# Tutorial: Isocenter shifting imageguidance in SlicerRT

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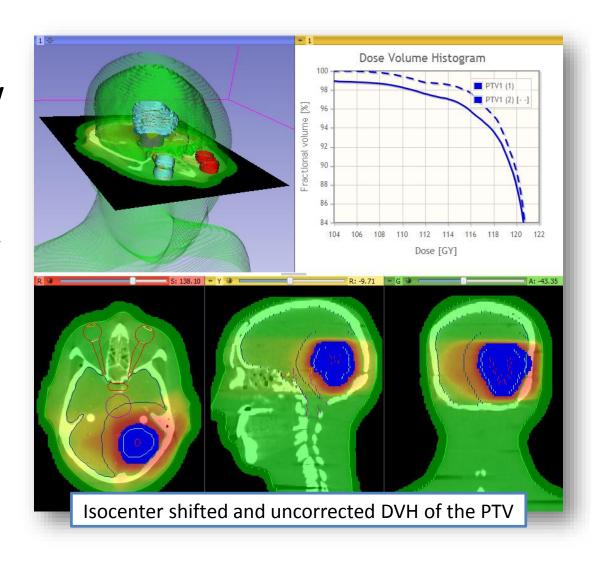




### Learning objective

This tutorial demonstrates how to perform a radiation therapy research workflow using the SlicerRT toolkit:

Isocenter shifting image-guidance









#### **Material**

- Connect to tutorial wifi: SSID: SlicerRT1, Password: tutorial (Disregard popup warning about no internet)
- Access download page:
   Enter in your web browser: 130.15.7.247
- Follow the instructions
- Supported platforms:
  - Windows, Mac OSX, 🐧 Linux
  - 32-bit is not supported!







#### **Overview**

- 1. Install SlicerRT extension
- 2. Load data from DICOM and nrrd files
- 3. Perform rigid registration on CT images
- 4. Transform day 2 dose volume
- 5. Accumulate dose distributions
- 6. Compute dose volume histogram
- +1. Create isodose lines and surfaces
- +2. Compare dose distributions using gamma







# 1/1. Install 3D Slicer

 Find the downloaded 3D Slicer package on your computer

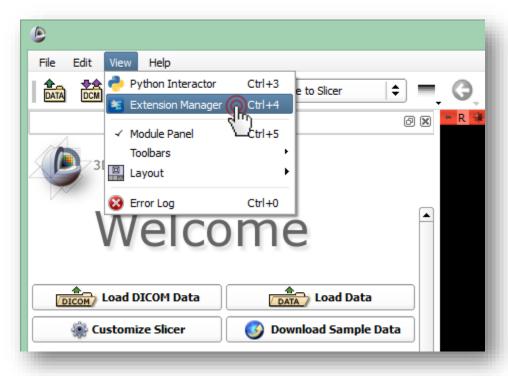
- Follow the usual steps to install an application
  - Different for each operating system







### 1/2. Install SlicerRT extension





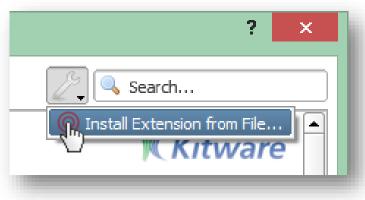








# 1/3. Install SlicerRT extension





Find the downloaded SlicerRT zip file on your computer



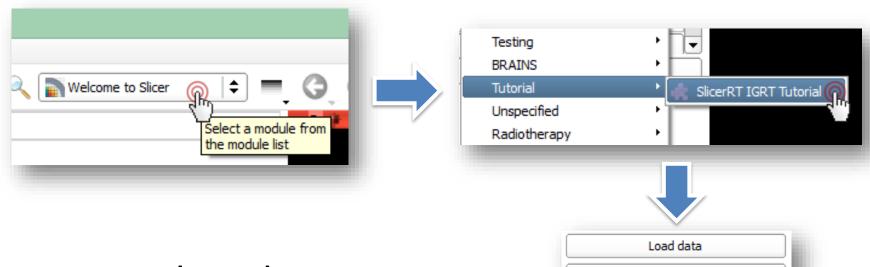




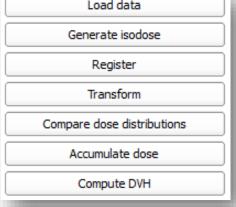


### In case of problems

If you fall behind, or encounter an error you can keep up using the tutorial module



Select the steps you want to automatically perform to catch up









# 2/1. Unpack tutorial datasets

 Find the dataset you downloaded named SlicerRT\_WorldCongress\_TutorialIGRT\_Dataset.zip

- Unpack it to a local folder of your choice
  - Different for each operating system



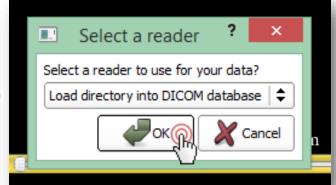




# 2/2. Import planning DICOM data

Drag&drop folder named EclipseEntPhantomRtData onto Slicer



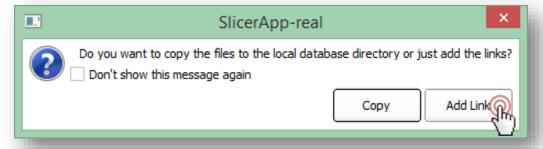


You'll be prompted for database folder here









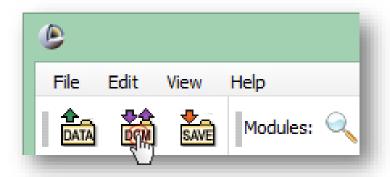






#### **Note**

If not importing via drag&drop, DICOM data can be imported and loaded from the DICOM browser that can be opened from the toolbar

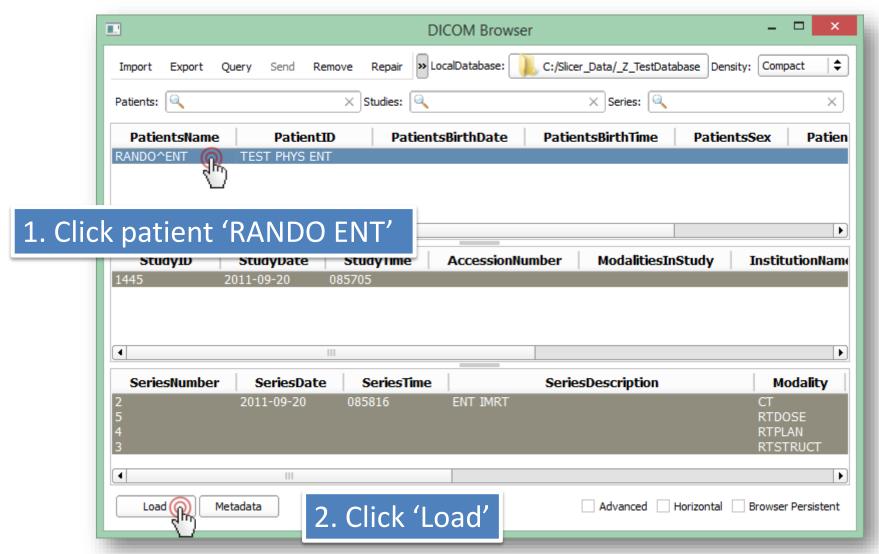








# 2/3. Load planning study

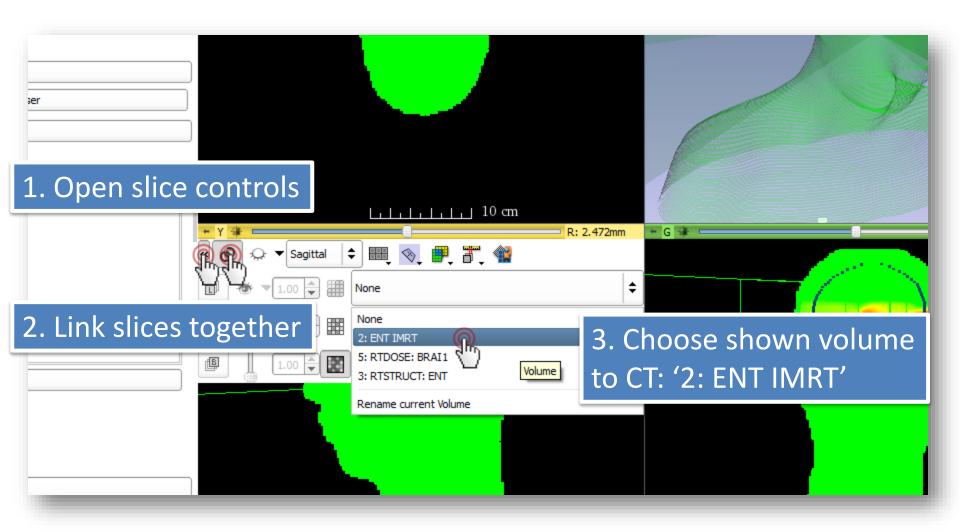








# 2/4. Change shown volumes

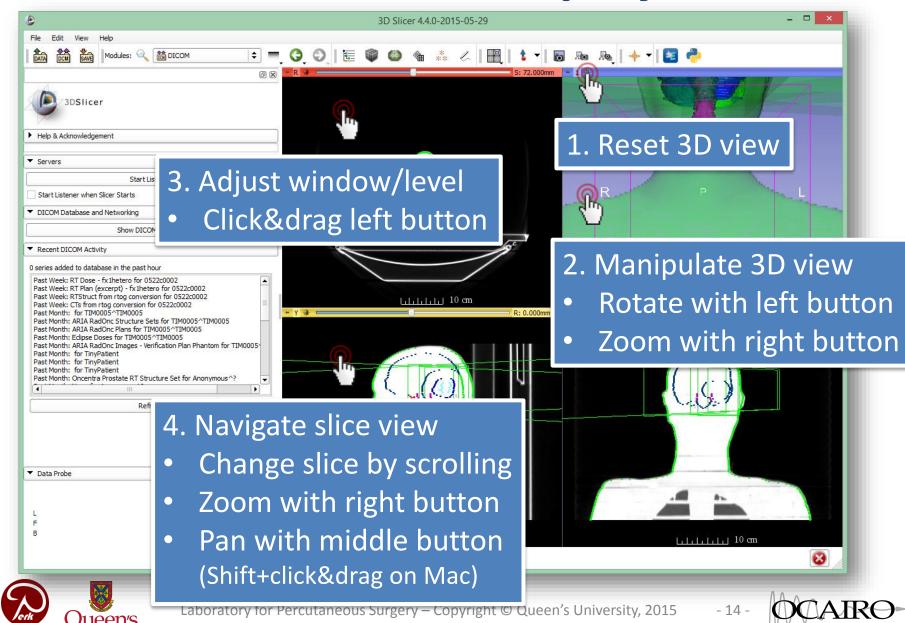




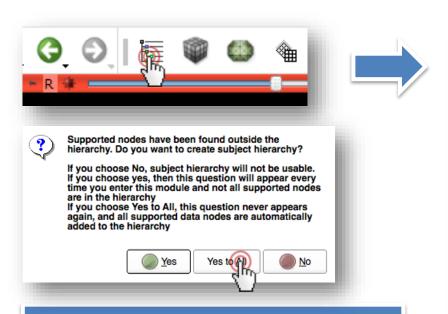




# 2/5. Tweak display

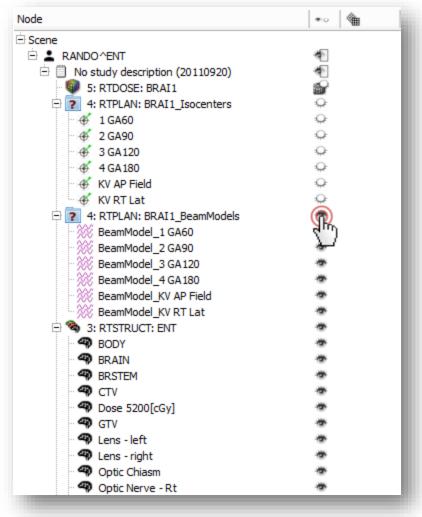


# 2/6. Explore loaded data



#### Subject hierarchy:

- Explore data in tree view
- Show/hide branches clicking the eye buttons
- Access options by right click



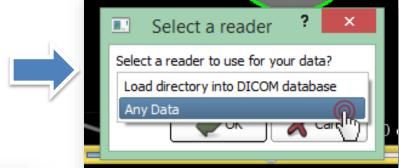


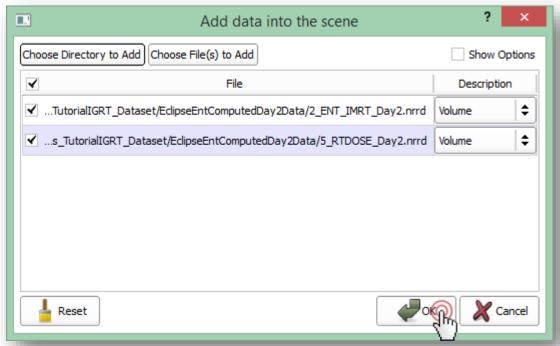


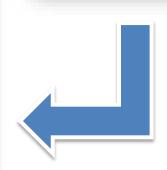


# 2/7. Load day 2 data

#### Drag&drop folder named EclipseEntComputedDay2Data onto Slicer













#### **Note**

- Data type selection dialog does not appear if you drag&drop files rather than folders, as files will be handled as non-DICOM
- Non-DICOM data can be also loaded in the dialog that appears after clicking



Data can be saved using

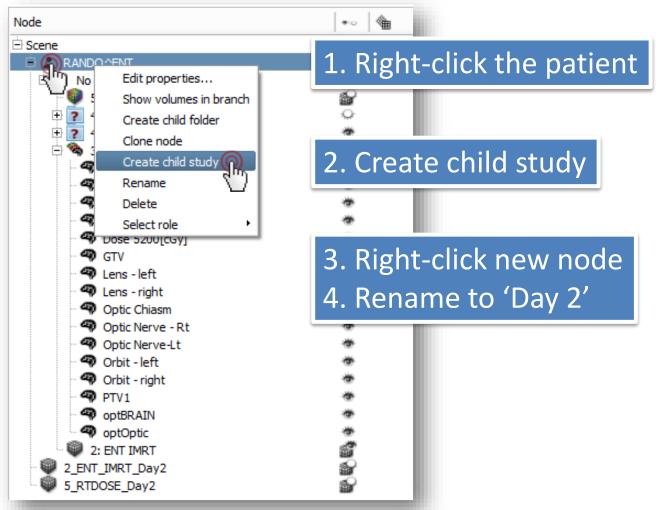








# 2/8. Add day 2 non-DICOM data to subject hierarchy



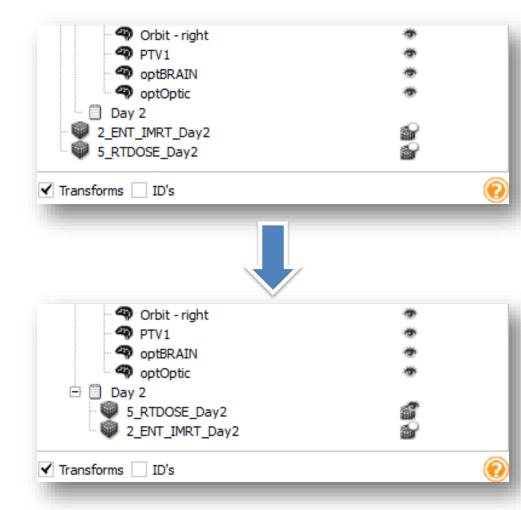






# 2/9. Add day 2 non-DICOM data to subject hierarchy

- 1. Drag&drop '2\_ENT\_IMRT\_ Day2' on the study 'Day 2'
- 2. Do the same with'5\_RTDOSE-Day2'

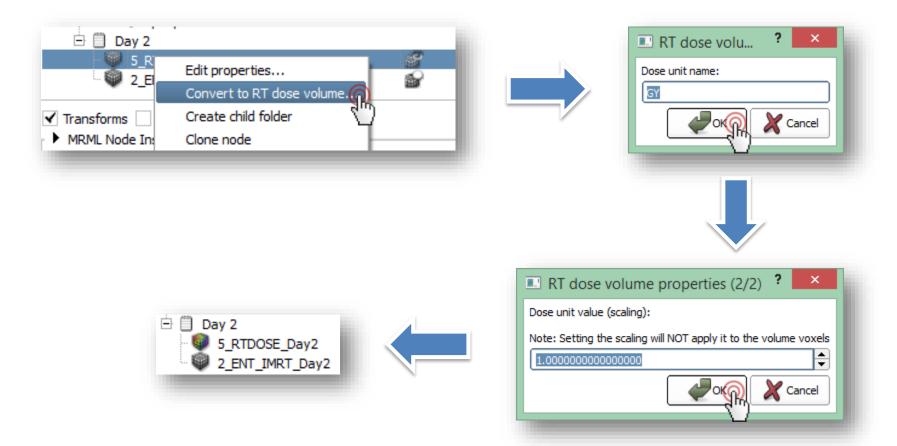








# 2/10. Convert day 2 dose volume actually a dose

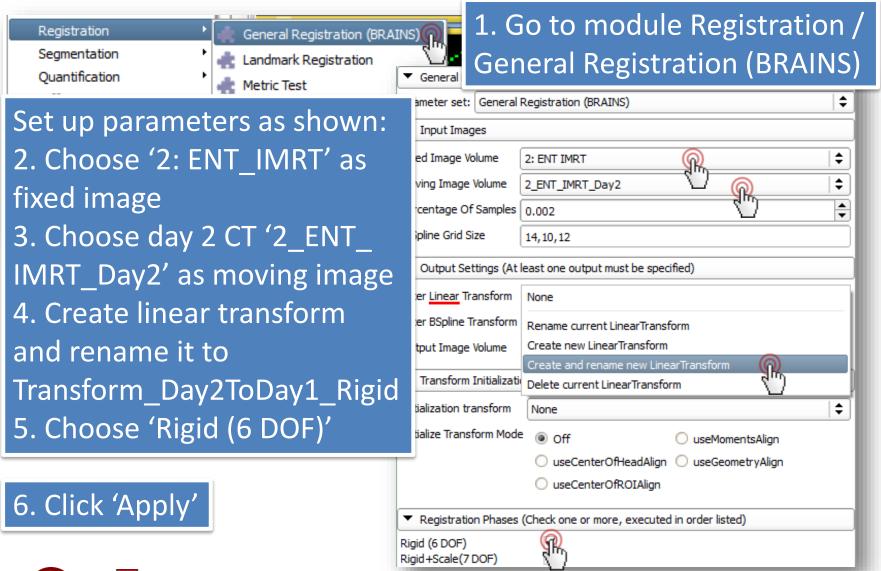








# 3/1. Register CT volumes



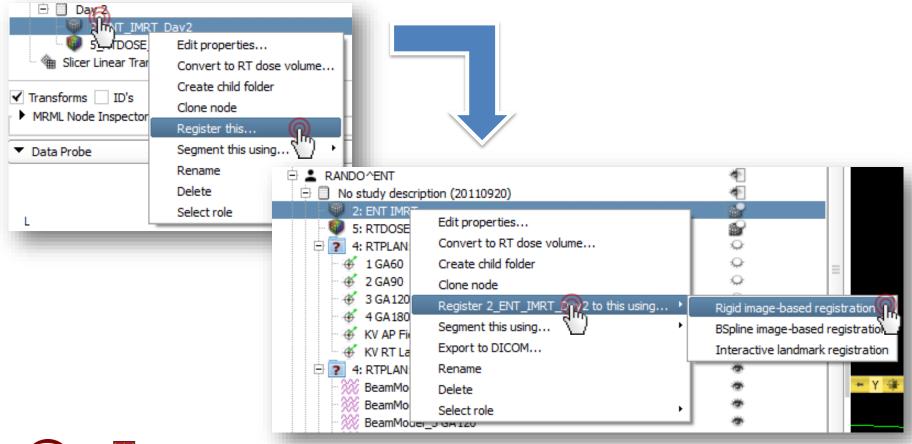






#### Note

You can also initiate registration from subject hierarchy (alternative way for previous step):

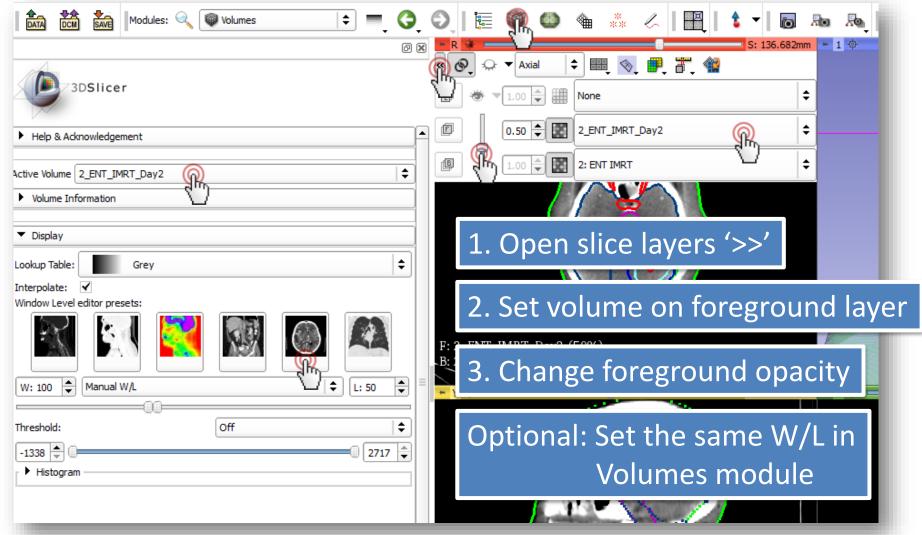








# 3/2. Explore volume differences

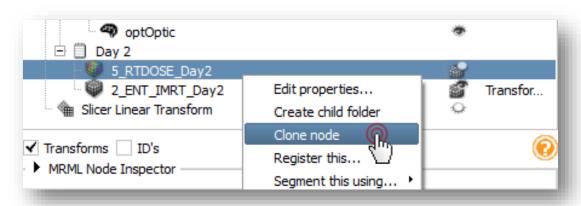






# 4/1. Clone dose volume

To be able to compare the non-registered (= uncorrected) and the registered (= isocenter shifted) results.



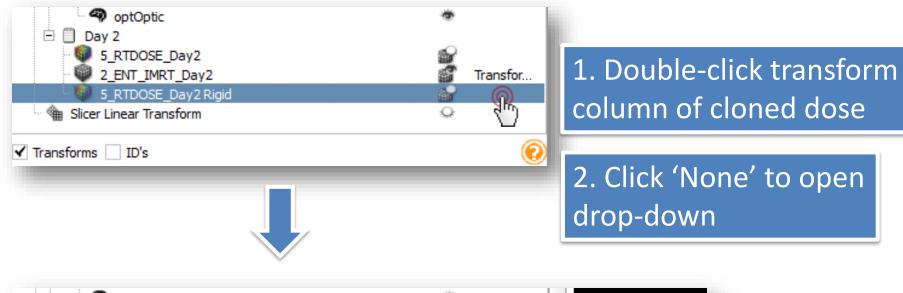
- 1. Switch to Subject hierarchy
- 2. Select 'Clone node' in context menu for day 2 dose
- 3. Rename it to '5\_RTDOSE\_Day2 Rigid'

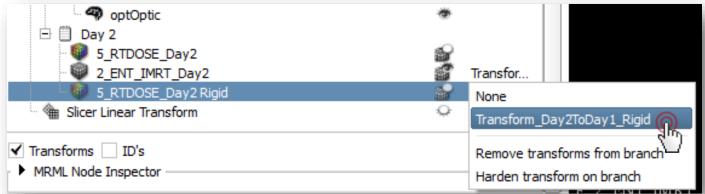






### 4/2. Transform cloned dose volume





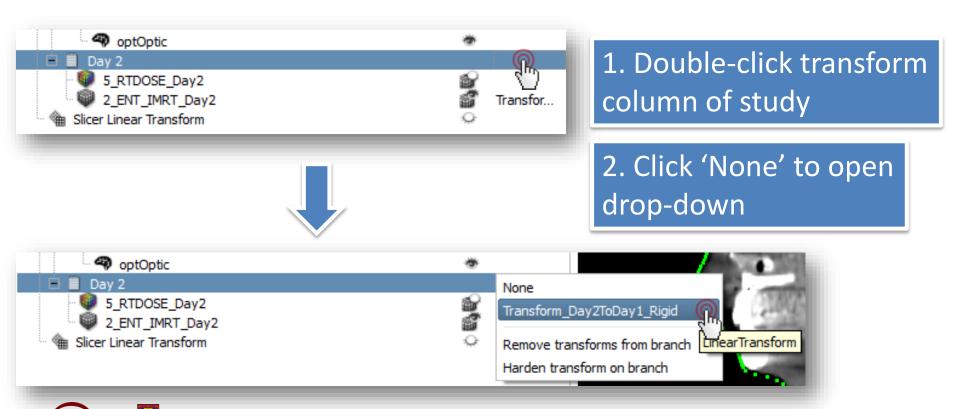






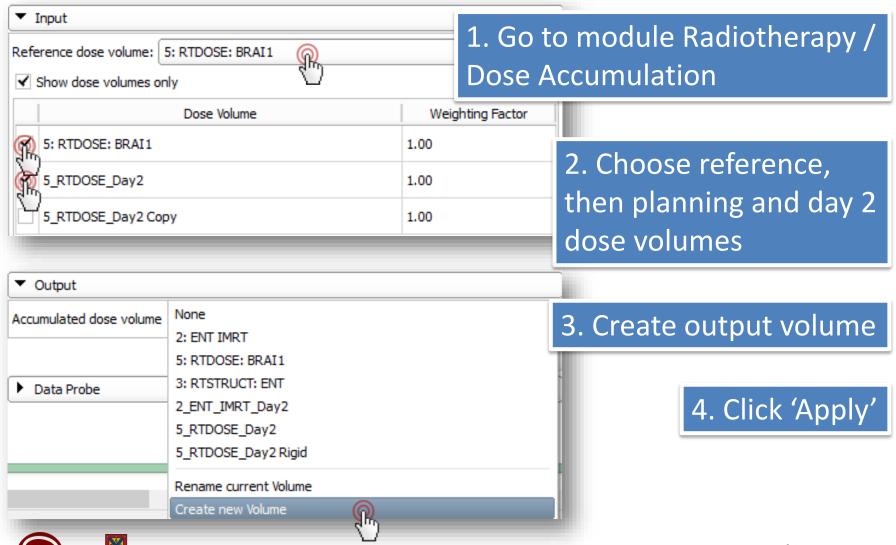
#### **Note**

We can transform the whole study too if we want to transform multiple objects (an alternative way for previous step):



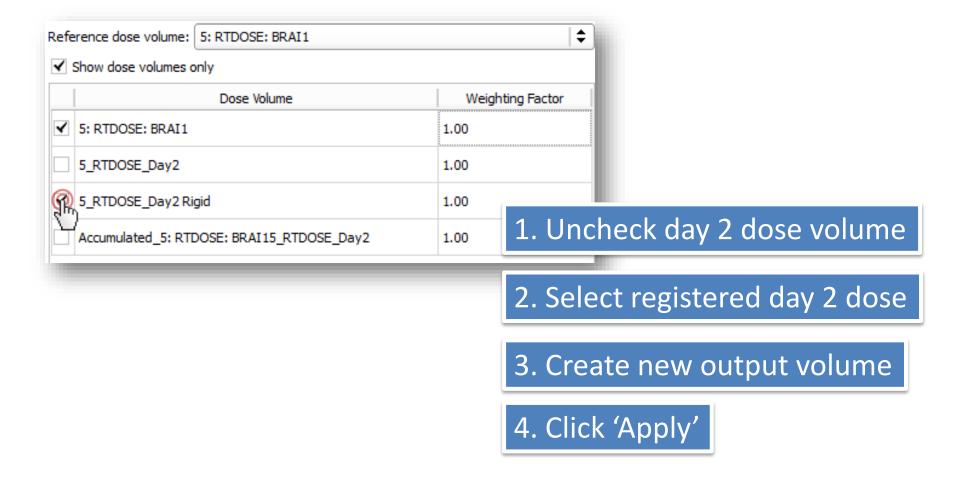


# 5/1. Accumulate dose distributions





# 5/2. Accumulate dose distributions

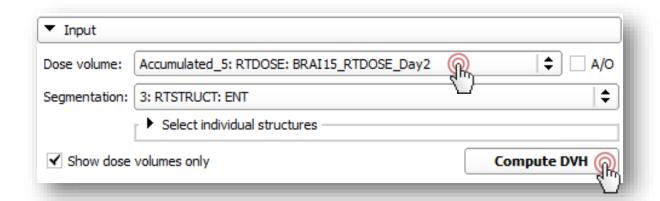








# 6/1. Compute DVH for unregistered



- 1. Go to module Radiotherapy / Dose Volume Histogram
- 2. Choose unregistered accumulated dose
- 3. Choose '3: RTSTRUCT: ENT'

Optional: Choose individual structures to speed up computation

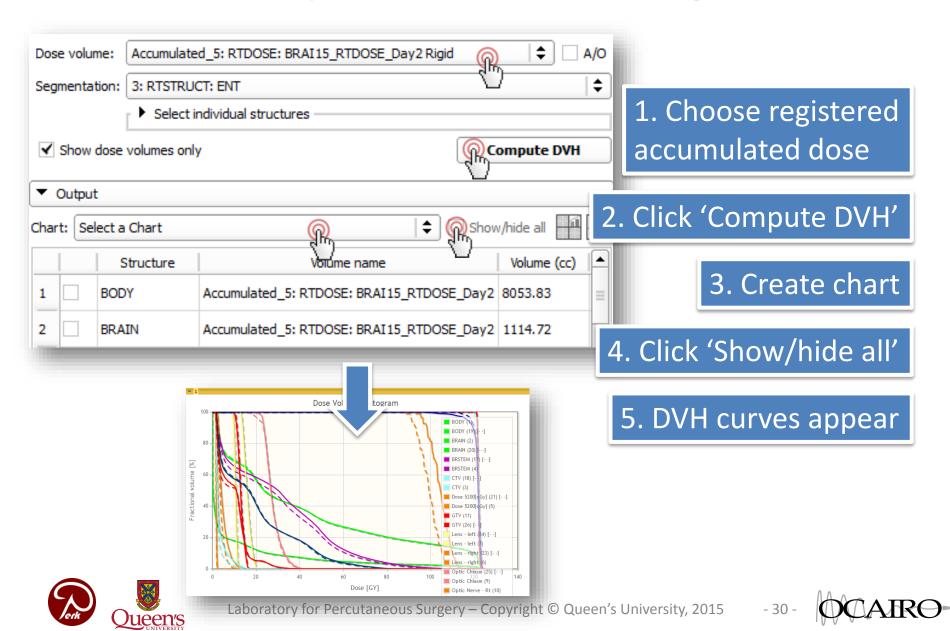
4. Click 'Compute DVH'



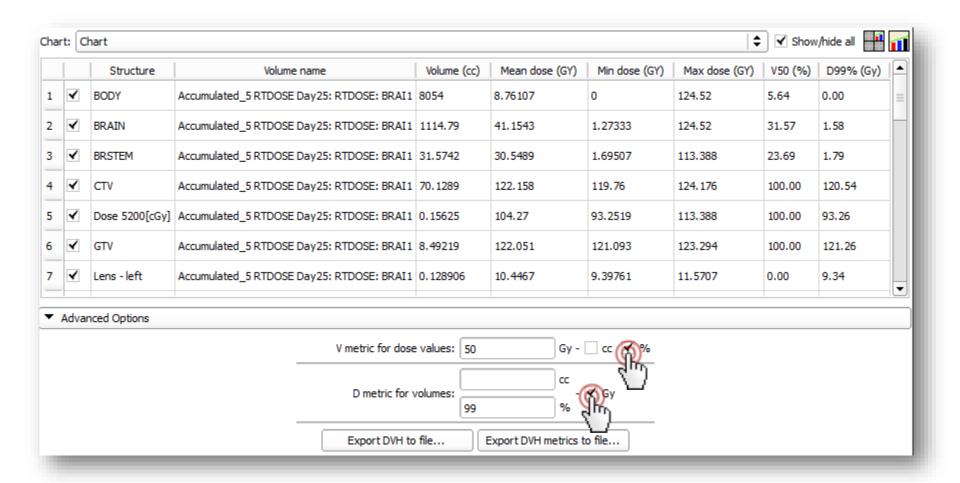




# 6/2. Compute DVH for registered



# 6/3. Quantify improvement

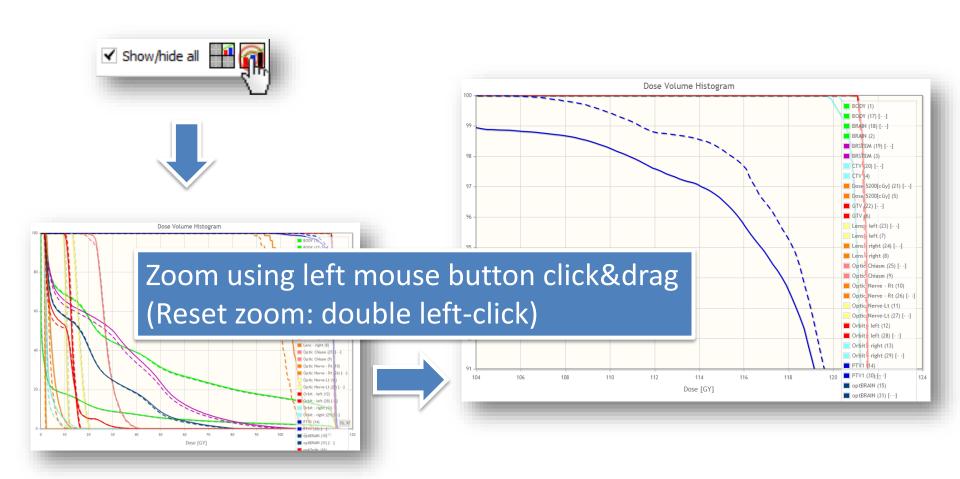








# 6/4. Visualize improvement

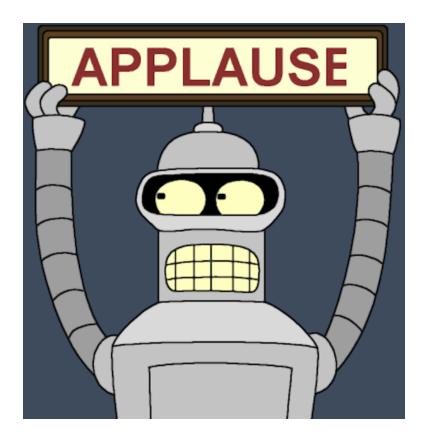








# Congratulations!



# Thanks for attending!







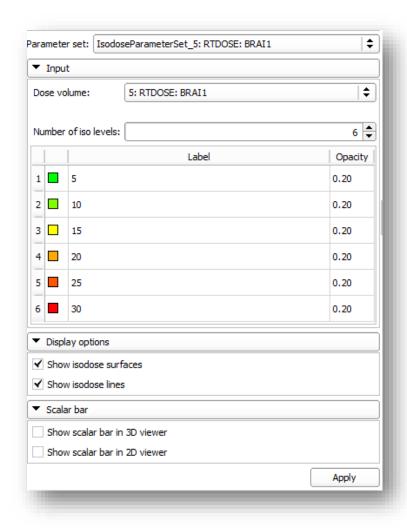
# **Appendix: Optional steps for IGRT**







# **Optional 1/1. Isodose lines/surfaces**



1. Choose Radiotherapy / Isodose module

3. Select '5: RTDOSE: BRAI1' volume as Dose volume

2. Click Apply

3. Select '5\_RTDOSE\_Day2' volume as Dose volume

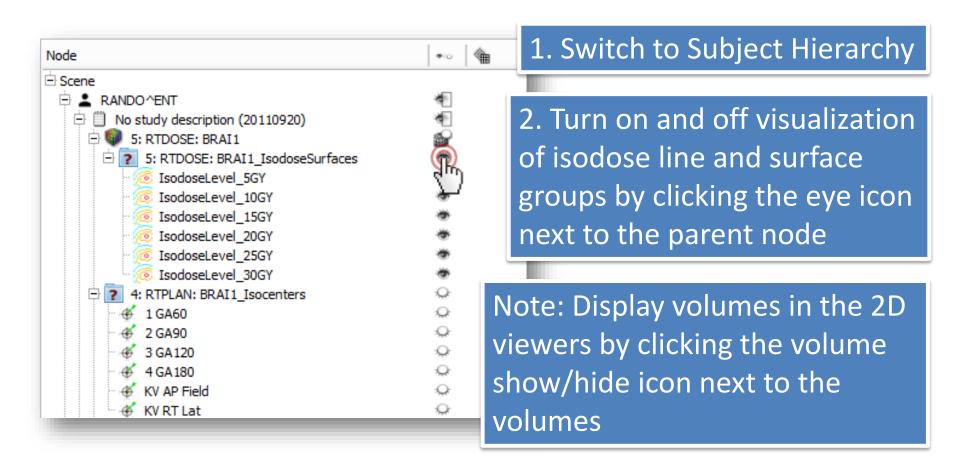
4. Click Apply







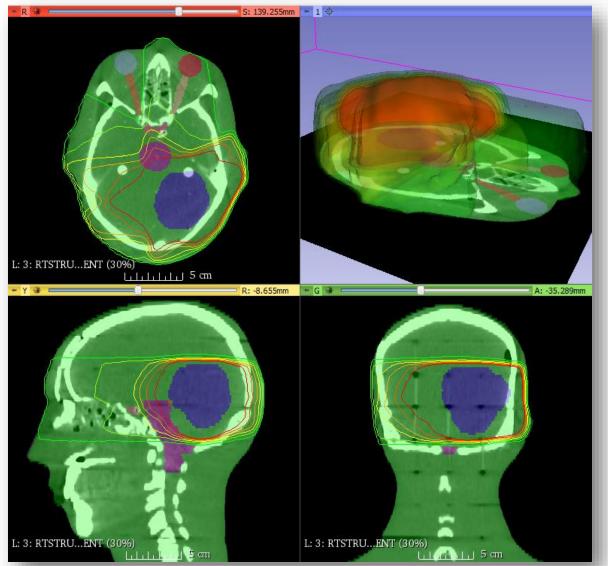
# Optional 1/2. Visualize isodose lines







# Optional 1/3. Isodose lines/surfaces

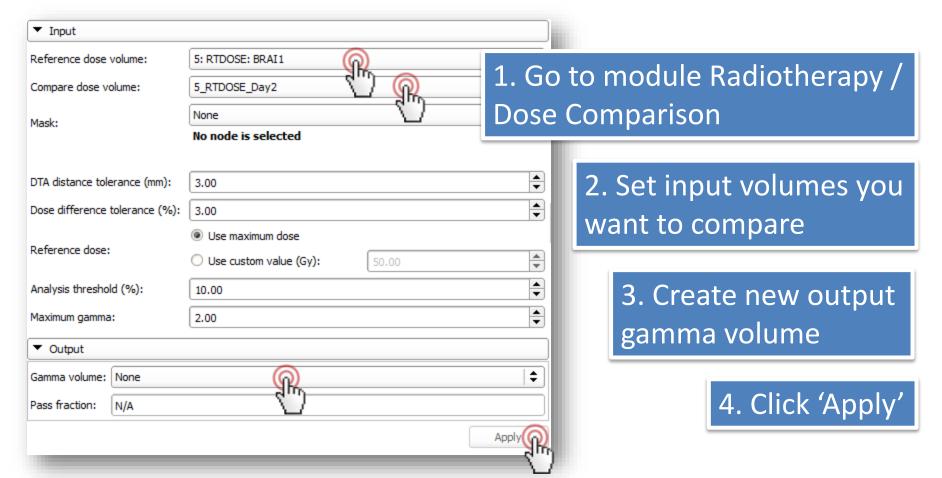








# Optional 2/1. Compare dose volumes using gamma comparison

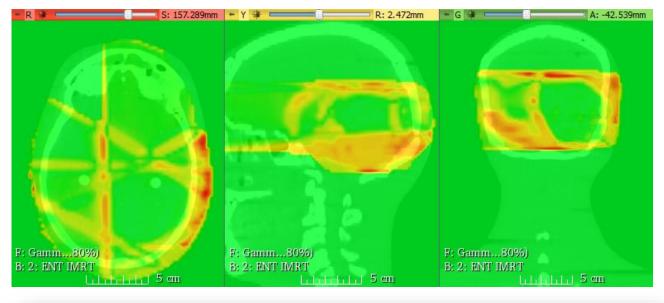






# **Optional 2/2. Evaluate improvement**

Plan dose vs Uncorrected Day 2 dose



Plan dose vs Corrected Day 2 dose







