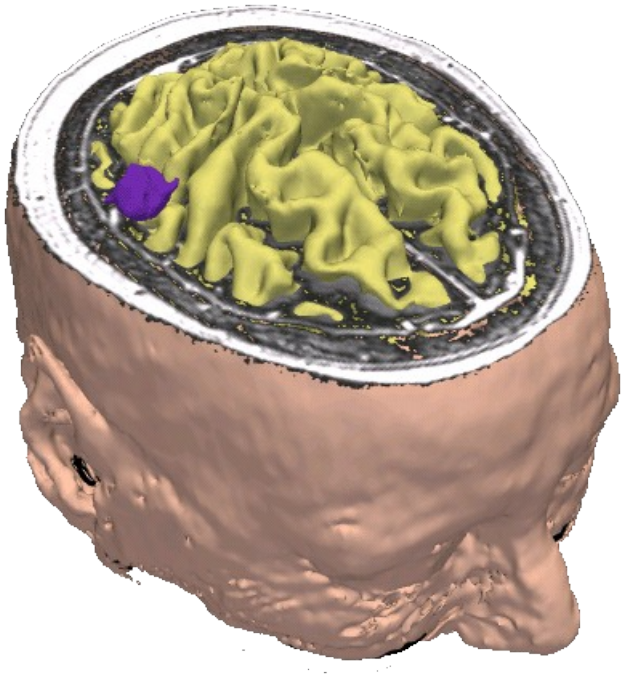
Measuring Volume Change in Tumors



Kilian M Pohl, PhD
Ender Konugolu, PhD
Andriy Fedorov, PhD

Material

This course requires the following installation:

- 3DSlicer version 3.4.1 Software, which can be installed from
http://www.slicer.org/pages/Special:SlicerDownloads
- A reliable internet connection for downloading the data

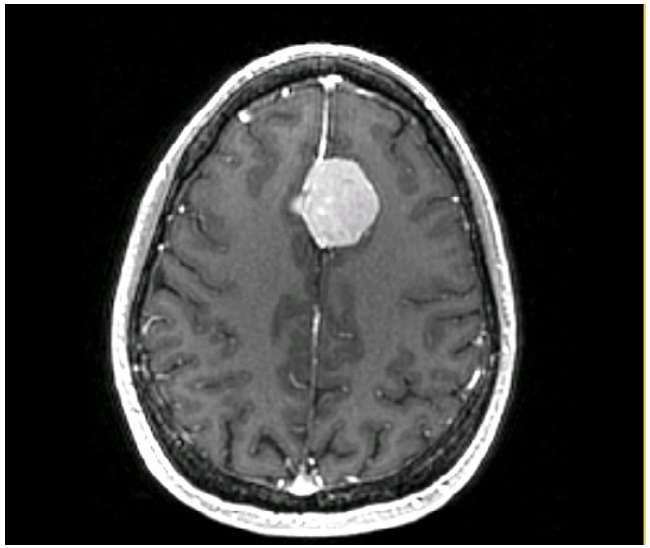
Disclaimer

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.

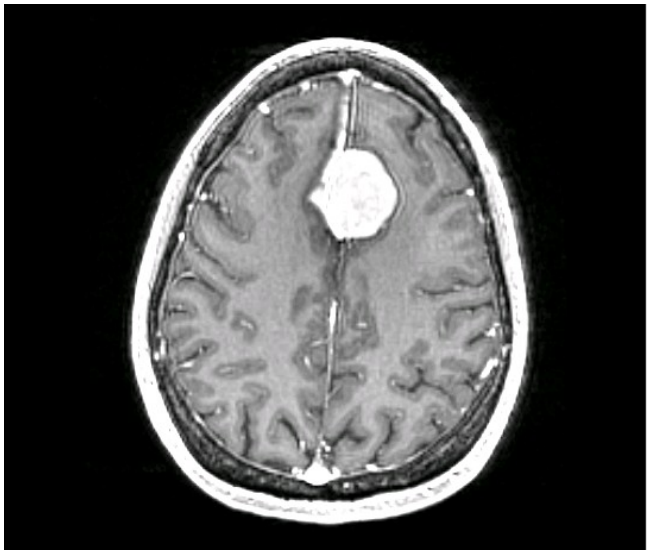
The module described in this tutorial was tested on Axial 3D SPGR T1 post Gadolinium scans (Voxel dimension: 0.94mm x 0.94mm x 1.20mm, FOV: 240mm, Matrix: 256 x 256)

Data

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



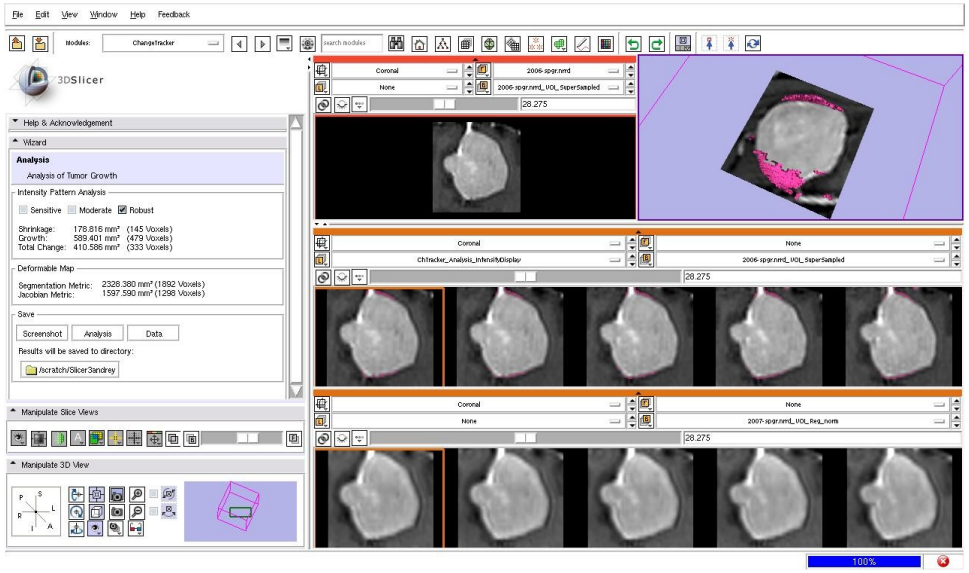
MR Scan 1



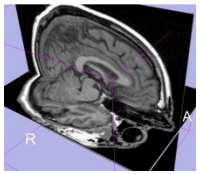
MR Scan 2

Learning objective

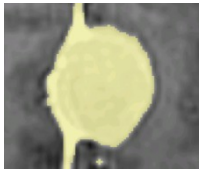
Following this tutorial, you'll be able to load two scans into Slicer3, and measure and visualize the change in volume of the meningioma between the two scans.



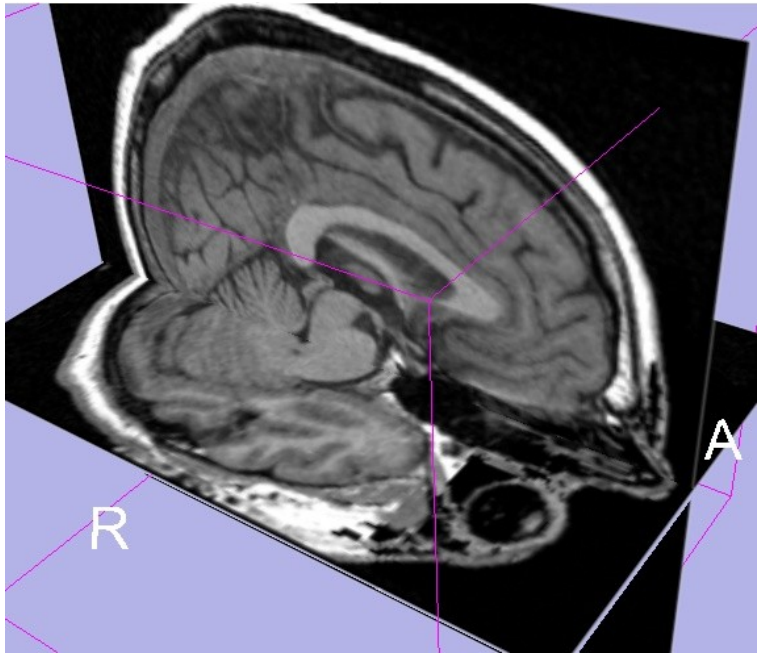
Overview



Loading tutorial data



Measuring volume change in tumors

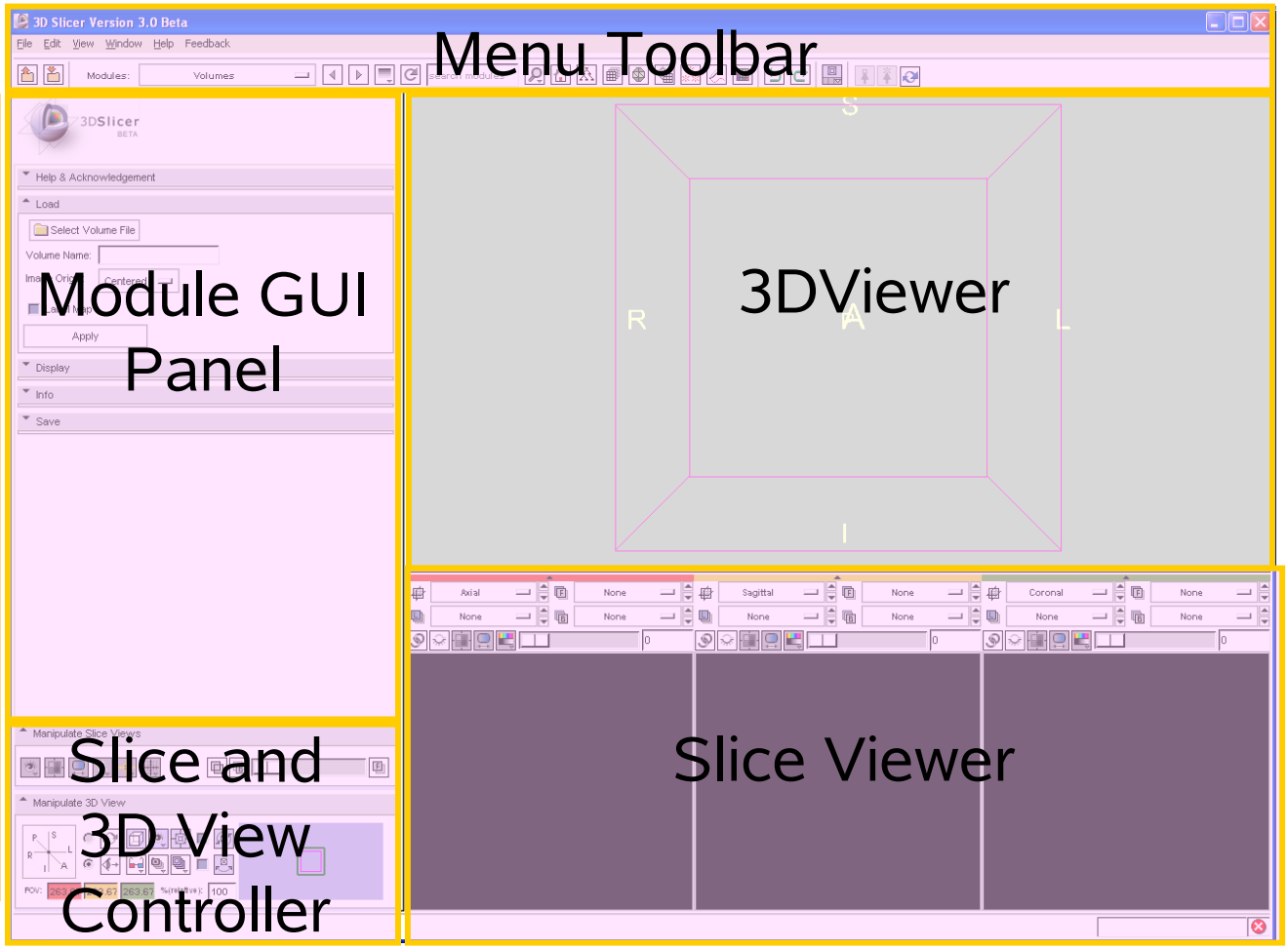


Part 1: Loading the tutorial data

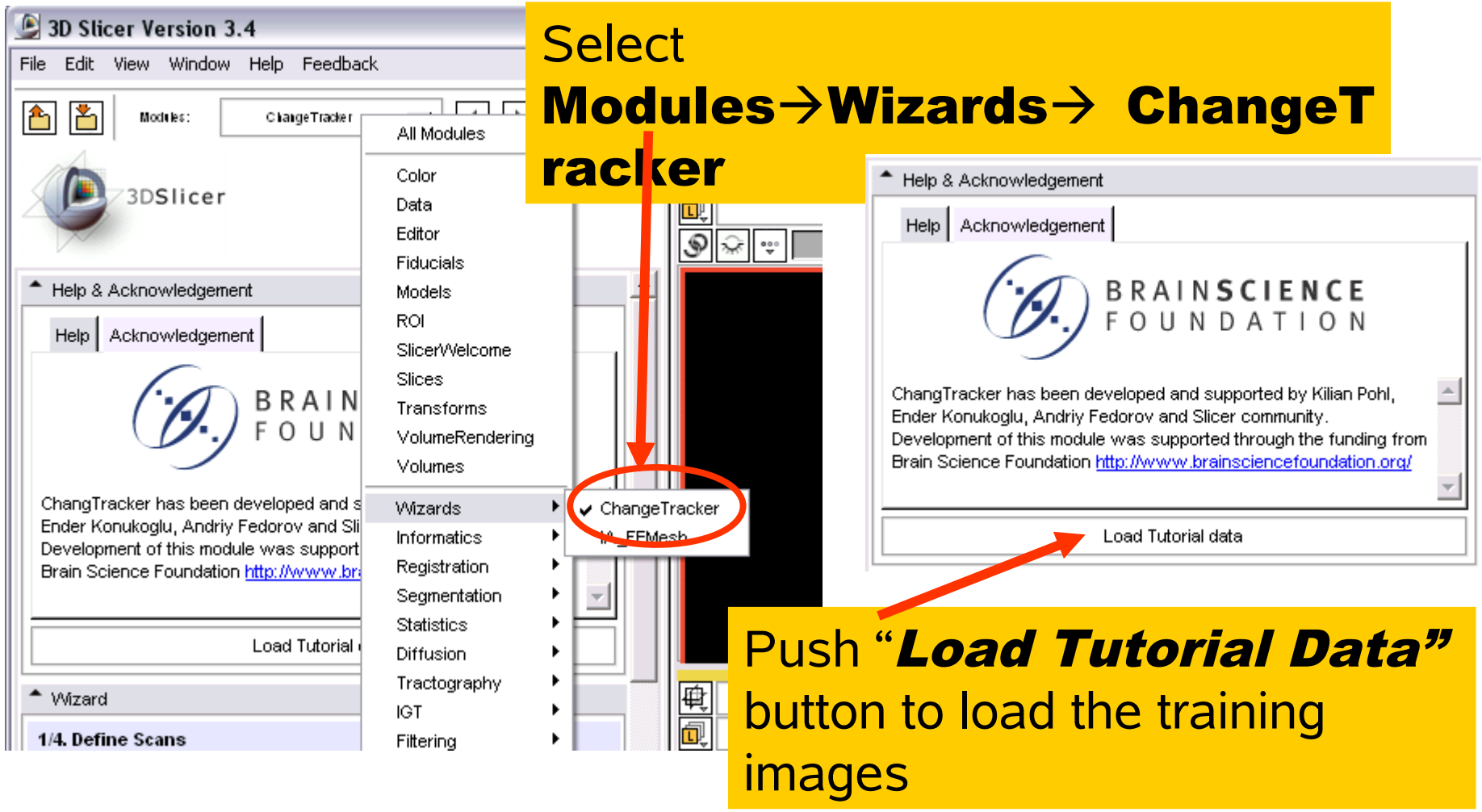
Slicer3 GUI

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

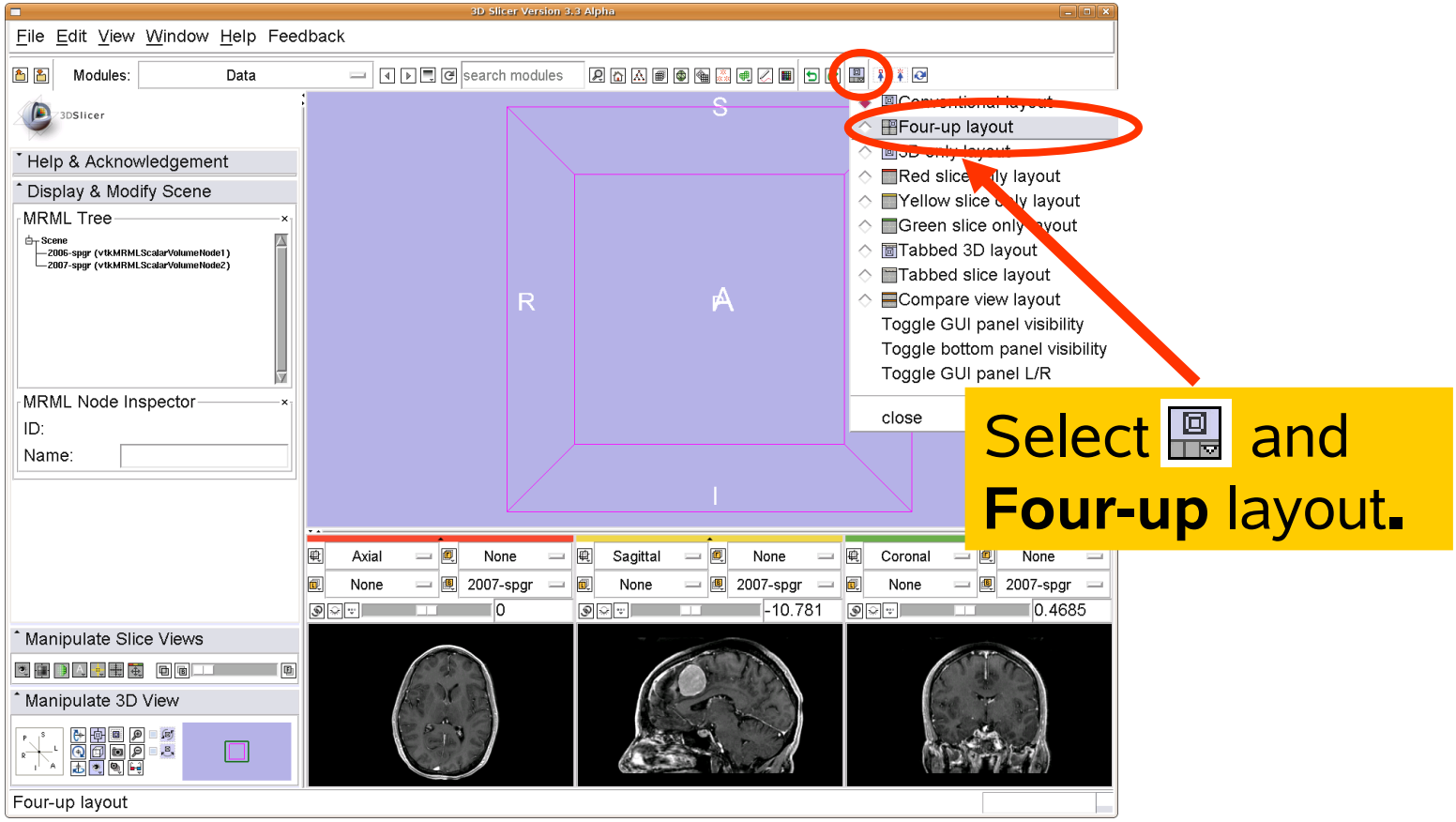
- the Menu Toolbar
- the Module GUI Panel
- the 3D Viewer
- the Slice Viewer
- the Slice and 3D View Controller



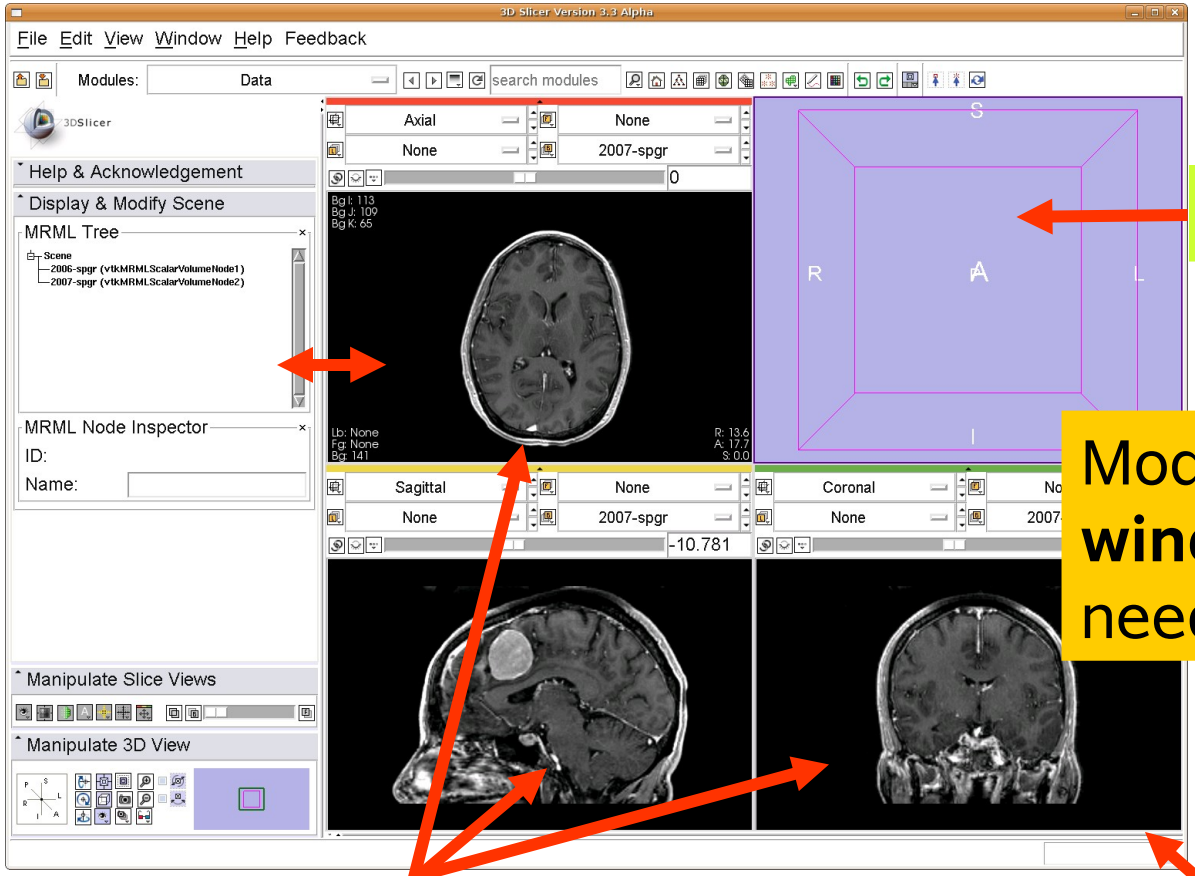
Loading training dataset

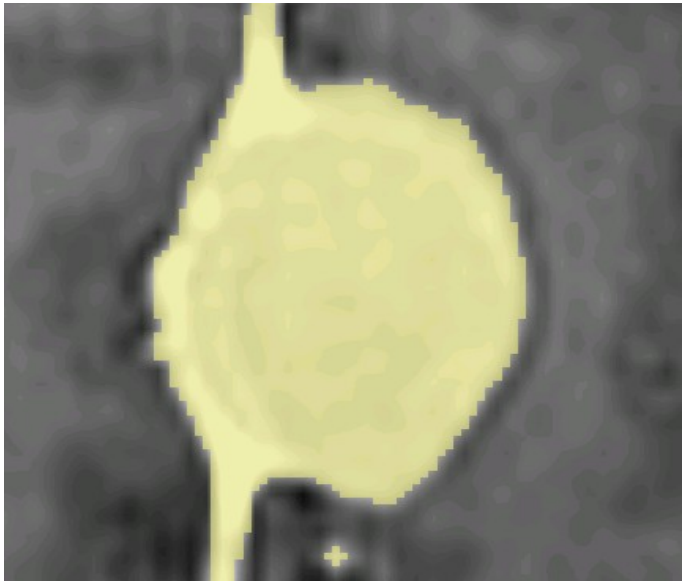


Modifying Layout of Viewer



New Layout of Viewer



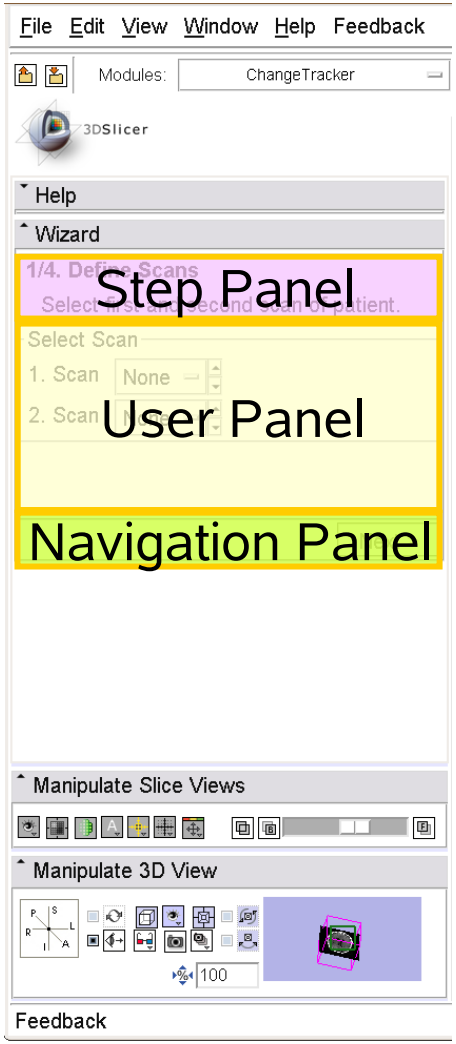


Part 2: Measuring Volume Change

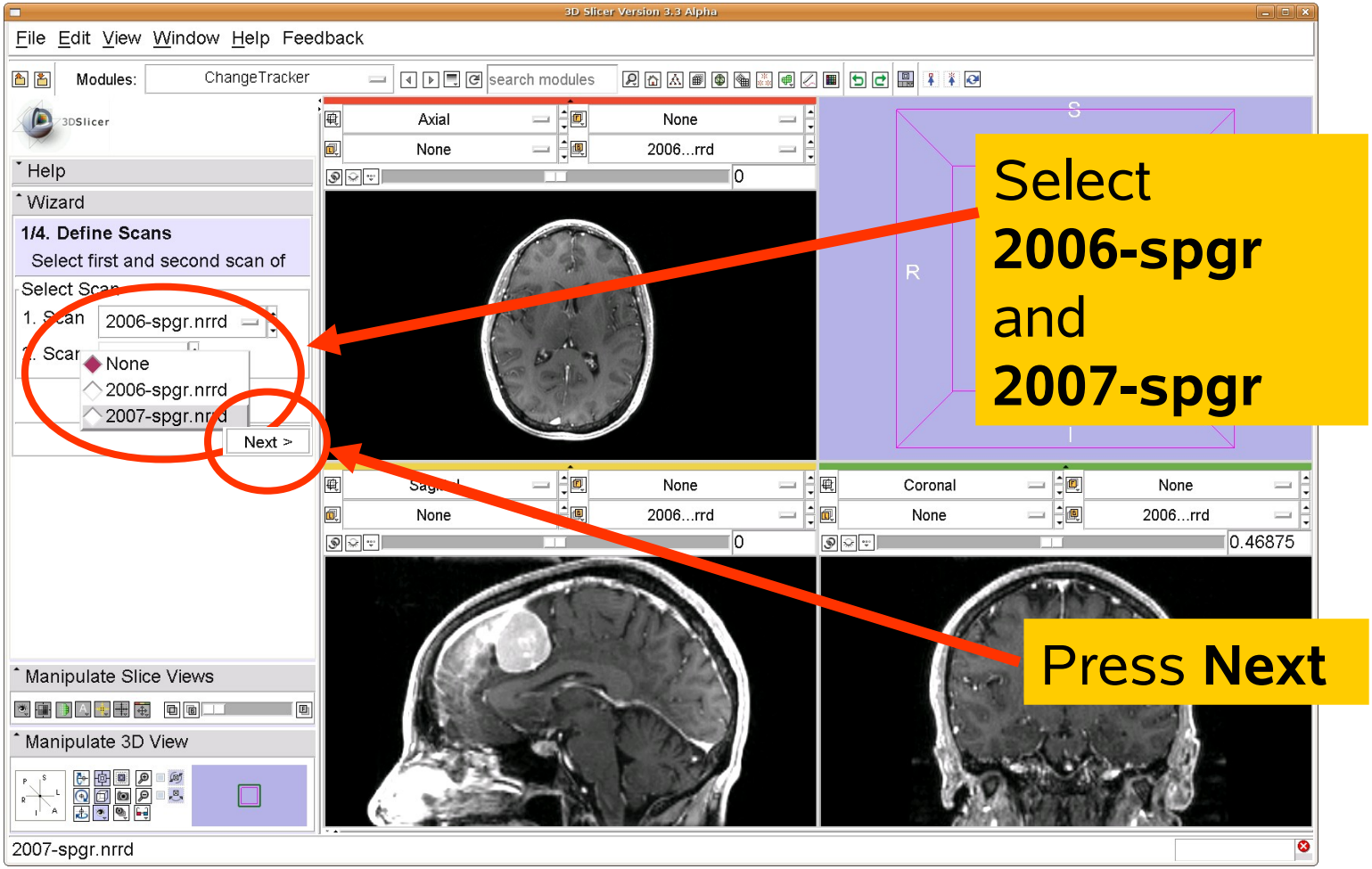
Workflow Wizard

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

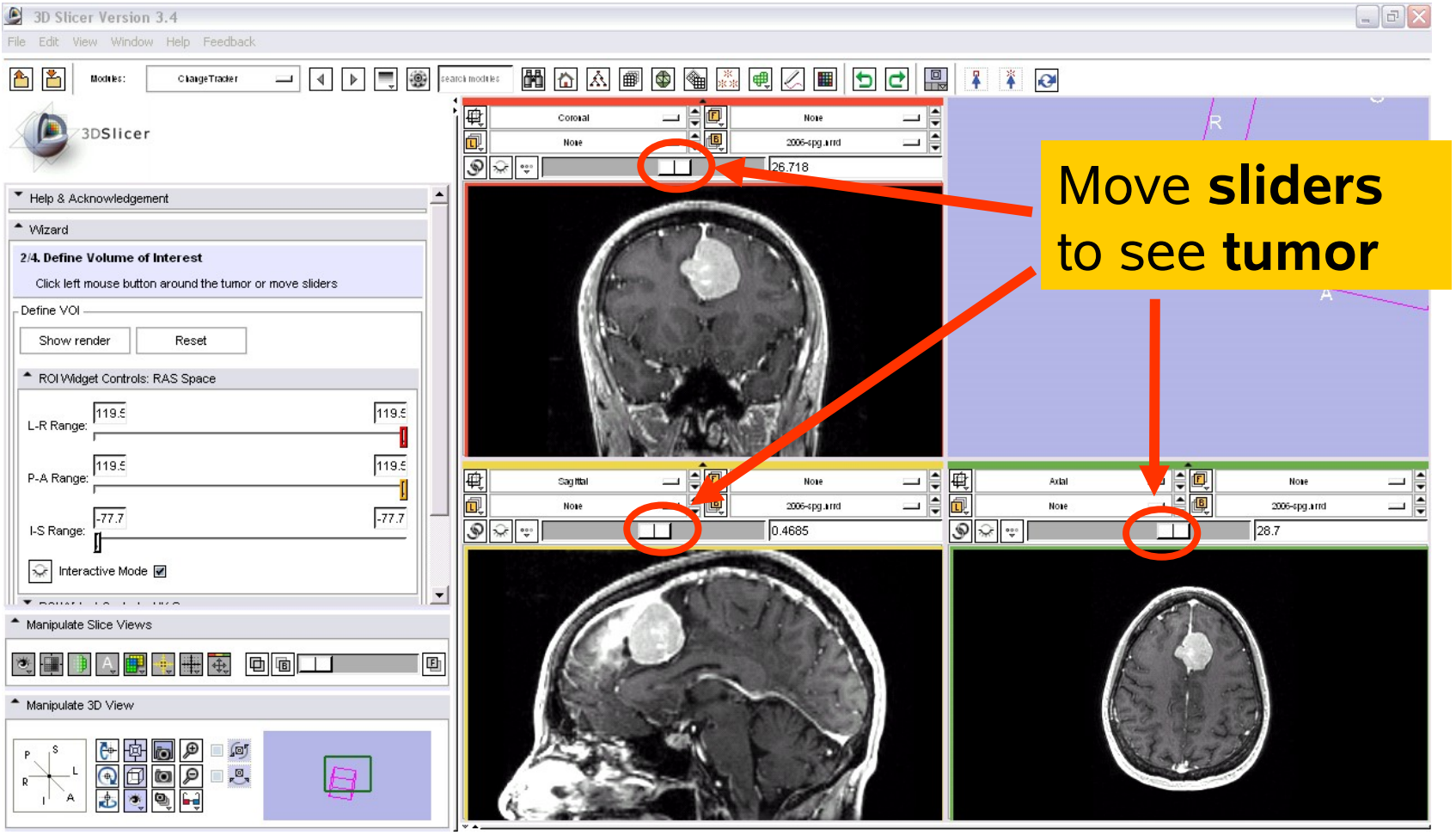
- the Step Panel
- the User Panel
- the Navigation Panel



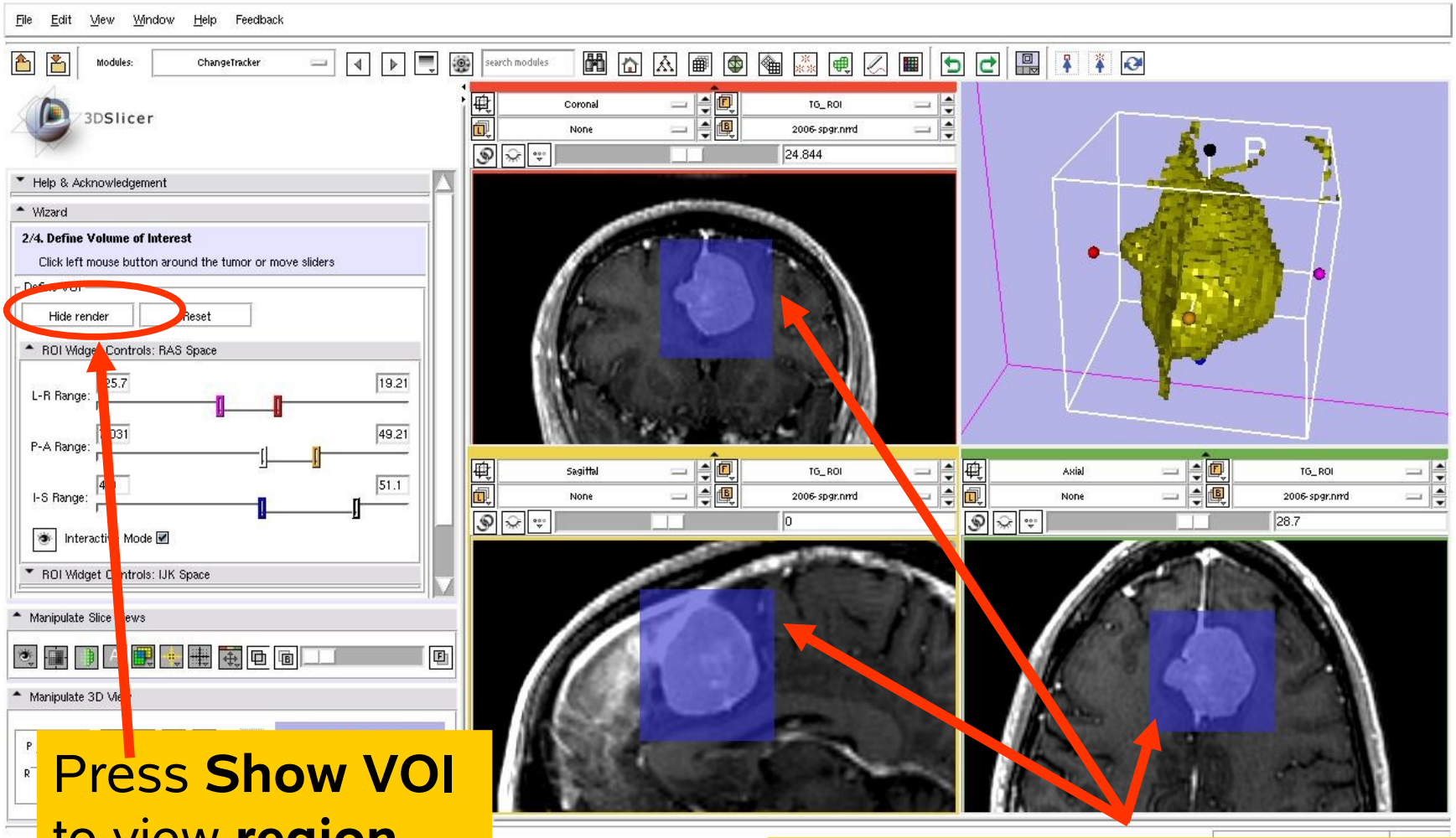
Select Scans



Identify Volume of Interest

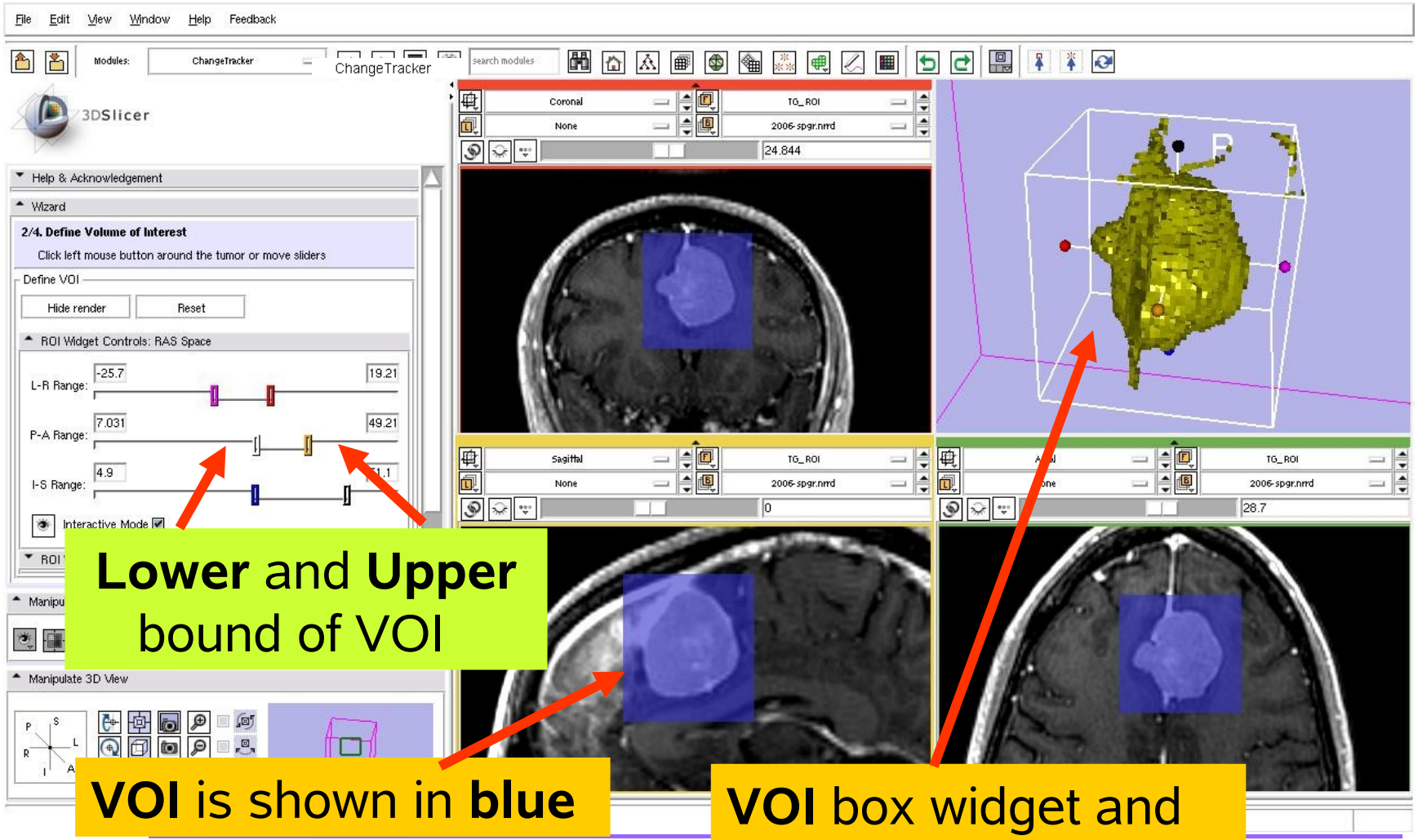


Identify Volume of Interest



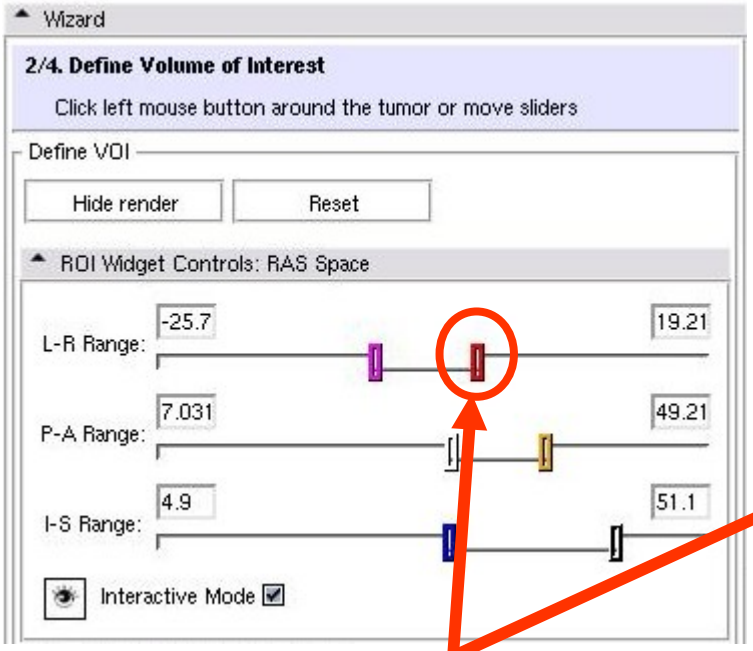
Pohl K, Konukoglu E, Fedorov A
National Alliance for Medical Image Computing

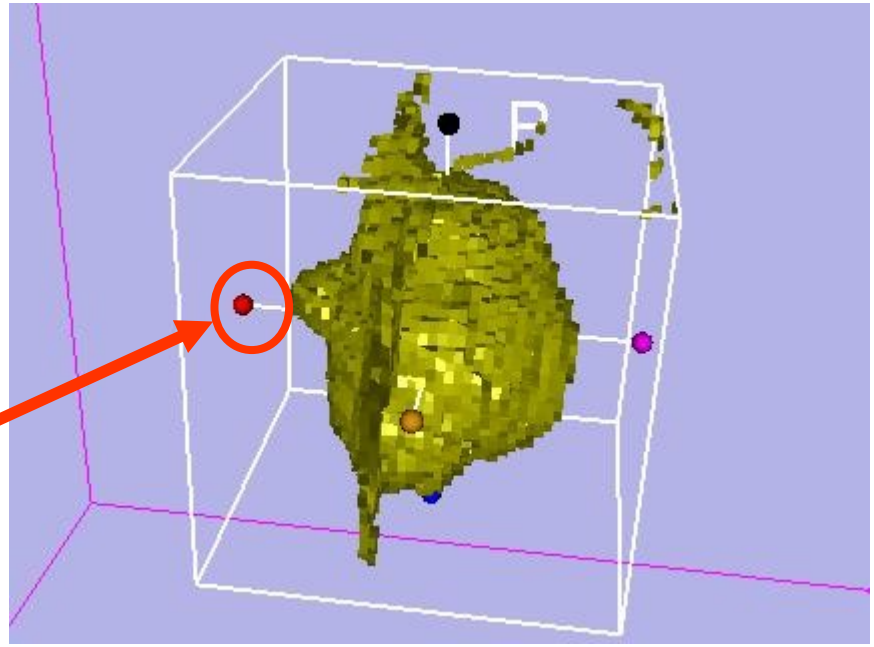
Identify Volume of Interest



Identify Volume of Interest

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

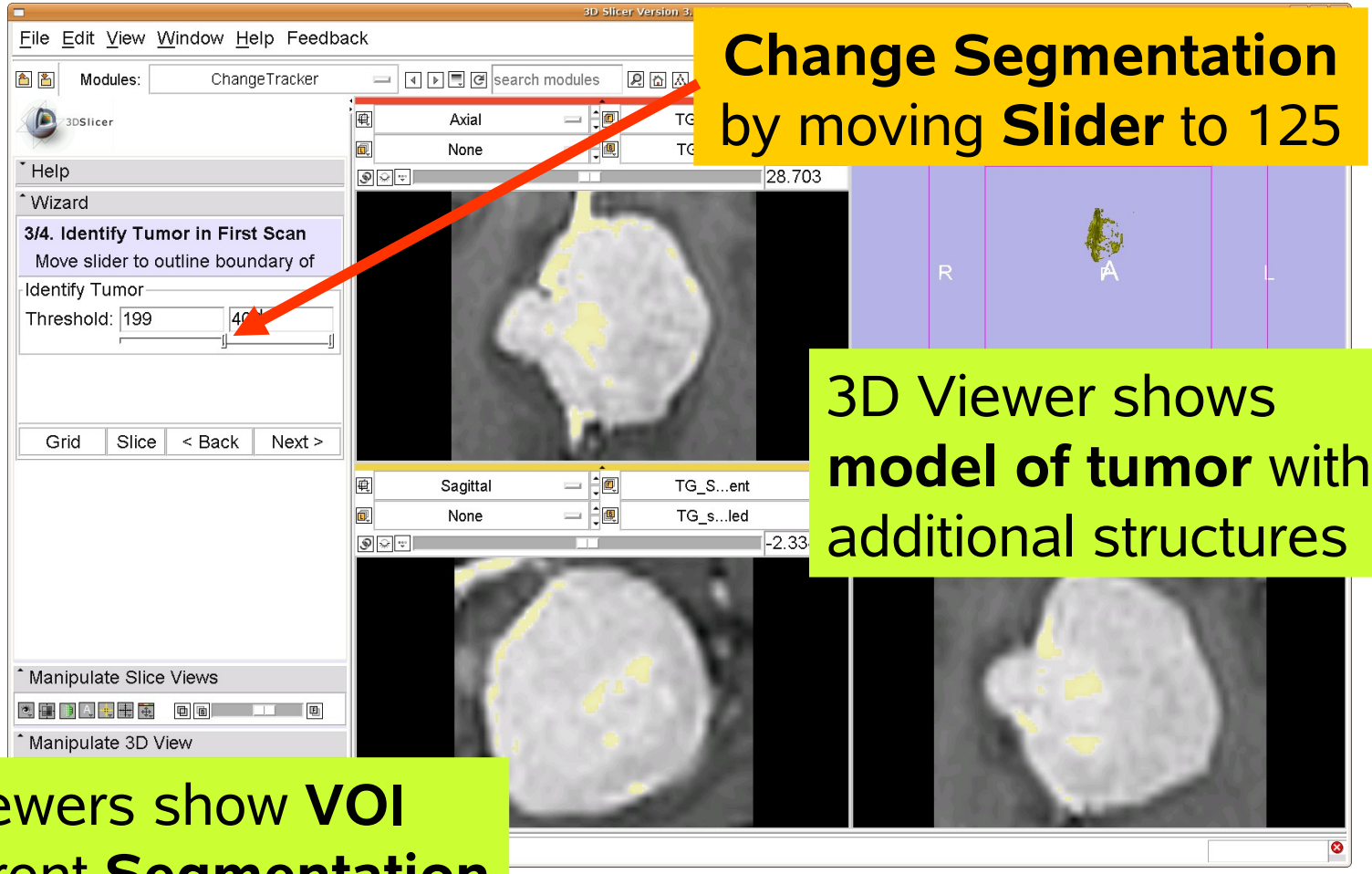




ROI Widget Control sliders are color-coded to match ROI Widget handles in 3D Viewer

Press **Next**

Segment Tumor

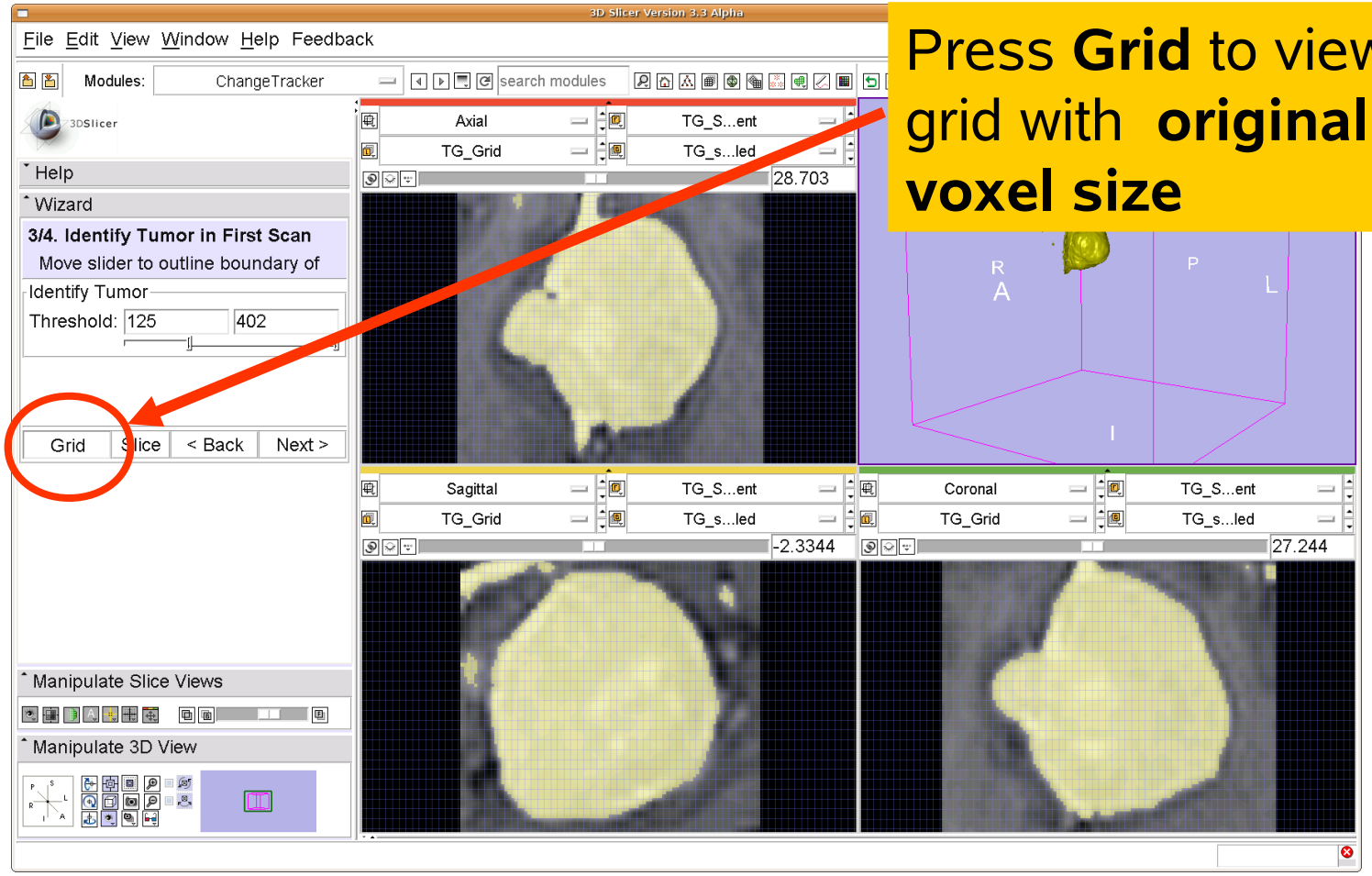


Change Segmentation by moving **Slider** to 125

3D Viewer shows **model of tumor** with additional structures

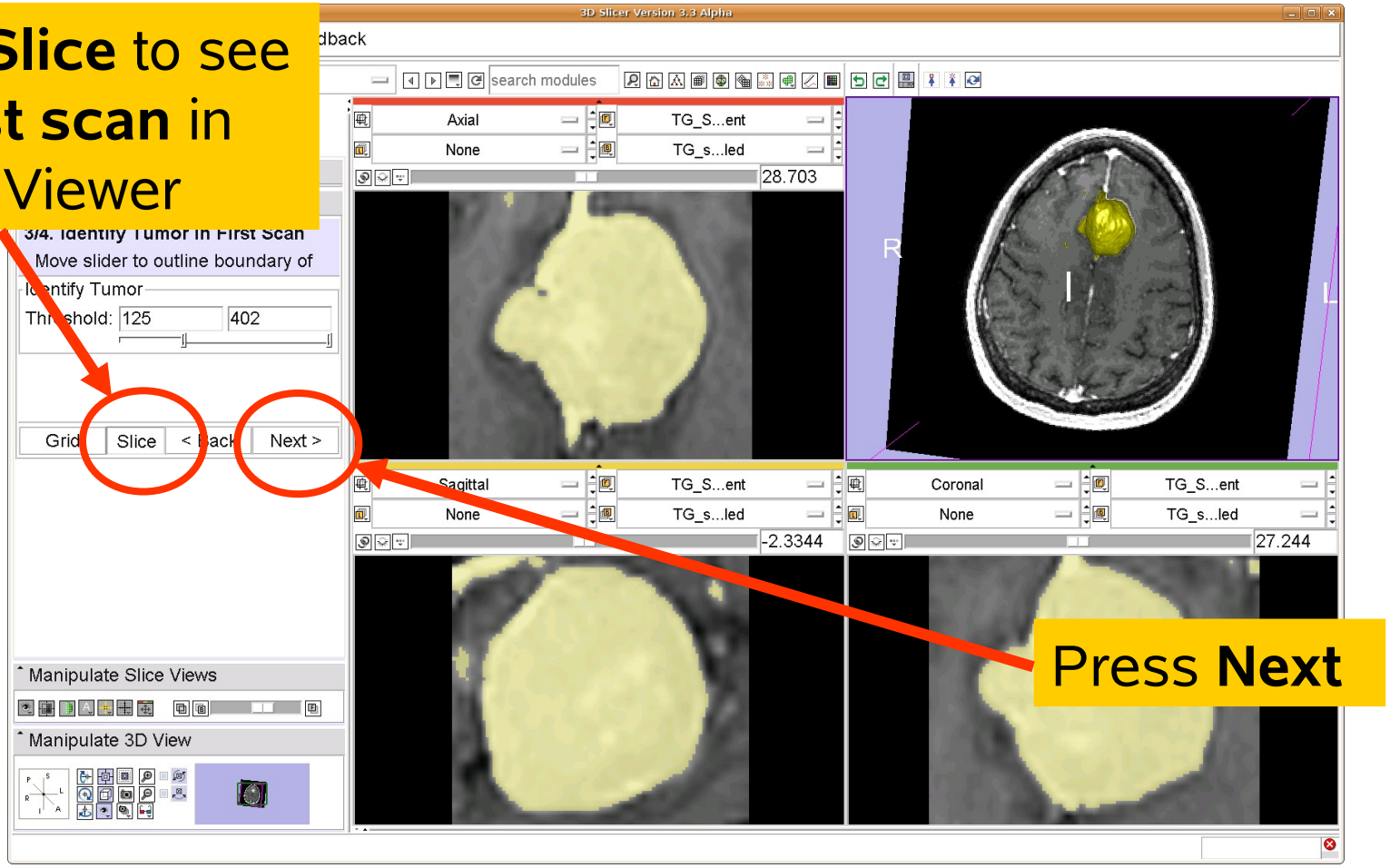
Slice Viewers show **VOI** with current **Segmentation**

Segment Tumor

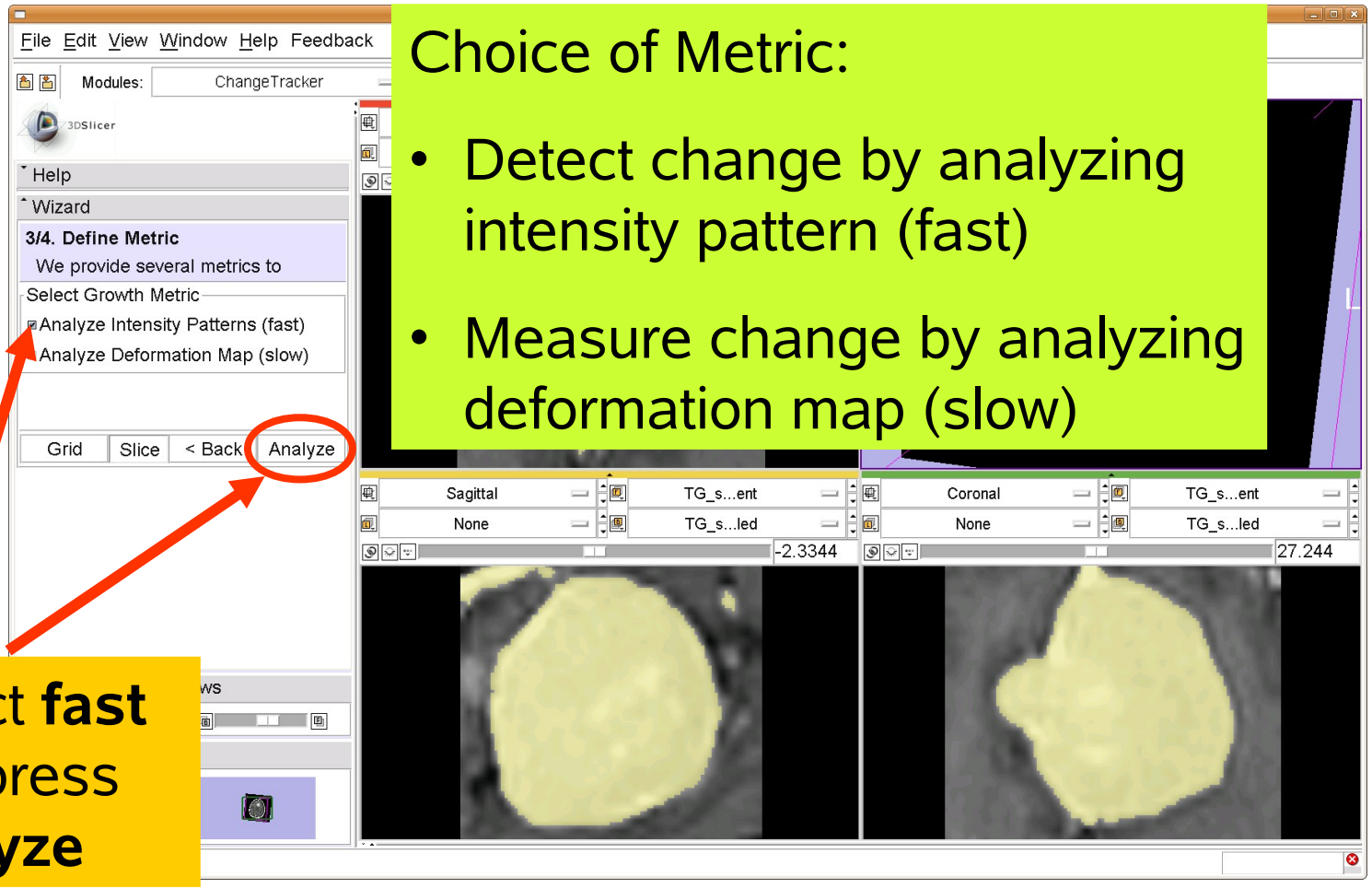


Segment Tumor

Press **Slice** to see the **first scan** in the 3D Viewer



Choose Metric Type

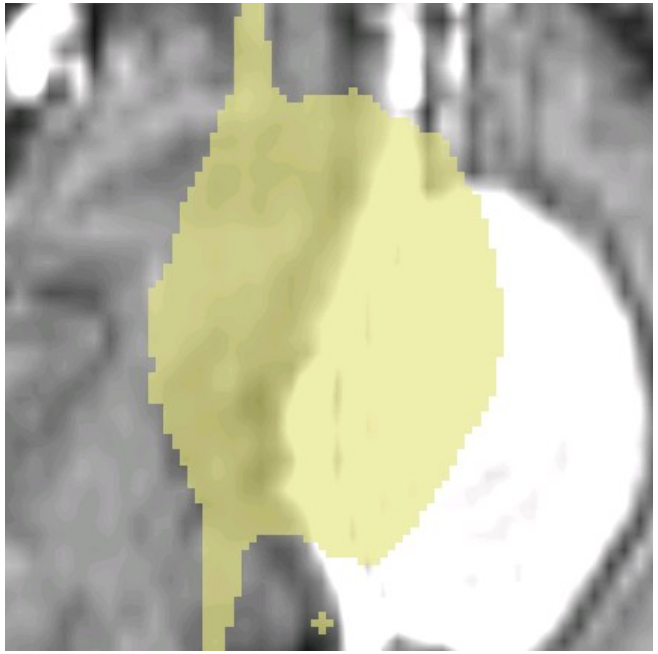


Choice of Metric:

- Detect change by analyzing intensity pattern (fast)
- Measure change by analyzing deformation map (slow)

Select **fast** and press **Analyze**

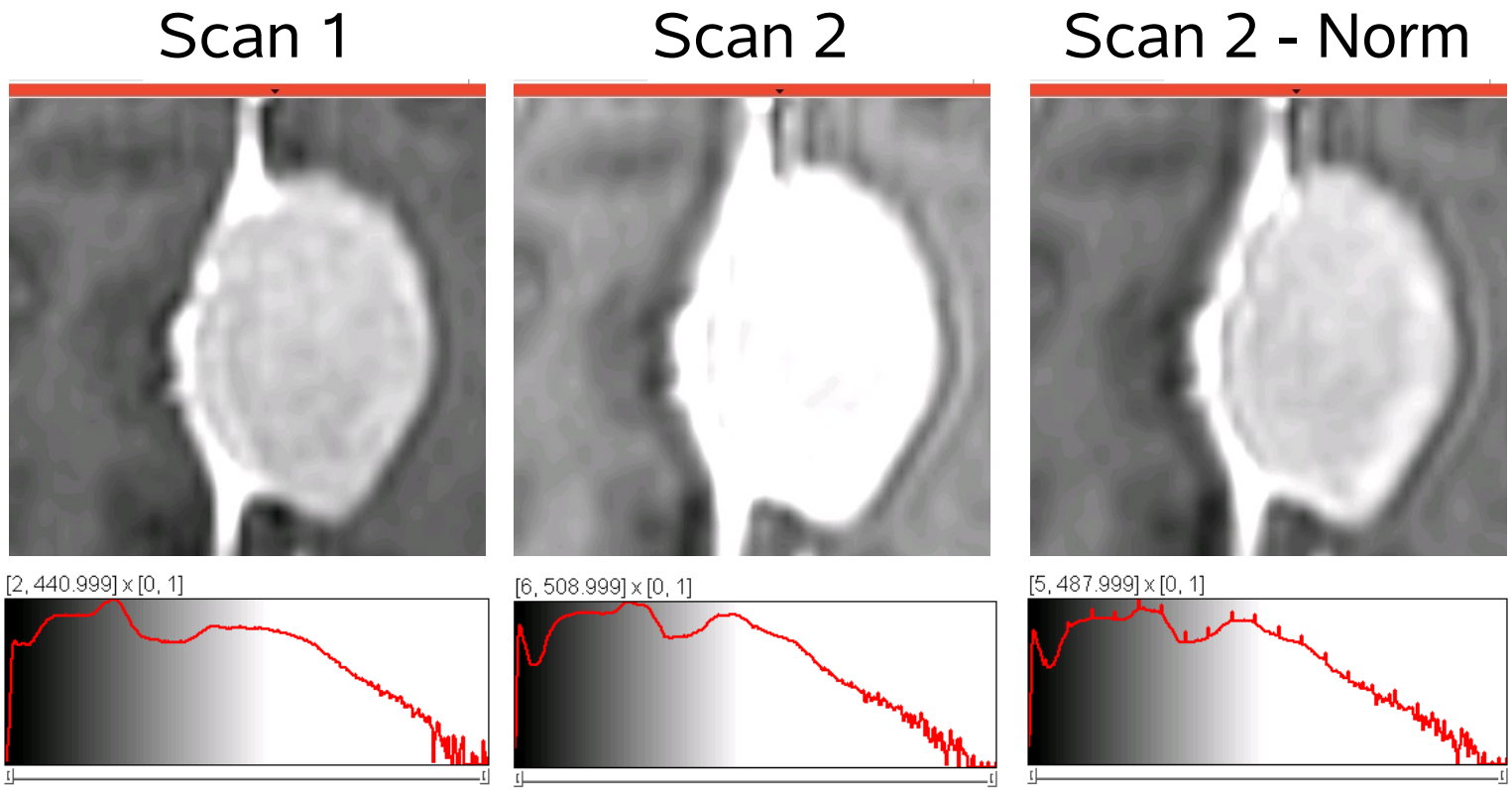
Analysis I: Volume Preserving Registration



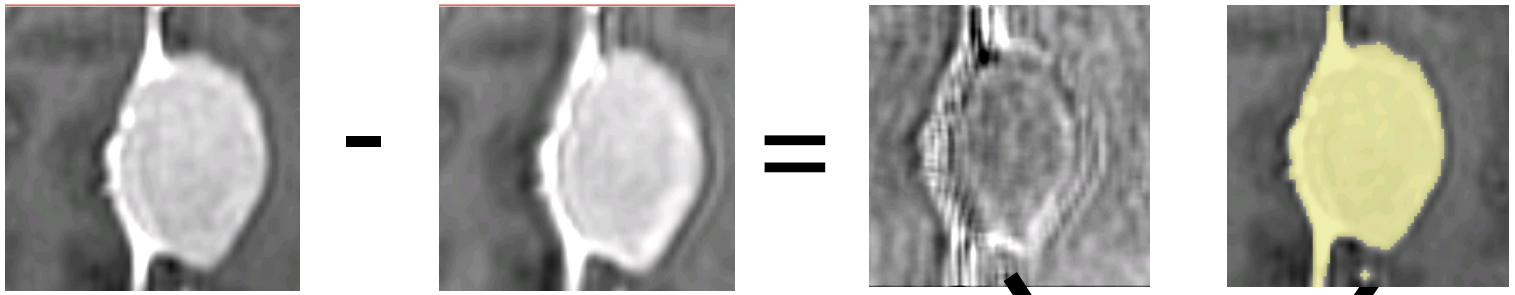
Before

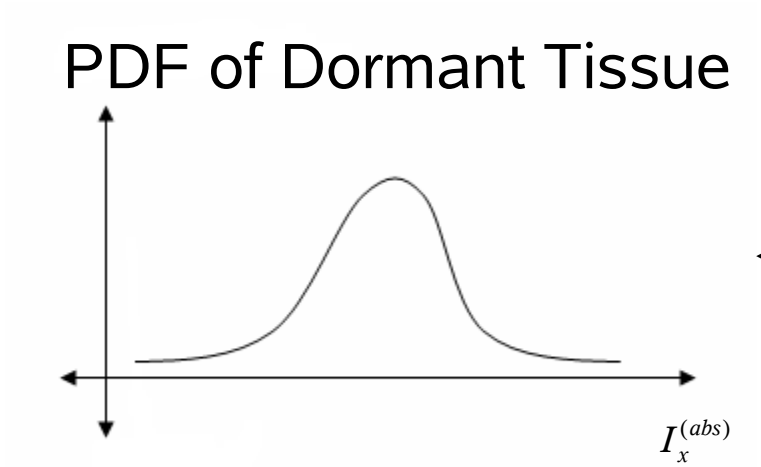
Afterwards

Analysis II: Normalize Intensities



Analysis III: Detect Change – Intensity Based

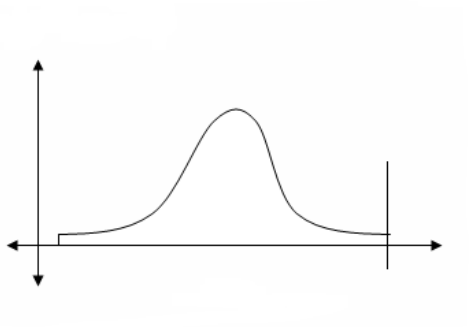


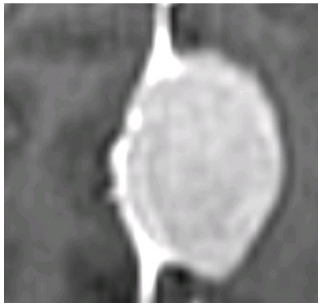


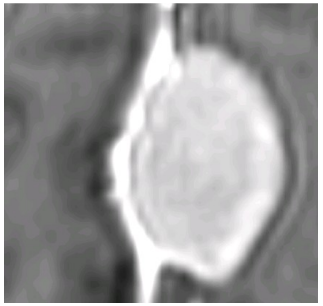
Compute statistics of static tissue:	Subtract Scan2 from Scan1	Define region of static tissue	Determine probability density function

Analysis III: Detect Change – Intensity Based

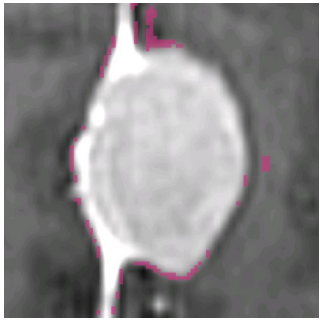
Data







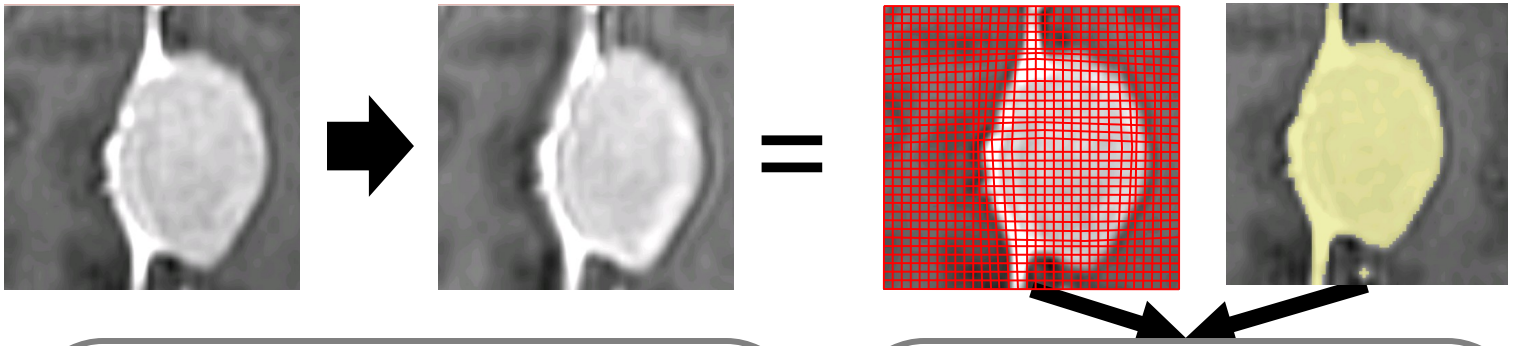
Analysis



Mode : Sensitive
False Positive Risk: High
Growth (mm^3) : 1301
Growth (voxel): 1057

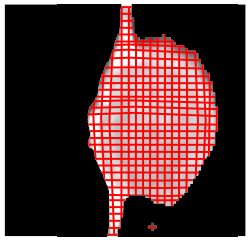
Konukoglu et al. ,"Monitoring Slowly Evolving Tumors", ISBI, 2008

Analysis III: Detect Change–Deformation Map



Mode : Segmentation
Growth (mm^3) : 1386

Growth (voxels): 1126



Mode : Jaccobian
Growth (mm^3) : 1291

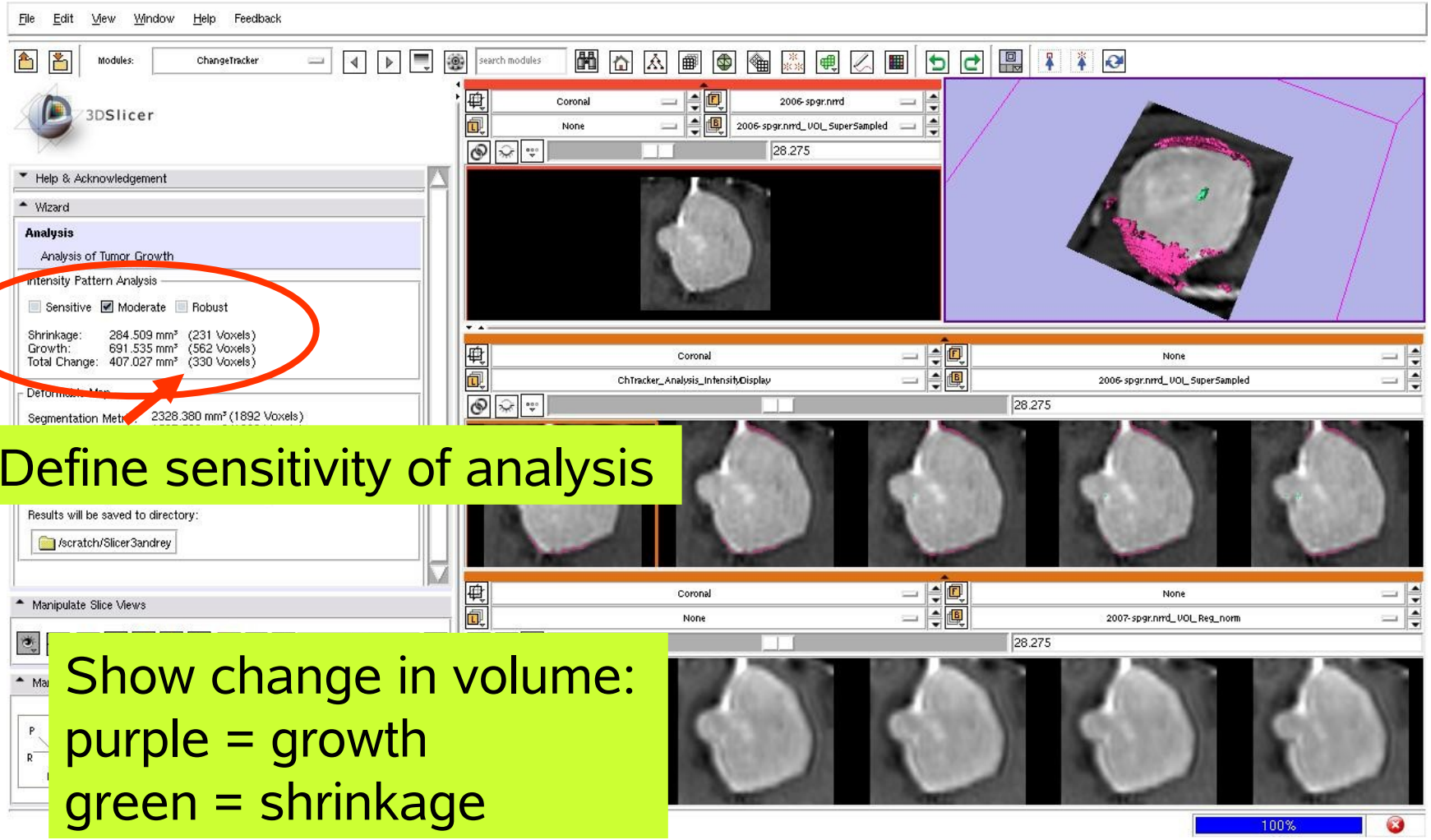
Growth (voxels): 1049

Determine change via mapping:

Compute diffeomorphic mapping between scans

Compute Jacobian or Apply mapping to segmentation

Show Change in Pathology

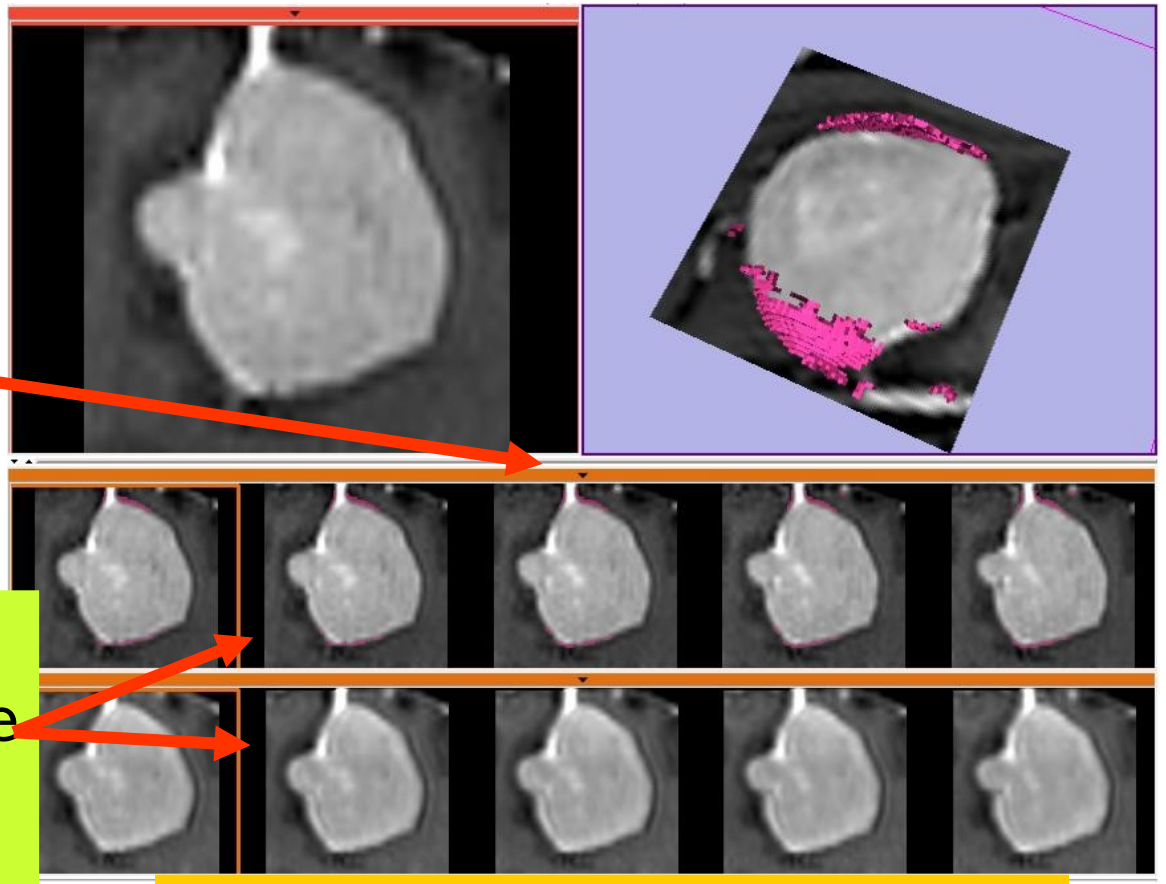


Define sensitivity of analysis

Show change in volume:
purple = growth
green = shrinkage

Show Change in Pathology

You can hide slice controls to have more screen space for viewing the images

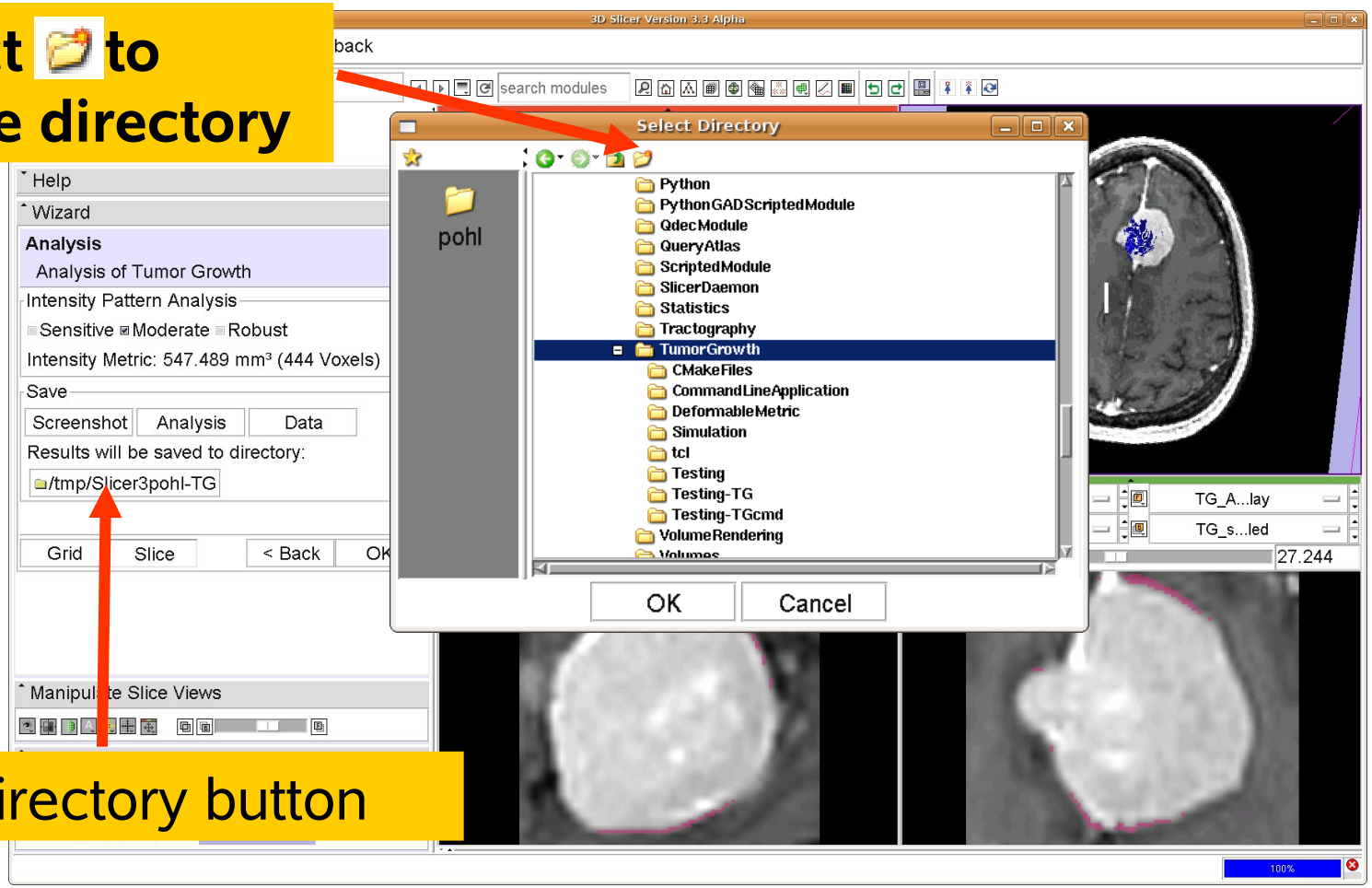


Lightbox View shows 5 consecutive slices for the ROI in the first scan (top), and in the second scan (bottom)

You can switch to **Four-up** layout or other layout at any time.

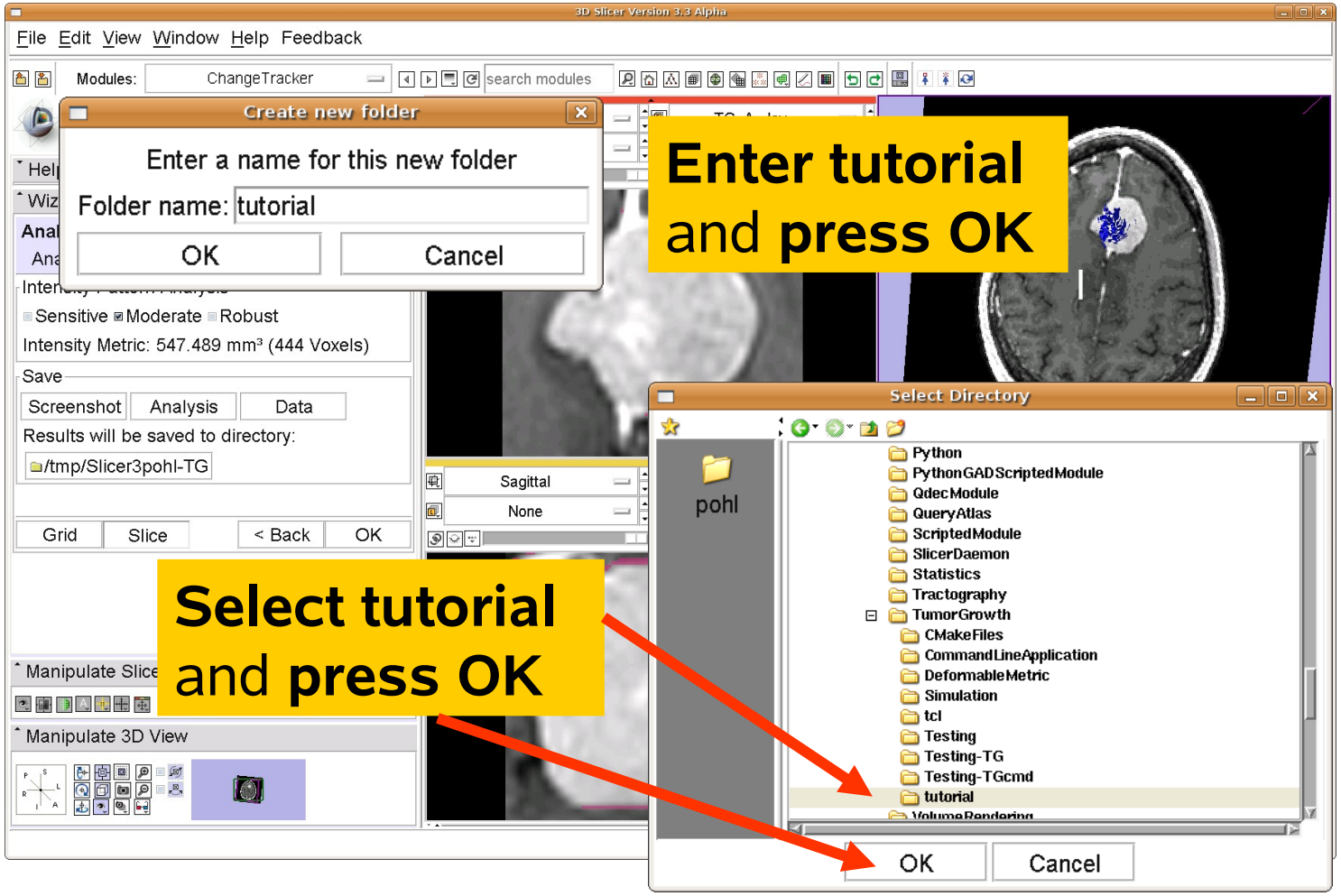
Define Working Directory

Select to create directory

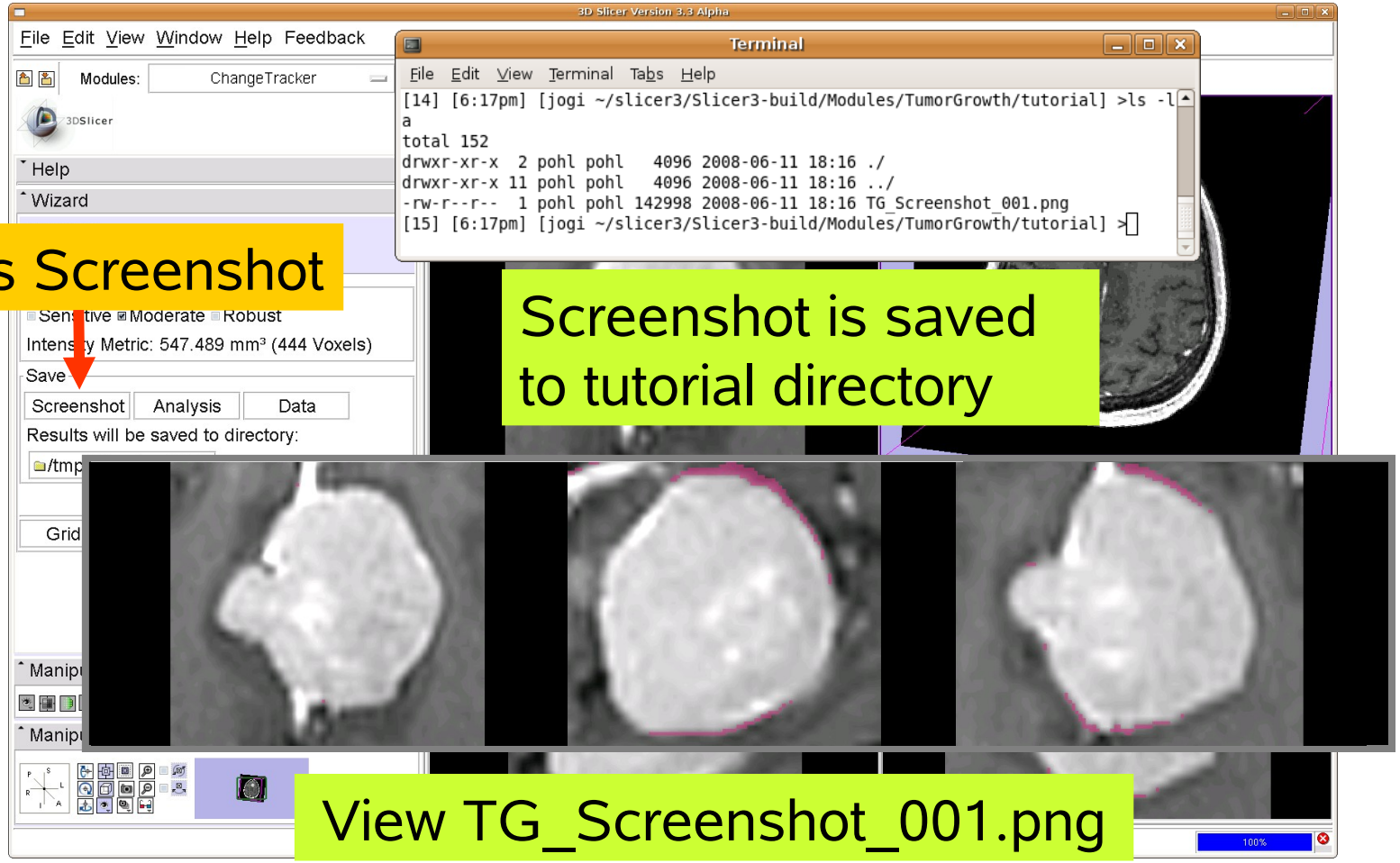


Press directory button

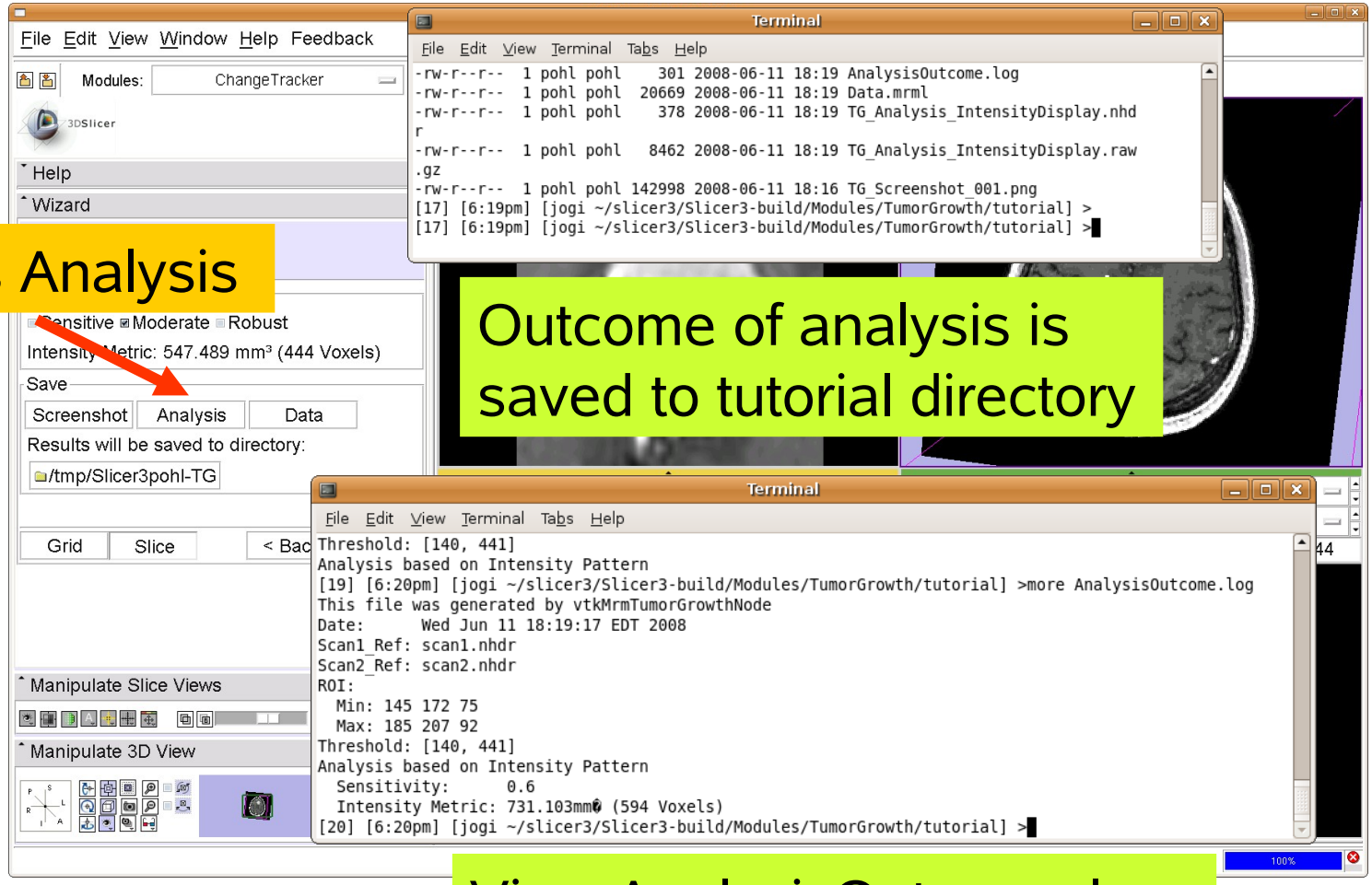
Define Working Directory



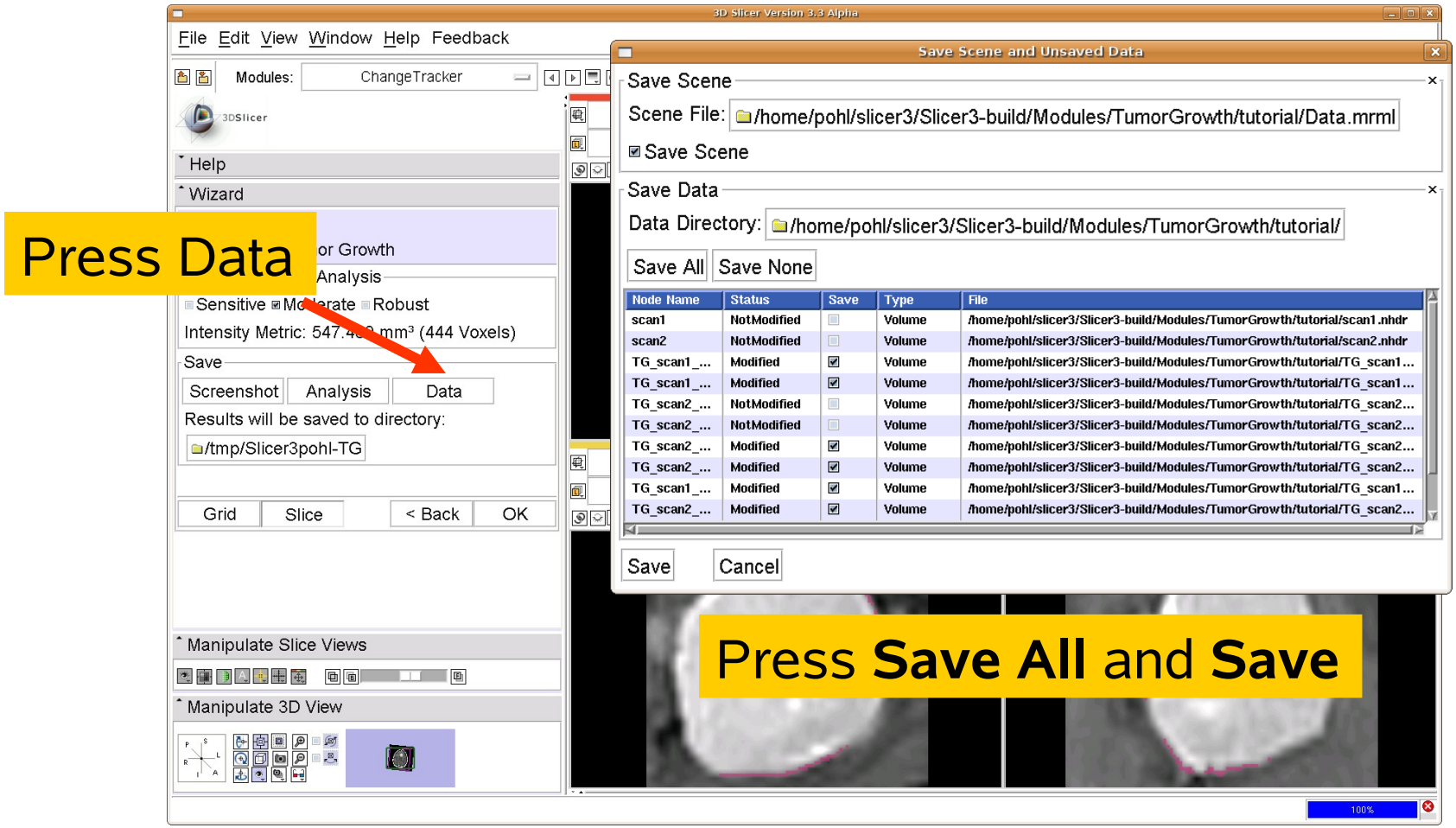
Save Screenshot



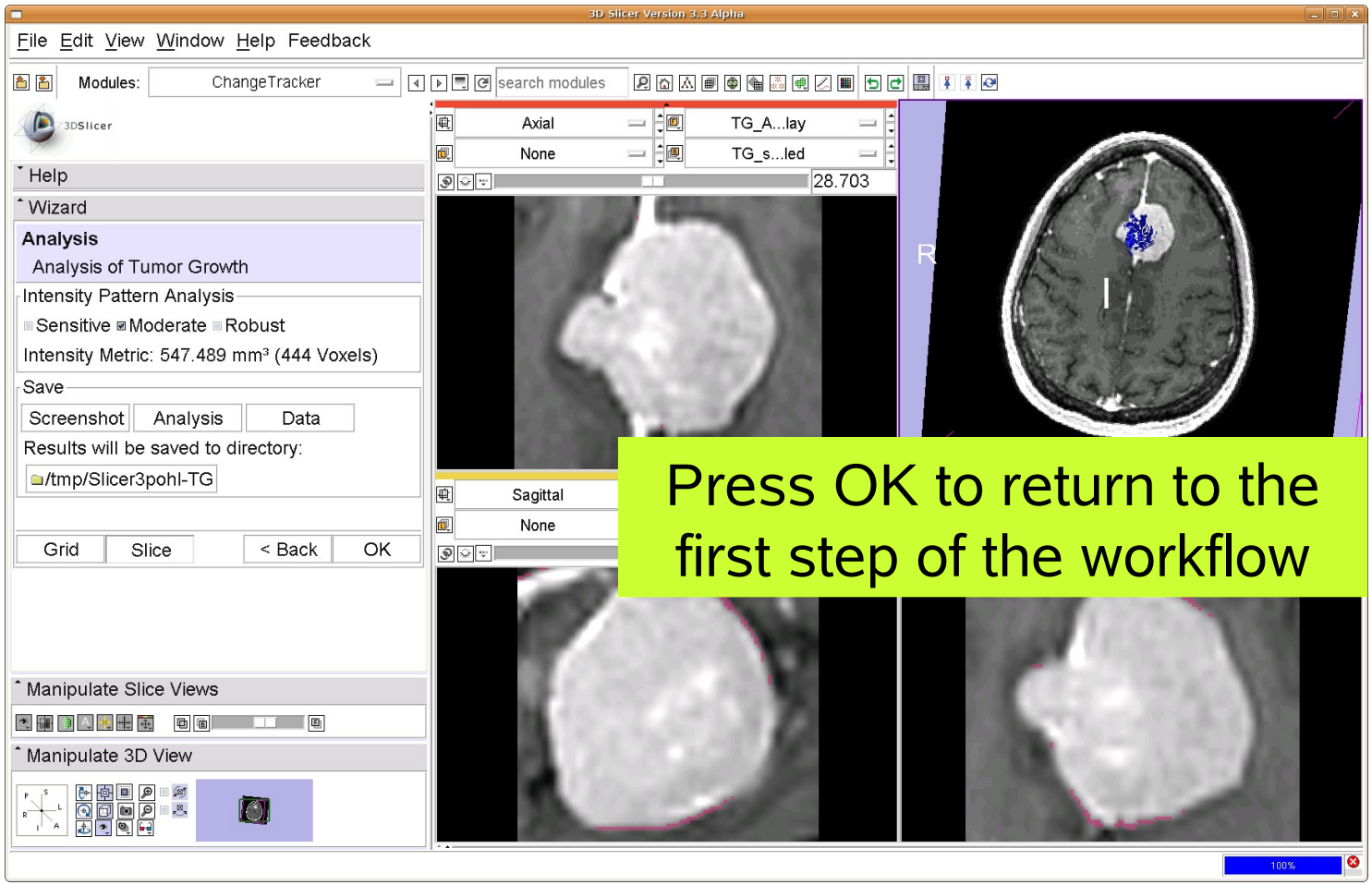
Save Analysis Result



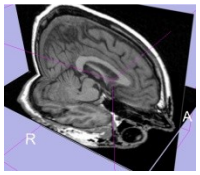
Save Intermediate Results



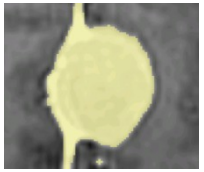
Finish Workflow



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

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add nac, ncigt and spl acknowledgements

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