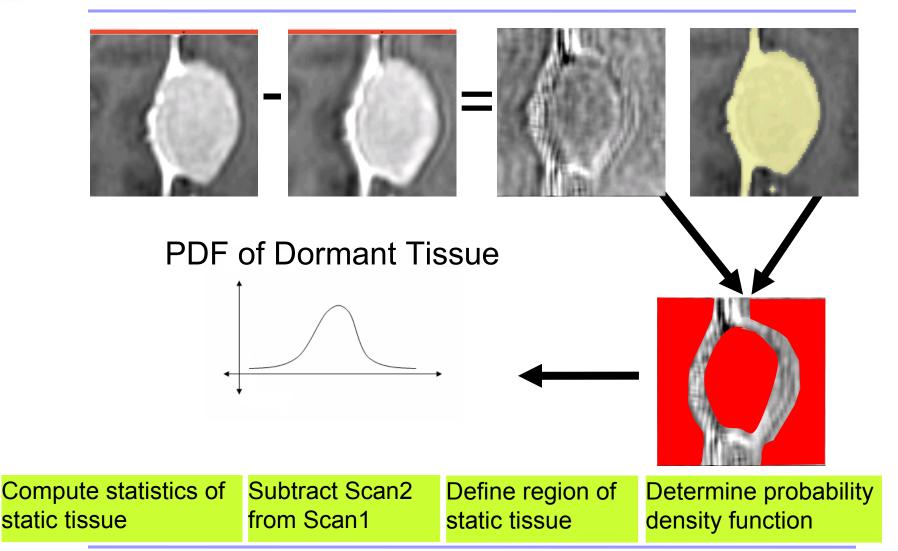


Analysis II: Normalize Intensities



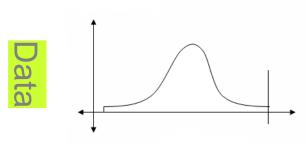


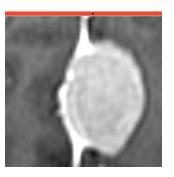
Analysis III: Detect Change – Intensity Based

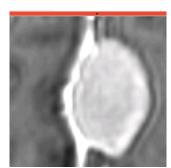




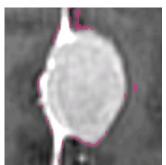
Analysis III: Detect Change – Intensity Based







Analysis

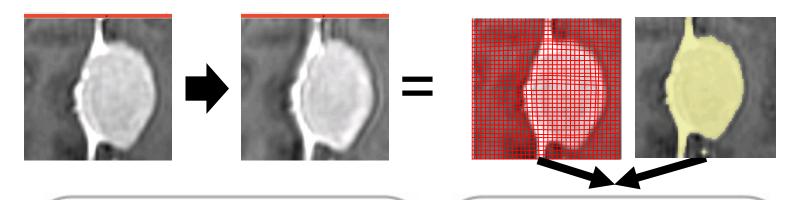


Mode: Sensitive Moderate Robust False Positive Risk: High Medium Low Growth (mm³): 1301 547 100 Growth (voxel): 1057 444 81

Konukoglu et al., "Monitoring Slowly Evolving Tumors", ISBI, 2008



Analysis III: Detect Change–Deformation Map

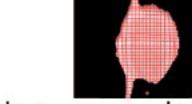




Mode:

Segmentation

Growth (mm³): 1386 Growth (voxels): 1126



Mode: Jaccobian

Growth (mm³): 1291 Growth (voxels): 1049

Determine change via mapping:

Compute diffeomorphic mapping between scans

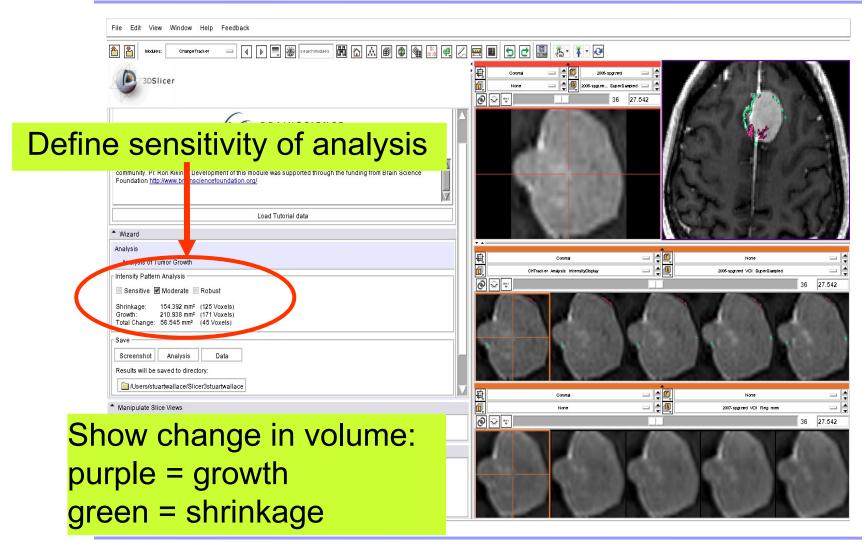
Compute Jacobian

or

Apply mapping to segmentation



Show Change in Pathology



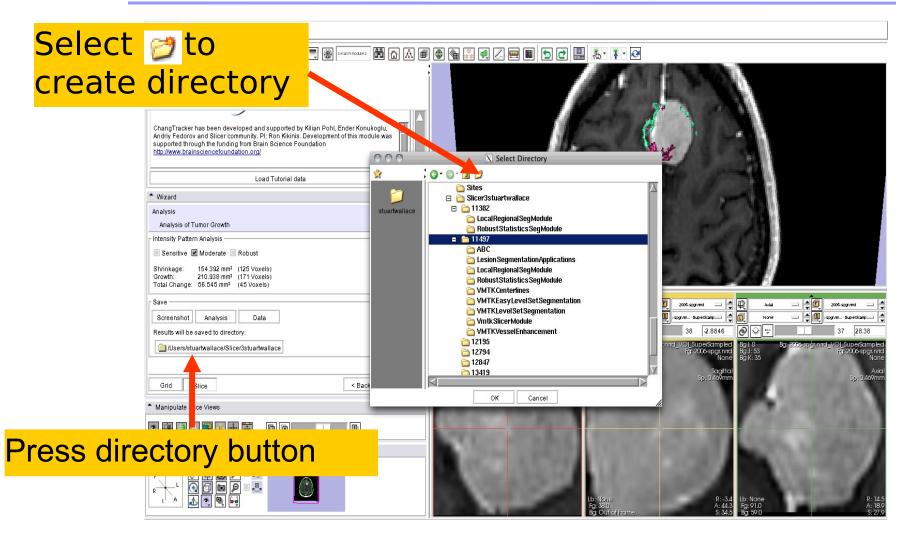


Show Change in Pathology

You can hide slice controls to have more screen space for viewing the images Lightbox View shows 5 consecutive slices for the ROI in the first scan (top), and in the second scan 27.542 **⊚** 💝 👓 (bottom) Screenshot Analysis Results will be saved to directory: I /Users/stuartwallace/Slicer3stuartwallace You can switch to 2007, sported WOL Red north 36 27.542 Four-up layout or other layout at any time.

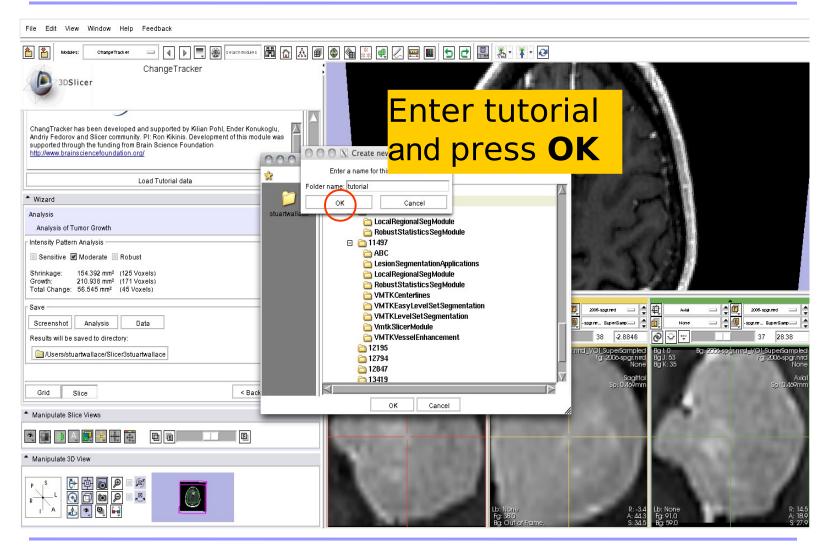


Define Working Directory



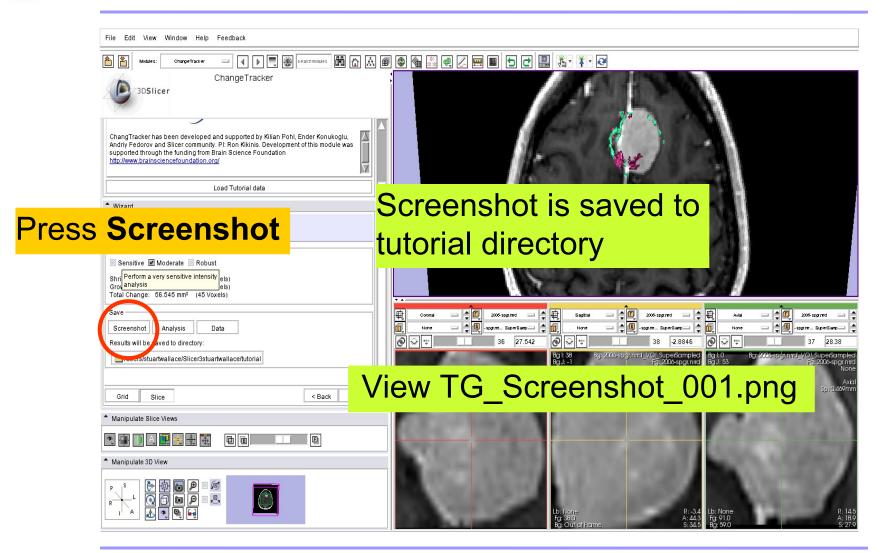


Define Working Directory



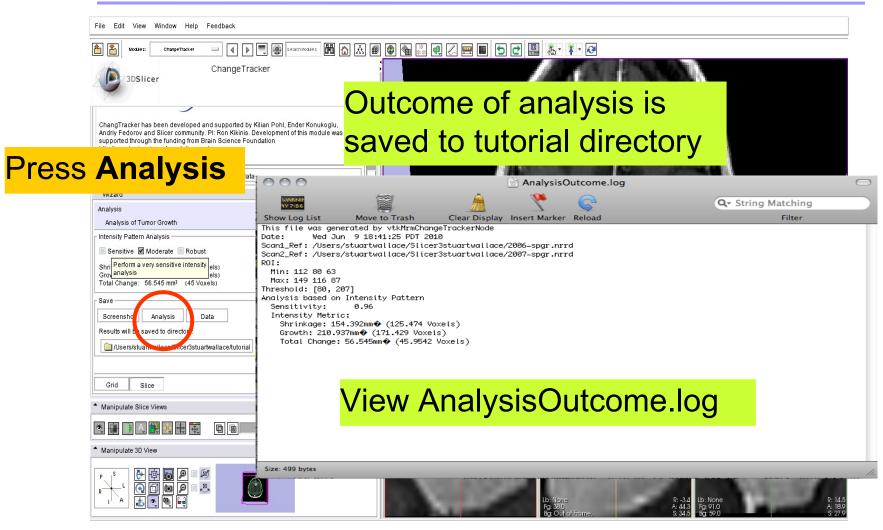


Save Screenshot



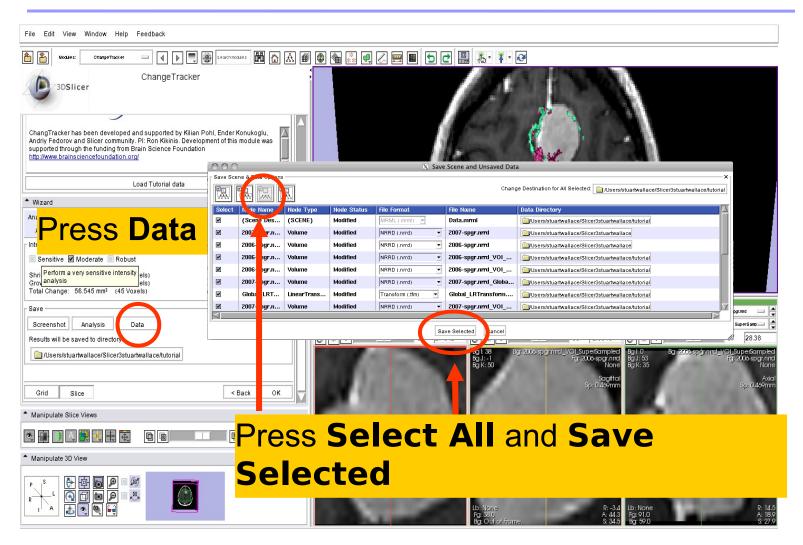


Save Analysis Result



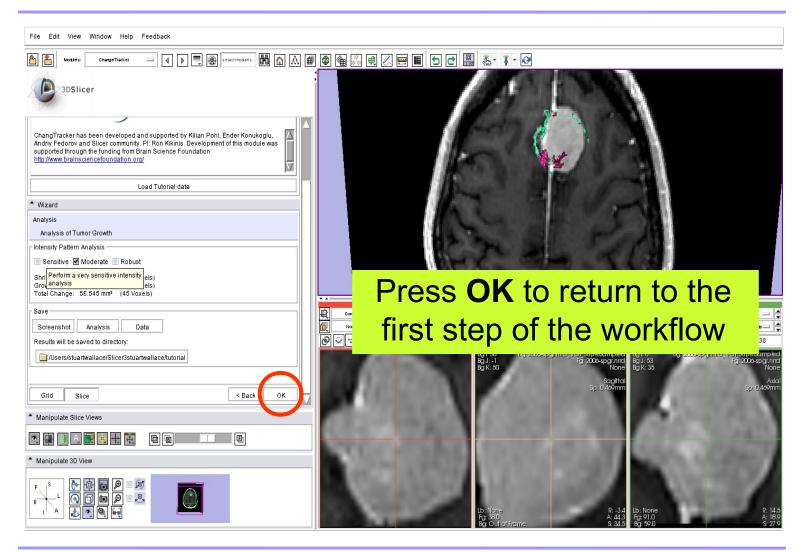


Save Intermediate Results



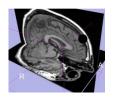


Finish Workflow

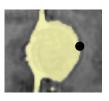




Overview



Loading a scene



Measuring volume change in tumors



Information

- We tested the tool on Axial 3D SPGR T1 post Gadolinium scans (Voxel dimension: 0.94mm x 0.94mm x 1.20mm, FOV: 240mm, Matrix: 256 x 256)
- We expect the tool to work also on set of scans that fulfill the following requirements:
- same patient was scanned
- scans were acquired with the same acquisition protocol
- scans have isotropic resolution
- pathology appearance is hyper-intense



Acknowledgments



National Alliance for Medical Image Computing
NIH U54EB005149



Brain Science Foundation



INRIA, France