

Tutorial

on SlicerRT and 3D Slicer modules



Agenda

- Acquire and build SlicerRT (on an existing Slicer)
- Use python console in Slicer
- Create extension and module skeletons
 - Use extension template
 - Use scripted module template
- Use SlicerRT
- Write a simple scripted module
- Write a simple CLI module



Acquire and build SlicerRT

- We must have a 3D Slicer built in Release mode before starting
- Get SlicerRT from the repository
 - `mkdir SlicerRT`
 - `svn co https://subversion.assembla.com/svn/slicerrt/trunk/SlicerRt/src SlicerRT`
- Configure SlicerRT
 - `mkdir SlicerRT_R64-bin`
 - `cd SlicerRT_R64-bin`
 - `../cmake-2.8.10.2-Linux-i386/cmake -DCMAKE_CXX_COMPILER=/usr/bin/g++-4.6 -DCMAKE_BUILD_TYPE=Release -DSlicer_DIR=/home/slicer/Support/Slicer4-SuperBuild-Debug/Slicer-build ../SlicerRT`
- Build SlicerRT
 - `make`



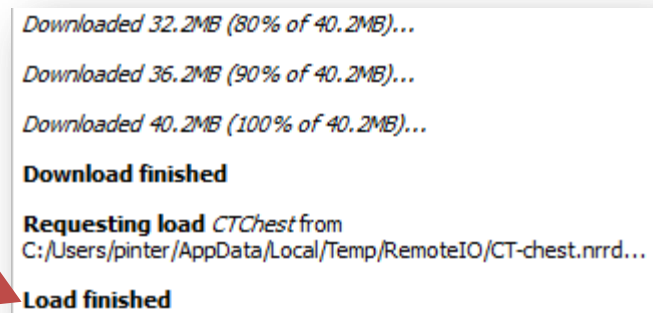
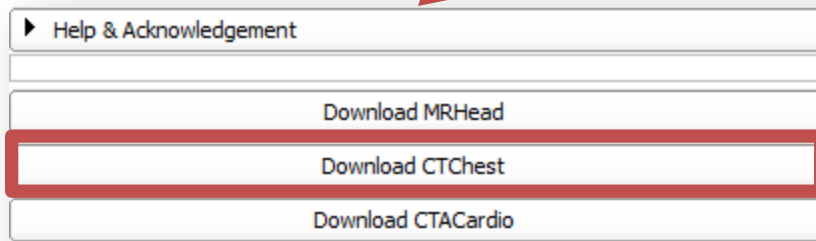
Next item on agenda

- Acquire and build SlicerRT (on an existing Slicer)
- **Use python console in Slicer**

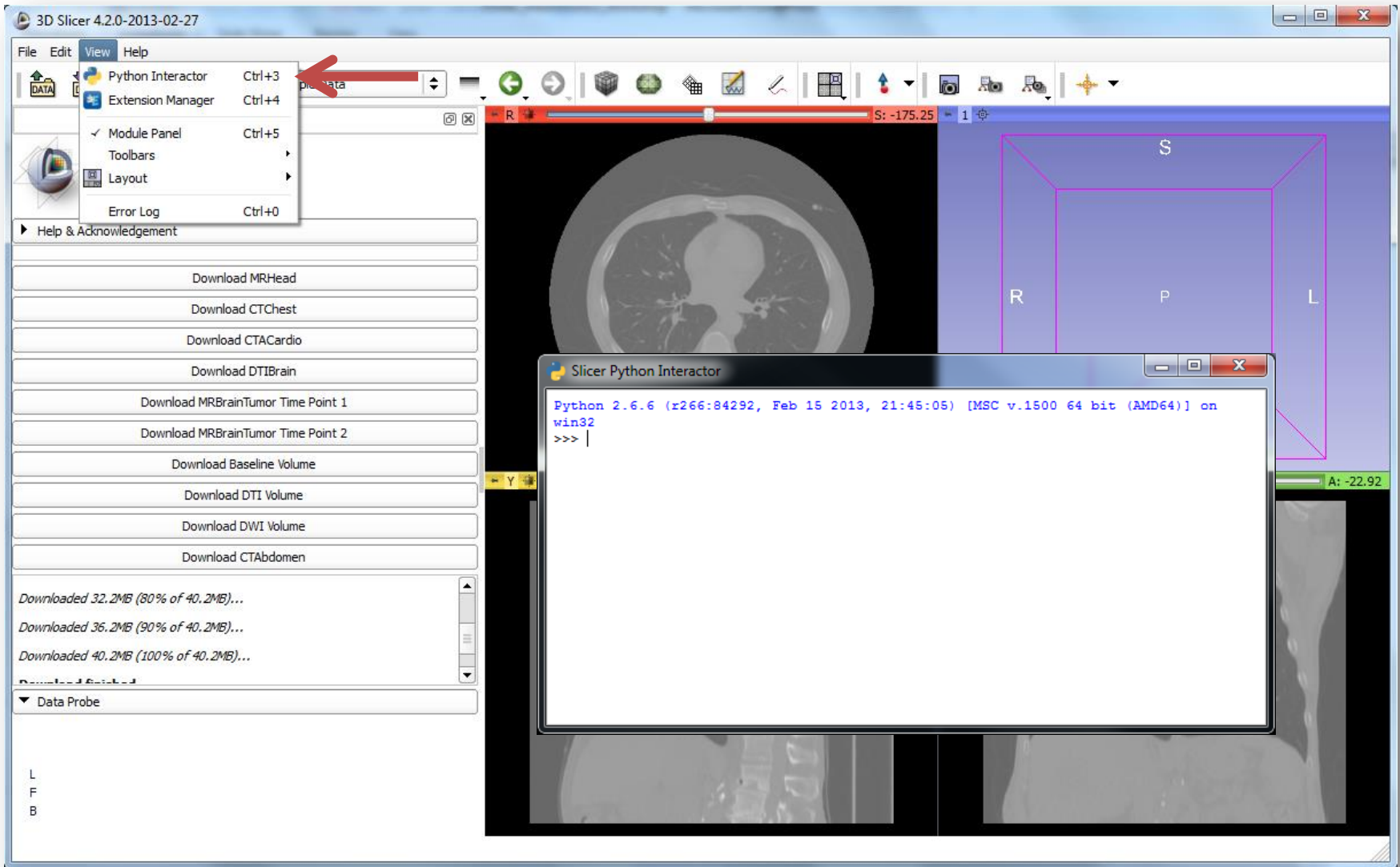


Launch Slicer and load data

- `cd ~/Support/Slicer4-SuperBuild-Debug/Slicer-build`
- `./Slicer`



Introducing the python console



Accessing the MRML scene and nodes

- Accessing MRML scene directly

- `c = slicer.mrmlScene.GetNodesByName('CTChest')`
- `c.GetNumberOfItems()`
- `v = c.GetItemAsObject(0)`

Also useful: `slicer.mrmlScene.GetNodeByID()`

- Using utility functions

- `v = slicer.util.getNode('CTChest')`

OR

- `v = slicer.util.getNode('CT*')`

Also useful (returns a list!): `slicer.util.getNodes('CT*')`



Manipulating MRML objects

- Setting window/level values programmatically
 - `d = v.GetDisplayNode()`
 - `d.SetAutoWindowLevel(0)`
 - `d.SetWindowLevel(350, 40)`
- Setting color palette programmatically
 - `c = slicer.util.getNode('Ocean')`
 - `d.SetAndObserveColorNodeID(c.GetID())`
- Slicer classes reference

<http://www.slicer.org/doc/html/classes.html>



Next item on agenda

- Acquire and build SlicerRT (on an existing Slicer)
- Use python console in Slicer
- **Create extension and module skeletons**
 - Use extension template
 - Use scripted module template



Create extension skeleton

- Setting window/level values programmatically
 - `cd ~/Support`
 - `python`
`~/Support/Slicer/Utilities/Scripts/ModuleWizard.py`
`--template`
`~/Support/Slicer/Extensions/Testing/SuperBuildExtensionTemplate --target ./MyExtension MyExtension`
 - `cd MyExtension`
 - `python`
`~/Support/Slicer/Utilities/Scripts/ModuleWizard.py`
`--template`
`~/Support/Slicer/Extensions/Testing/ScriptedLoadableExtensionTemplate/ScriptedLoadableModuleTemplate --target ./MyScriptedModule MyScriptedModule`



Create extension skeleton cont'd #1

- Edit super-build Cmake file
 - `gedit CMakeLists &`
Add under the other `add_subdirectory...` lines
 - `add_subdirectory(MyScriptedModule)`
 - Quit editor
 - `cd ..`
 - `mkdir MyExtension_R64-bin`
 - `cd MyExtension_R64-bin`



Create extension skeleton cont'd #2

- Configure extension skeleton

- ```
../cmake-2.8.10.2-Linux-i386/cmake
-DCMAKE_CXX_COMPILER=/usr/bin/g++-4.6 --
DSlicer_DIR=/home/slicer/Support/Slicer4-
SuperBuild-Debug/Slicer-build -
DMyExtension_SUPERBUILD=0 ../MyExtension
```

- Build extension skeleton

- ```
make
```



Add new modules to the paths

- Start Slicer
- Add modules to the additional module paths
 - Go to *Edit / Application settings / Modules*
 - Click the “>>” button next to the *Additional module paths* section
 - Add the following directories:
 - *~/Support/MyExtension_R64-bin/lib/Slicer-4.2/qt-loadable-modules*
 - same location but the directory *qt-scripted-module*



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- Use python console in Slicer
- Create extension and module skeletons
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 - Use scripted module template
- **Use SlicerRT**





Add SlicerRT to the paths

- Start Slicer
- Add SlicerRT to the additional module paths
 - Go to *Edit / Application settings / Modules*
 - Click the “>>” button next to the *Additional module paths* section
 - Add all sub-directories from:
 - */home/slicer/Support/SlicerRT_R64-bin/inner-build/lib/Slicer-4.2*
 - Press *Restart*



Load RT data

- Go to *module list / Testing / TestCases / SlicerRT Demo RSNA2012 Self Test*
- Click *Load data*
- Click 
(This step is not necessary, it just shows the DICOM browser which has to be used when loading one's own data, not downloaded automatically from script)
- When the data is loaded, save the scene for faster future loading using 

Registering CT volumes

- Go to *module list / Registration / General Registration (BRAINS)*
- Set the module parameters as seen in this image and click *Apply*

▼ Input Images

Fixed Image Volume 2: ENT IMRT

Moving Image Volume 2_ENT_IMRT_Day2

▼ Output Settings (At least one output must be specified.)

Slicer BSpline Transform None

Slicer Linear Transform Day1ToDay2RigidTransform

Output Image Volume None

▼ Initialization of registration

Initialization transform None

Intitialize Transform Mode Off

useMomentsAlign

useCenterOfHeadAlign

useGeometryAlign

useCenterOfROIAlign

▼ Registration Phases (Check one or more, executed in order listed)

Rigid (6 DOF)

Select *Create and rename...*

Resampling the day 2 dose volume

- Go to *module list / Registration / Resample Image (BRAINS)*
- Set the module parameters as seen in this image and click *Apply*

The screenshot shows the configuration interface for the 'Resample Image' module. It is divided into three main sections: Inputs, Outputs, and Warping Parameters. Red arrows point to the following fields:

- Inputs:**
 - Image To Warp: 5_RTDOSE_Day2
 - Reference Image: 5_RTDOSE_Day2
- Outputs:**
 - Output Image: 5_RTDOSE_Day2_Resampled_Rigid
 - Pixel Type: float (selected)
- Warping Parameters:**
 - Displacement Field (deprecated): None
 - Transform file: Day1ToDay2RigidTransform

Accumulating doses

- Go to *module list / Radiotherapy / Dose Accumulation*
- Un-check *Show dose volumes only* checkbox
- Select reference dose volume *5: RTDOSE*
- Check volumes
 - 5: RTDOSE
 - 5_RTDOSE_Day2
- Open combobox in the *Output* section
- Select *Create new Volume*
- Click *Apply*
- Accumulate these volumes in the same way
 - 5: RTDOSE
 - 5_RTDOSE_Day2_Resampled_Rigid

Computing dose volume histograms

- Go to *Radiotherapy / Dose Volume Histogram*
- Select the first accumulated dose volume from the list
- Select *PTV* as the structure set
- Click *Compute DVH*
- Select second accumulated dose volume
- Compute DVH
- Do it for other structures, such as *BRSTEM* or *optOptic*
- Click *Select Chart*
- Click *Create new Chart*
- Check *Show/hide all*



Next item on agenda

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- Use python console in Slicer
- Create extension and module skeletons
 - Use extension template
 - Use scripted module template
- Use SlicerRT
- **Write a simple scripted module**



Write our scripted module #1

- Open *MyExtension/MyScriptedModule/MyScriptedModule.py*
- Rename the module and put it in our extension category

```
class MyScriptedModule:  
    def __init__(self, parent):  
        parent.title = "Center of Masses" | # T  
        parent.categories = ["MyExtension"]
```

- Create widgets to specify inputs:

Under `def setup(self):` look for *input volume selector*, and change

```
self.inputSelector.addAttribute( "vtkMRMLScalarVolumeNode", "LabelMap", 1 )  
self.inputSelector.selectNodeUponCreation = False
```

and

```
self.inputSelector.setToolTip( "Pick the first input to the algorithm." )  
parametersFormLayout.addRow("Input Volume 1: ", self.inputSelector)
```

- Create second input selector
 - Rename *outputSelector* to *input2Selector* everywhere
setUp and *onApplyButton* and *onSelect* functions
 - Set the same settings as for the first one



Write our scripted module #2

- Look for *class MyScriptedModuleLogic* on the bottom
- Insert this code above the *Run* function

```
def getCenterOfMass(self, volumeNode):
    centerOfMass = [0,0,0]

    numberOfStructureVoxels = 0
    sumX = sumY = sumZ = 0

    volume = volumeNode.GetImageData()

    for z in xrange(volume.GetExtent()[4], volume.GetExtent()[5]+1):
        for y in xrange(volume.GetExtent()[2], volume.GetExtent()[3]+1):
            for x in xrange(volume.GetExtent()[0], volume.GetExtent()[1]+1):
                voxelValue = volume.GetScalarComponentAsDouble(x,y,z,0)
                if voxelValue>0:
                    numberOfStructureVoxels = numberOfStructureVoxels+1
                    sumX = sumX + x
                    sumY = sumY + y
                    sumZ = sumZ + z

    if numberOfStructureVoxels > 0:
        centerOfMass[0] = sumX / numberOfStructureVoxels
        centerOfMass[1] = sumY / numberOfStructureVoxels
        centerOfMass[2] = sumZ / numberOfStructureVoxels

    print('Center of mass for \'' + volumeNode.GetName() + '\': ' + repr(centerOfMass))
    return centerOfMass
```



Write our scripted module #3

- Insert this code in the *Run* function above the *return* statement

```
def run(self, inputVolume, input2Volume):  
    """  
    Run the actual algorithm  
    """  
    center1 = self.getCenterOfMass(inputVolume)  
    center2 = self.getCenterOfMass(input2Volume)  
  
    self.translation = []  
    for i in [0,1,2]:  
        self.translation.append(center2[i] - center1[i])  
  
    return True
```

- Displaying the output

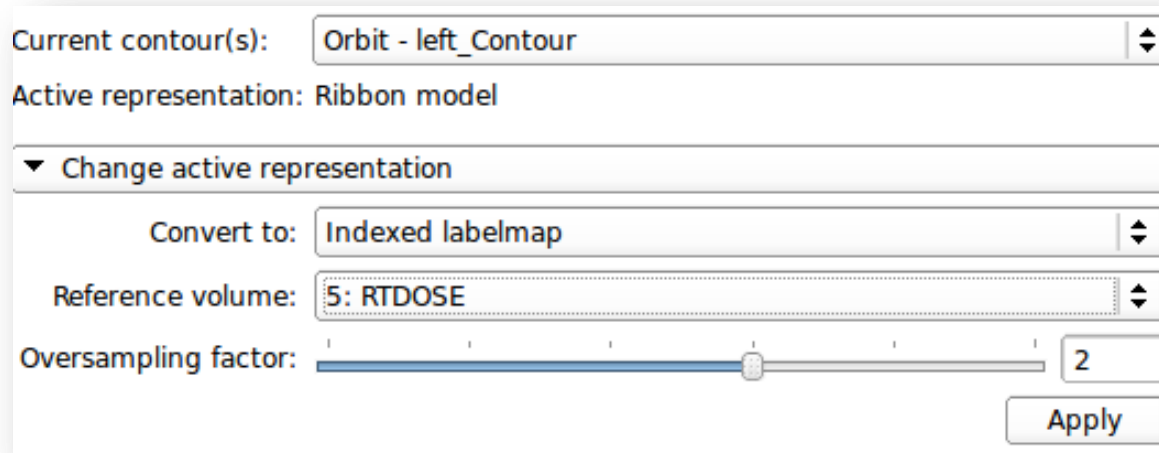
```
def onApplyButton(self):  
    logic = MyScriptedModuleLogic()  
    print("Run the algorithm")  
    logic.run(self.inputSelector.currentNode(), self.outputSelector.currentNode())  
    self.outputField.setText('(' + repr(logic.translation[0]) + ', ' + repr(  
(logic.translation[1]) + ', ' + repr(logic.translation[2]) + ')')
```

- Create the text field under the *Apply* button

```
self.outputLabel = qt.QLabel()  
parametersFormLayout.addRow(self.outputLabel)
```


Try our scripted module

- Build the extension (go to its bin directory and *make*)
- Load data using the demo module (load the scene we saved)
- Convert two structures, for example the two eyeballs (*Orbit...*) using the *Radiotherapy / Contours* module



- Go to our module in *MyExtension / Center of Masses*
- Set the two labelmaps as input
- Press *Apply*

Last item on agenda

- Acquire and build SlicerRT (on an existing Slicer)
- Use python console in Slicer
- Create extension and module skeletons
 - Use extension template
 - Use scripted module template
- Use SlicerRT
- Write a simple scripted module
- **Write a simple CLI module**



Write our CLI module #1

- Open *MyExtension/SuperCLIModuleTemplate/SuperCLIModuleTemplate.xml*
- Rename the module and put it in our extension category

```
<executable>  
  <category>MyExtension</category>  
  <title>Morphology</title>
```

Write our CLI module #2

- Create controls

```
<parameters>
  <string-enumeration>
    <name>operation</name>
    <longflag>operation</longflag>
    <label>Operation</label>
    <description><![CDATA[Type of operation]]></description>
    <default>Dilate</default>
    <element>Dilate</element>
    <element>Erode</element>
  </string-enumeration>
  <label>IO</label>
  <description><![CDATA[Input/output parameters]]></description>
  <image>
    <name>inputVolume</name>
    <label>Input Volume</label>
    <channel>input</channel>
    <index>0</index>
    <description><![CDATA[Input volume]]></description>
  </image>
  <image>
    <name>outputVolume</name>
    <label>Output Volume</label>
    <channel>output</channel>
    <index>1</index>
    <description><![CDATA[Output Volume]]></description>
  </image>
</parameters>
```

Write our CLI module #3

- Open *MyExtension/SuperCLIModuleTemplate/SuperCLIModuleTemplate.cxx*
- Include needed headers

```
#include "itkBinaryBallStructuringElement.h"  
#include "itkGrayscaleDilateImageFilter.h"  
#include "itkGrayscaleErodeImageFilter.h"
```

- Define used types (replace *FilterType* definition)

```
typedef itk::BinaryBallStructuringElement<  
    InputPixelType, 3> StructuringElementType;  
typedef itk::GrayscaleDilateImageFilter<  
    InputImageType, OutputImageType, StructuringElementType> DilateFilterType;  
typedef itk::GrayscaleErodeImageFilter<  
    InputImageType, OutputImageType, StructuringElementType> ErodeFilterType;
```

Write our CLI module #4

- Add processing code

```
reader->SetFileName( inputVolume.c_str() );

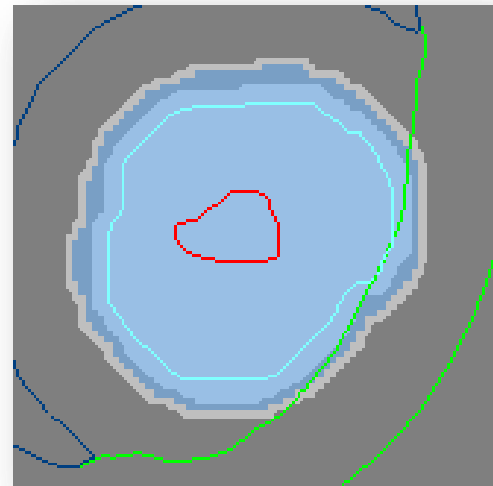
typename OutputImageType::Pointer outputImage = NULL;
if (operation == "Dilate")
{
    typename DilateFilterType::Pointer filter = DilateFilterType::New();
    filter->SetInput( reader->GetOutput() );
    filter->Update();
    outputImage = filter->GetOutput();
}
else if (operation == "Erode")
{
    typename ErodeFilterType::Pointer filter = ErodeFilterType::New();
    filter->SetInput( reader->GetOutput() );
    filter->Update();
    outputImage = filter->GetOutput();
}
else
{
    return EXIT_FAILURE;
}

typename WriterType::Pointer writer = WriterType::New();
writer->SetFileName( outputVolume.c_str() );
writer->SetInput( outputImage );
writer->SetUseCompression(1);
writer->Update();
```



Try our CLI module

- Load data using the demo module (load the scene we saved)
- Convert one structure, for example *PTV* using the *Contours* module
- Go to our module in *MyExtension / Morphology*
- Set the *PTV* labelmap as input
- Create output volume
- Press *Apply*
- Change *Operation to Erode*
- Create output volume
- Press *Apply*
- Show volumes in the viewers: access slice toolbar, expand and turn on lock (⏏), select output volumes and *PTV* labelmap



Congratulations!

Thanks for attending!

