



Paul Cézanne, *Moulin sur la Coulevre à Pontoise, 1881*,
Staatliche Museen zu Berlin, Nationalgalerie

Programming into Slicer3

Sonia Pujol, Ph.D.

Surgical Planning Laboratory
Harvard University





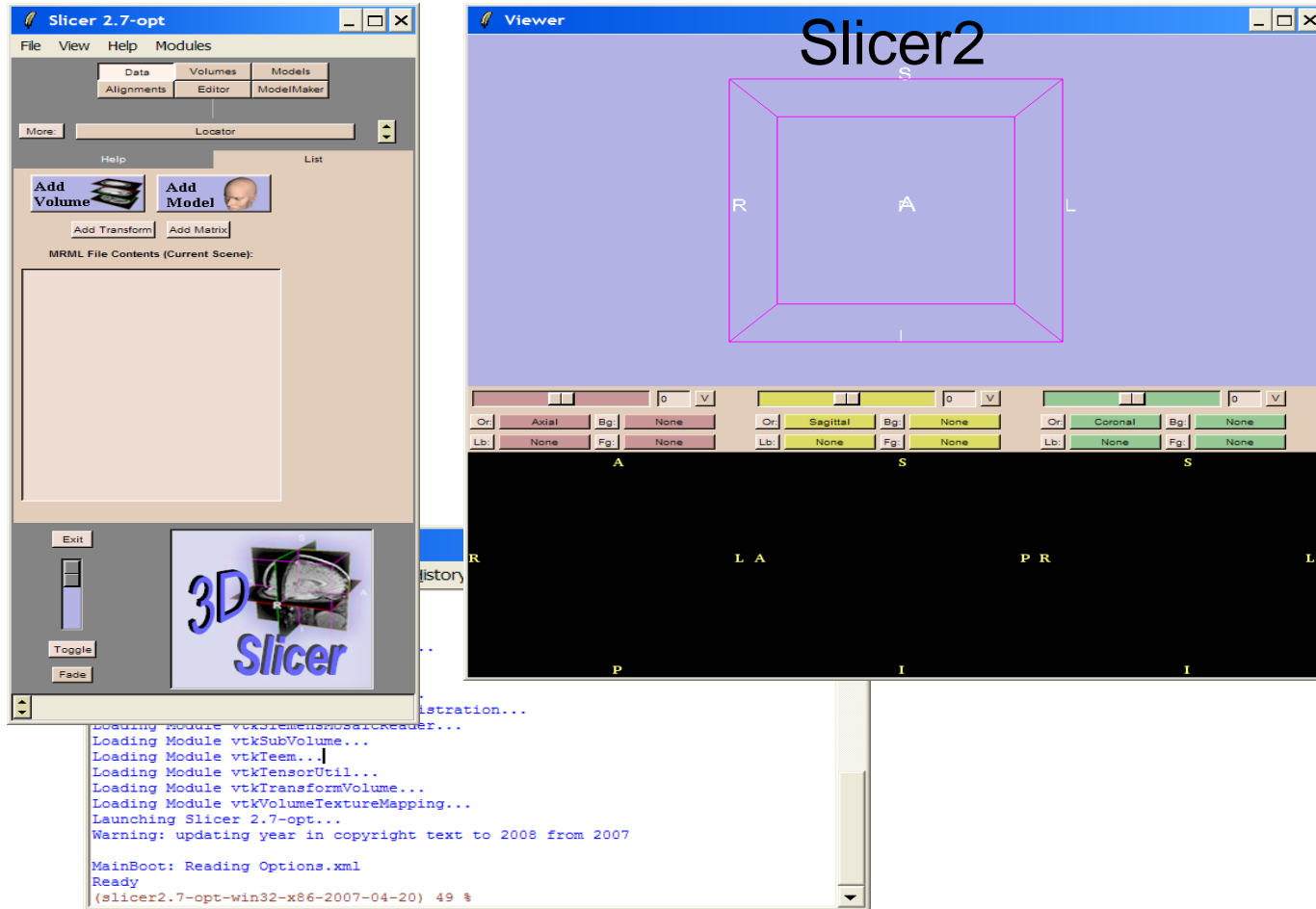
The NA-MIC Kit





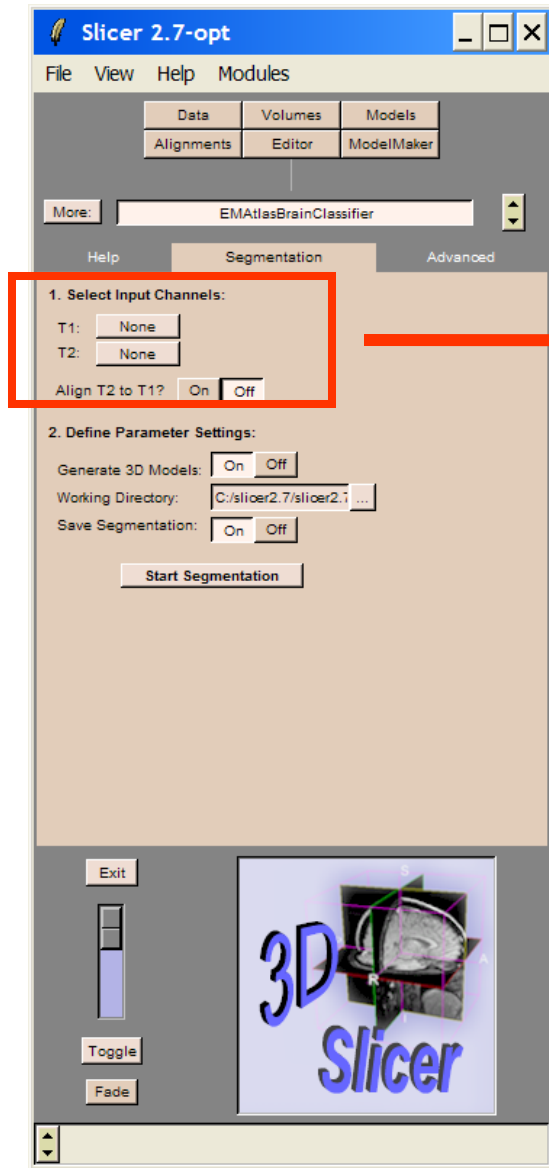
- An **end-user application** for image analysis
- An **open-source environment** for software development
- A software platform that is both **easy to use** for clinical researchers and **easy to extend** for programmers

Before Slicer3



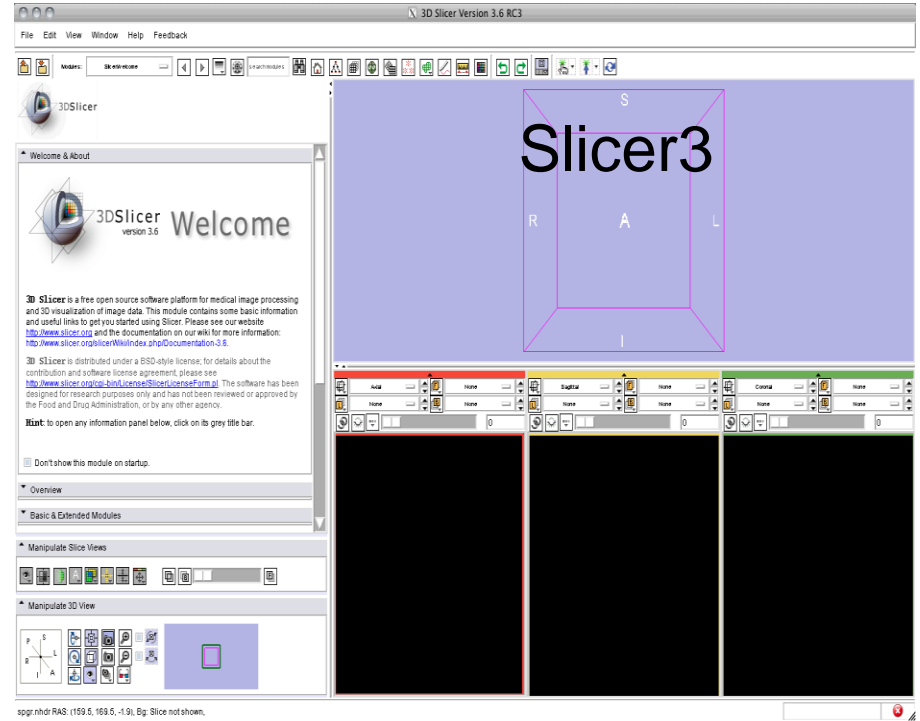
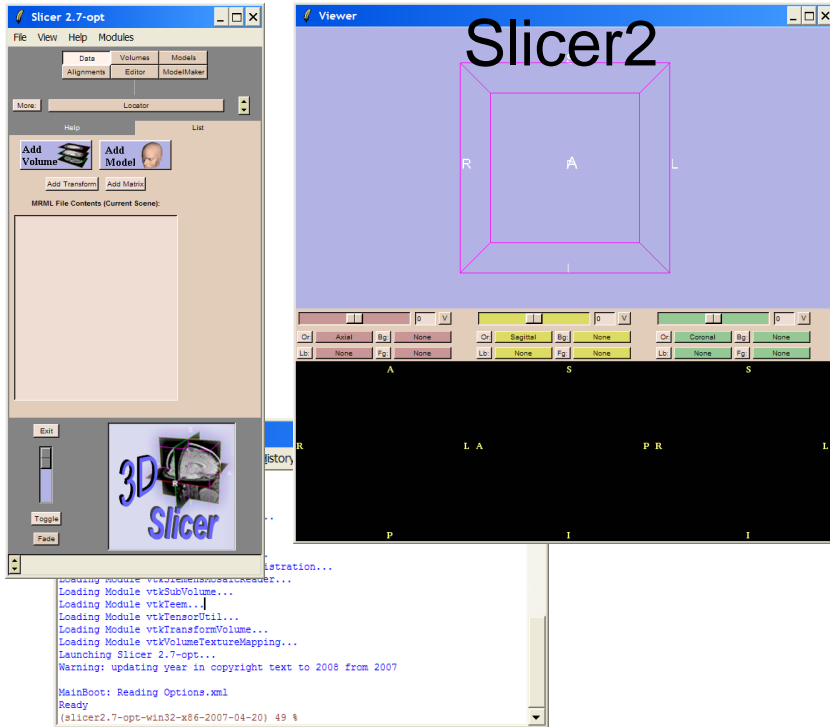


Programming into Slicer2



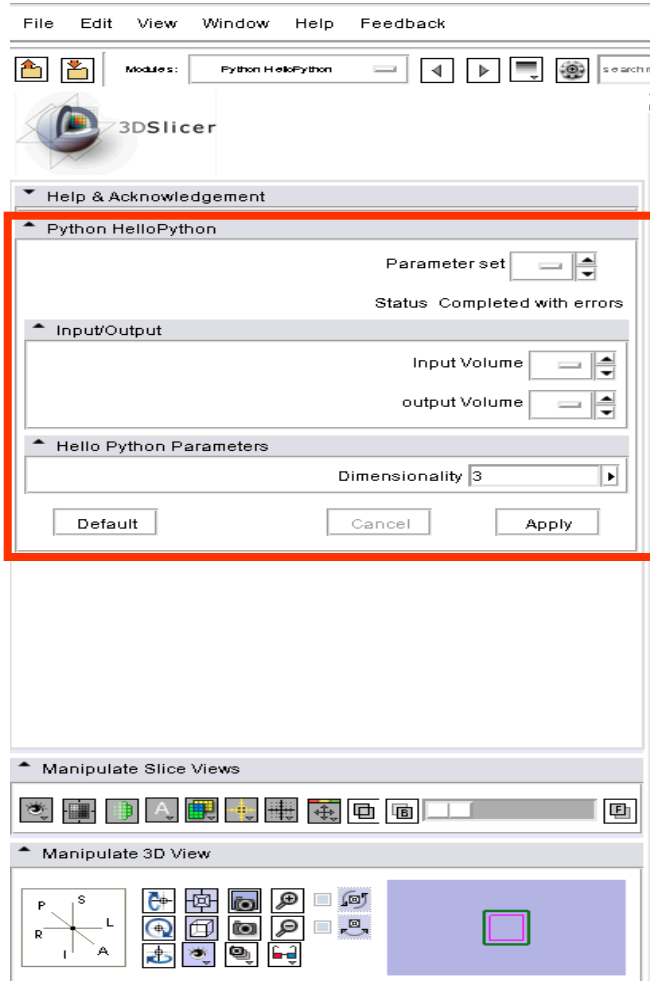
```
#-----  
# 1. Step  
#-----  
set f $fSeg.fStep1  
DevAddLabel $f.ITitle "1. Select Input Channels: " WTA  
pack $f.ITitle -side top -padx $Gui(pad) -pady 1 -anchor w  
frame $f.InInput -bg $Gui(activeWorkspace)  
pack $f.InInput -side top -padx 0 -pady 0 -anchor w  
foreach frame "Left Right" {  
    frame $f.InInput.f$frame -bg $Gui(activeWorkspace)  
    pack $f.InInput.f$frame -side left -padx 0 -pady $Gui(pad) }  
foreach LABEL "T1 T2" Input "SPGR T2W" {  
    DevAddLabel $f.InInput.fLeft.I$Input " ${LABEL}:"  
    pack $f.InInput.fLeft.I$Input -side top -padx $Gui(pad) -pady 1 -anchor w  
    set menubutton $f.InInput.fRight.m${Input}Select  
    set menu $f.InInput.fRight.m${Input}Select.m  
eval {menubutton $menubutton -text [Volume($EMAtlasBrainClassifier(Volume,${Input}),node) GetName] -  
relief raised -bd 2 -width 9 -menu $menu} $Gui(WMBA)  
    eval {menu $menu} $Gui(WMA)  
    TooltipAdd $menubutton "Select Volume defining ${Input}"  
    set EMAtlasBrainClassifier(mbSeg-${Input}Select) $menubutton  
    set EMAtlasBrainClassifier(mSeg-${Input}Select) $menu  
    # Have to update at UpdateMRML too  
    DevUpdateNodeSelectButton Volume EMAtlasBrainClassifier Seg-${Input}Select Volume,$Input  
    pack $menubutton -side top -padx $Gui(pad) -pady 1 -anchor w }  
frame $f.Align -bg $Gui(activeWorkspace)  
    TooltipAdd $f.Align "If the input T1 and T2 are not aligned with each other set flag here"  
    pack $f.Align -side top -padx 0 -pady 2 -padx $Gui(pad) -anchor w  
    DevAddLabel $f.Align.I$Align "Align T2 to T1? "  
    pack $f.Align.I$Align -side left -padx $Gui(pad) -pady 1 -anchor w  
    foreach value "1 0" text "On Off" width "4 4" {  
        eval {radiobutton $f.Align.r$value -width $width -indicatoron 0  
            -text "$text" -value "$value" -variable EMAtlasBrainClassifier(AlignInput) } $Gui(WCA)  
        pack $f.Align.r$value -side left -padx 0 -pady 0 }
```

From Slicer2 to Slicer3





The New Execution Model



```
<?xml version="1.0" encoding="utf-8"?>
<executable>
  <category> Demonstration </category>
  <title> Python HelloPython </title>
  <description> Slicer Developer Course </description>
  <version> 1.0 </version>
  <documentation-url> </documentation-url>
  <license></license>
  <contributor>
    Sonia Pujol, Ph.D., Surgical Planning Laboratory, Harvard Medical School
  </contributor>
  <acknowledgements> National Alliance for Medical Image Computing (NAMIC), Grant
  U54 EB005149. </acknowledgements>
  <parameters>
    <label>Input/Output</label>
    <description>Input/output parameters</description>
    <image>
      <name>helloPython</name>
      <label>Input Volume</label>
      <channel>input</channel>
      <index>0</index>
      <default>None</default>
      <description>Input volume</description>
    </image>
    <image>
      <name>helloPythonOutputVolume</name>
      <label>Output Volume</label>
      <channel>output</channel>
      <index>1</index>
      <default>None</default>
      <description>Output filtered</description>
    </image>
  </parameters>
</executable>
```



Slicer3 Execution Model

- This course is based on the [Execution Model](#) which provides a mechanism for incorporating command line programs as Slicer modules.
- Jim Miller, Dan Blezek, Bill Lorensen (GE)
- This course uses the Python interpreter that has been integrated to Slicer.





Material

This course requires the following material

- Slicer3-3.6.3-2011-06-07

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

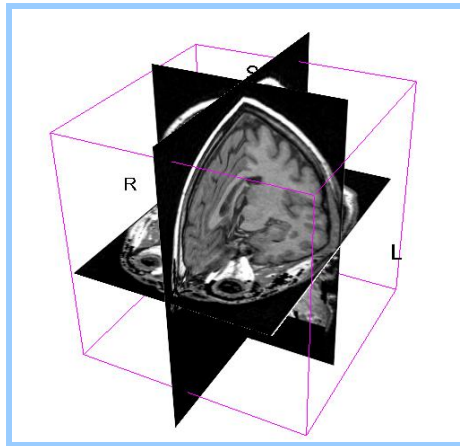
- HelloPython.zip

http://www.slicer.org/slicerWiki/index.php/Slicer3.6:TrainingSoftware_tutorials

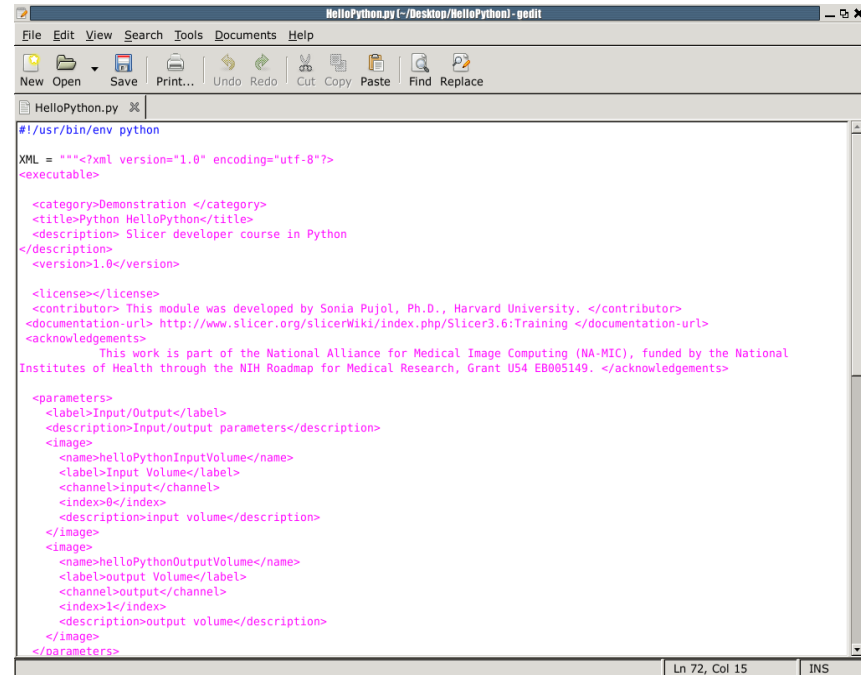
Disclaimer

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.

Unzip the HelloPython.zip archive



spgr.nhdr spgr.raw.gz
(124 SPGR images)



```
#!/usr/bin/env python

XML = """<?xml version="1.0" encoding="utf-8"?>
<executable>

<category>Demonstration </category>
<title>Python HelloPython</title>
<description> Slicer developer course in Python
</description>
<version>1.0</version>

<license></license>
<contributor> This module was developed by Sonia Pujol, Ph.D., Harvard University. </contributor>
<documentation-url> http://www.slicer.org/slicerWiki/index.php/Slicer3.6:Training </documentation-url>
<acknowledgements>
  This work is part of the National Alliance for Medical Image Computing (NA-MIC), funded by the National
  Institutes of Health through the NIH Roadmap for Medical Research, Grant U54 EB005149. </acknowledgements>

<parameters>
<label>Input/Output</label>
<description>Input/output parameters</description>
<image>
  <name>helloPythonInputVolume</name>
  <label>Input Volume</label>
  <channel>input</channel>
  <index>0</index>
  <description>input volume</description>
</image>
<image>
  <name>helloPythonOutputVolume</name>
  <label>output Volume</label>
  <channel>output</channel>
  <index>1</index>
  <description>output volume</description>
</image>
</parameters>

```

HelloPython.py

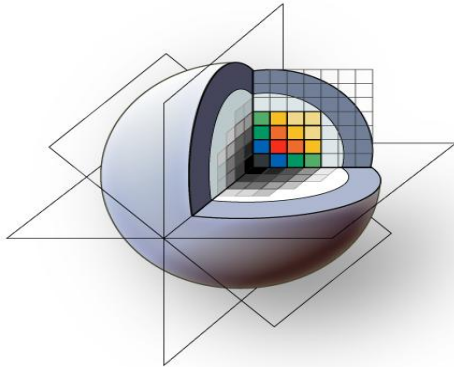


Overview

- Part A: Integration of the HelloPython.py program into Slicer3
- Part B: Implementation of the Laplace operator in the HelloPython module
- Part C: Image Sharpening using the Laplace operator



3DSlicer

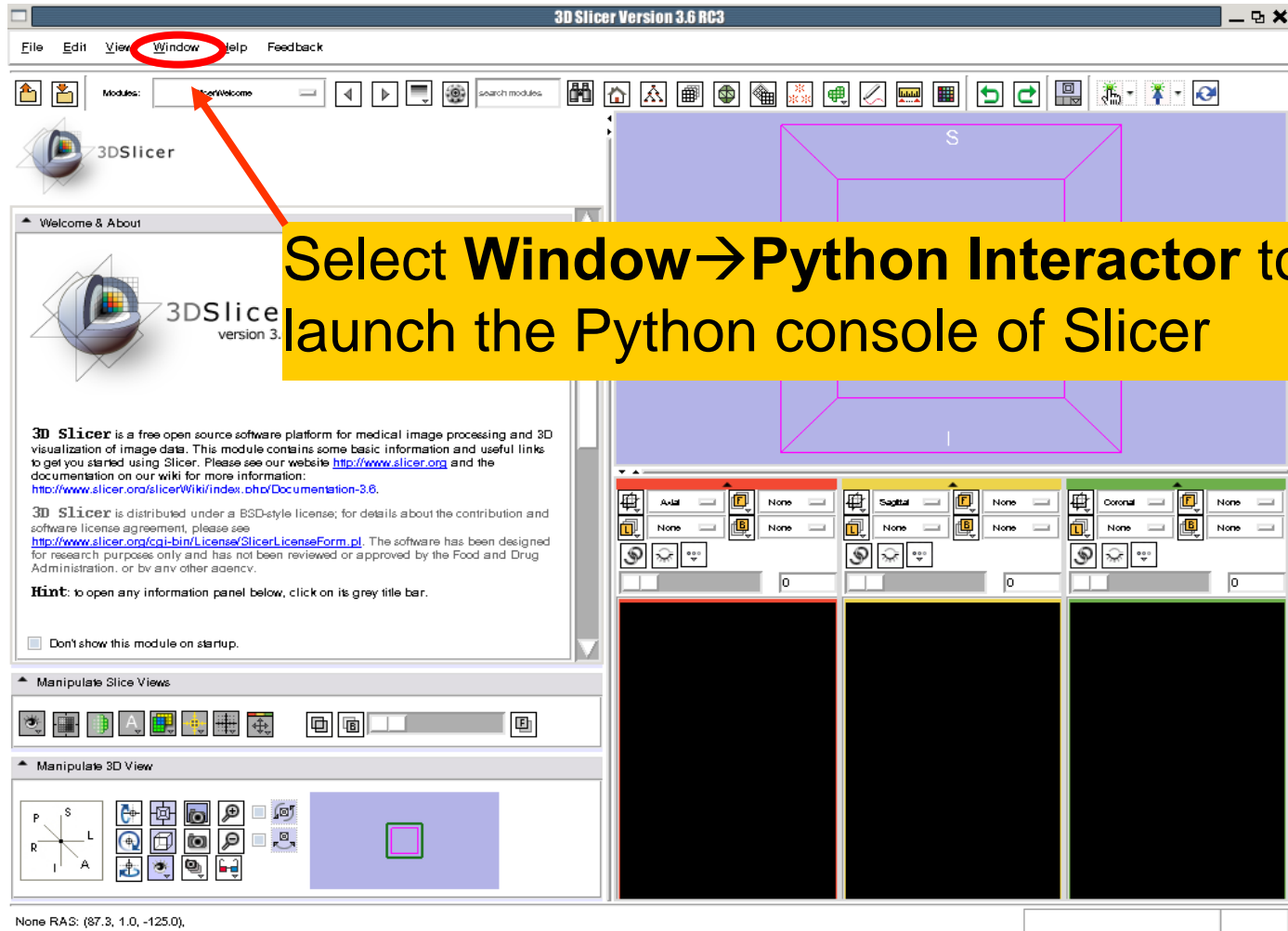


3DSlicer

Part A: Integrating HelloPython into Slicer3

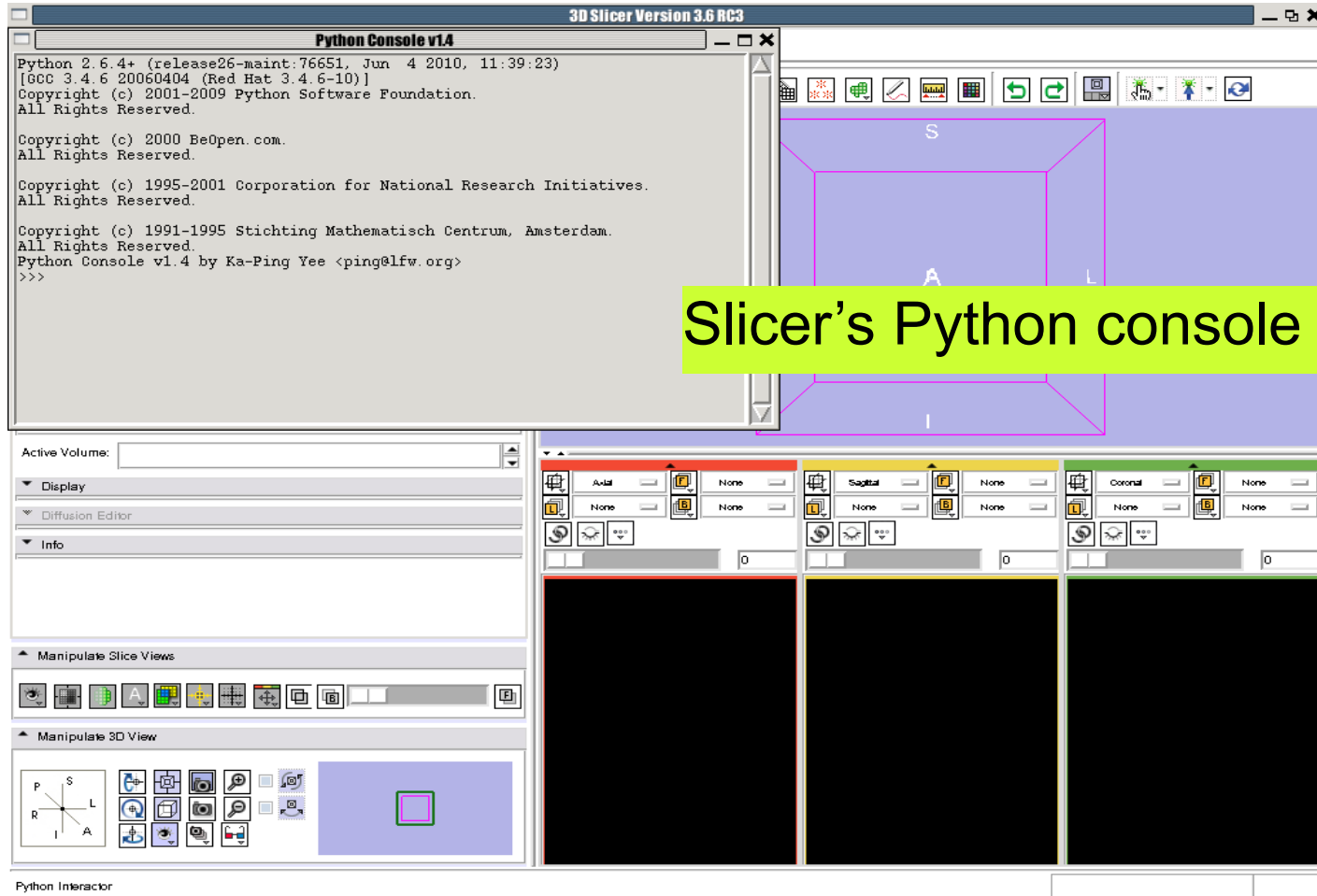
```
>HelloPython.py [C:/Desktop/HelloPython]-gedit
File Edit View Search Tools Documents Help
New Open Save Print... Undo Redo Cut Copy Paste Find Replace
HelloPython.py
#!/usr/bin/env python
XML - ***?xml version="1.0" encoding="utf-8"?
<executable>
<category>Demonstration </category>
<title>Python HelloPython</title>
<description> Slicer developer course in Python
</description>
<version>1.0</version>
<license></license>
<contributor> This module was developed by Sonia Pujol, Ph.D., Harvard University. </contributor>
<documentation-url> http://www.slicer.org/slicerWiki/index.php/Slicer3.6:Training </documentation-url>
<acknowledgements>
This work is part of the National Alliance for Medical Image Computing (NA-MIC), funded by the National
Institutes of Health through the NIH Roadmap for Medical Research, Grant US4 EB005149. </acknowledgements>
<parameters>
<label>Input/Output</label>
<description>Input/output parameters</description>
<image>
<name>helloPythonInputVolume</name>
<label>Input Volume</label>
<channel>input</channel>
<index>0</index>
<description>input volume</description>
</image>
<image>
<name>helloPythonOutputVolume</name>
<label>output Volume</label>
<channel>output</channel>
<index>1</index>
<description>output volume</description>
</image>
</parameters>
```

Python Console



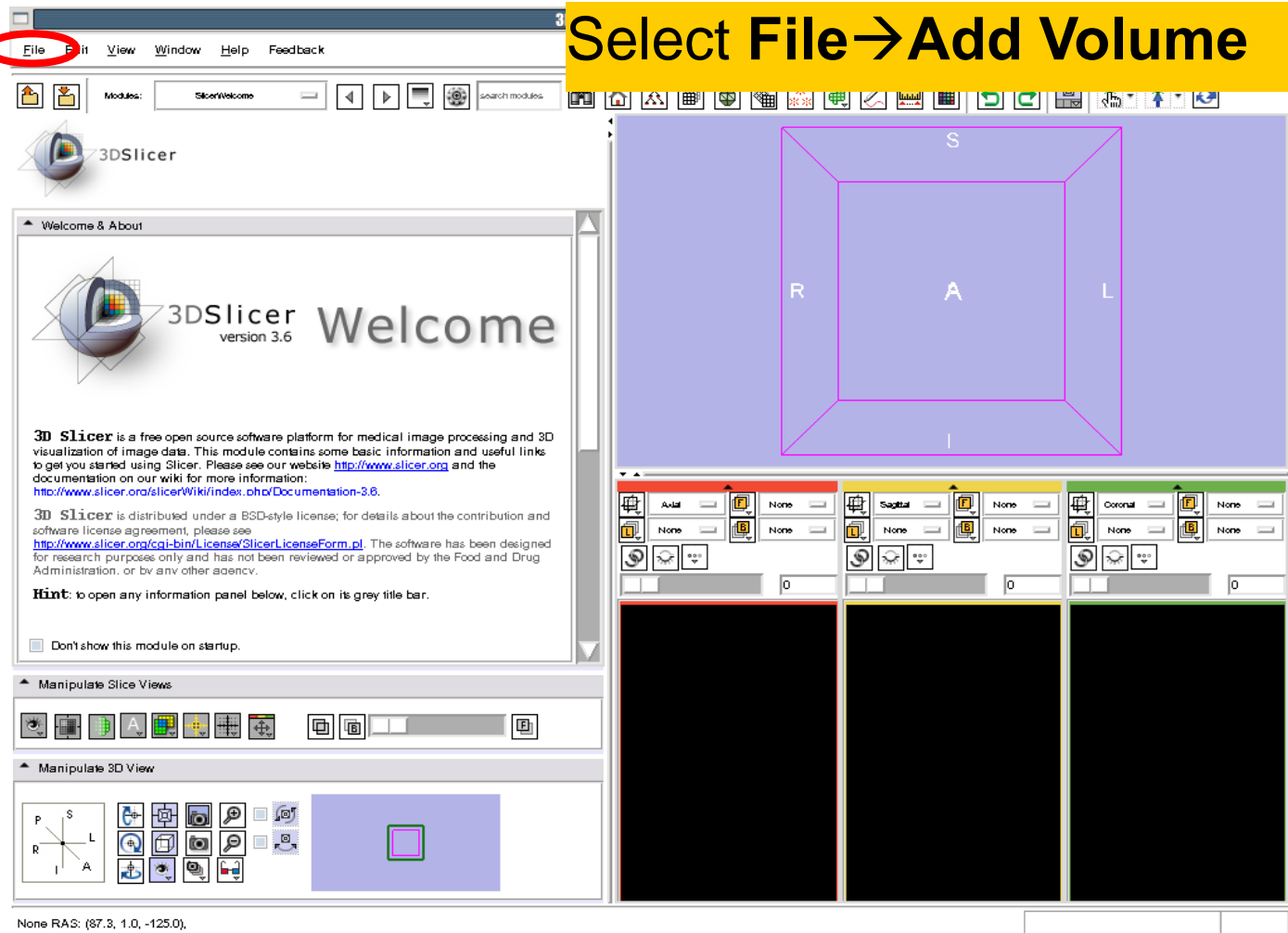
The screenshot shows the 3D Slicer Version 3.6 RC3 interface. The 'Window' menu is circled in red. A yellow callout box contains the text: **Select Window → Python Interactor to launch the Python console of Slicer**. The interface includes a menu bar (File, Edit, View, Window, Help, Feedback), a toolbar, and a main workspace with three viewports (Axial, Sagittal, Coronal) and a console area at the bottom.

Python Console



Python Console

Select File → Add Volume



File Edit View Window Help Feedback

Modules: SlicerWelcome

3DSlicer

Welcome & About

3DSlicer version 3.6 **Welcome**

3D Slicer is a free open source software platform for medical image processing and 3D visualization of image data. This module contains some basic information and useful links to get you started using Slicer. Please see our website <http://www.slicer.org> and the documentation on our wiki for more information: <http://www.slicer.org/SlicerWiki/index.php/Documentation-3.6>.

3D Slicer is distributed under a BSD-style license; for details about the contribution and software license agreement, please see <http://www.slicer.org/cgi-bin/License/SlicerLicenseForm.pl>. The software has been designed for research purposes only and has not been reviewed or approved by the Food and Drug Administration, or by any other agency.

Hint: to open any information panel below, click on its grey title bar.

Don't show this module on startup.

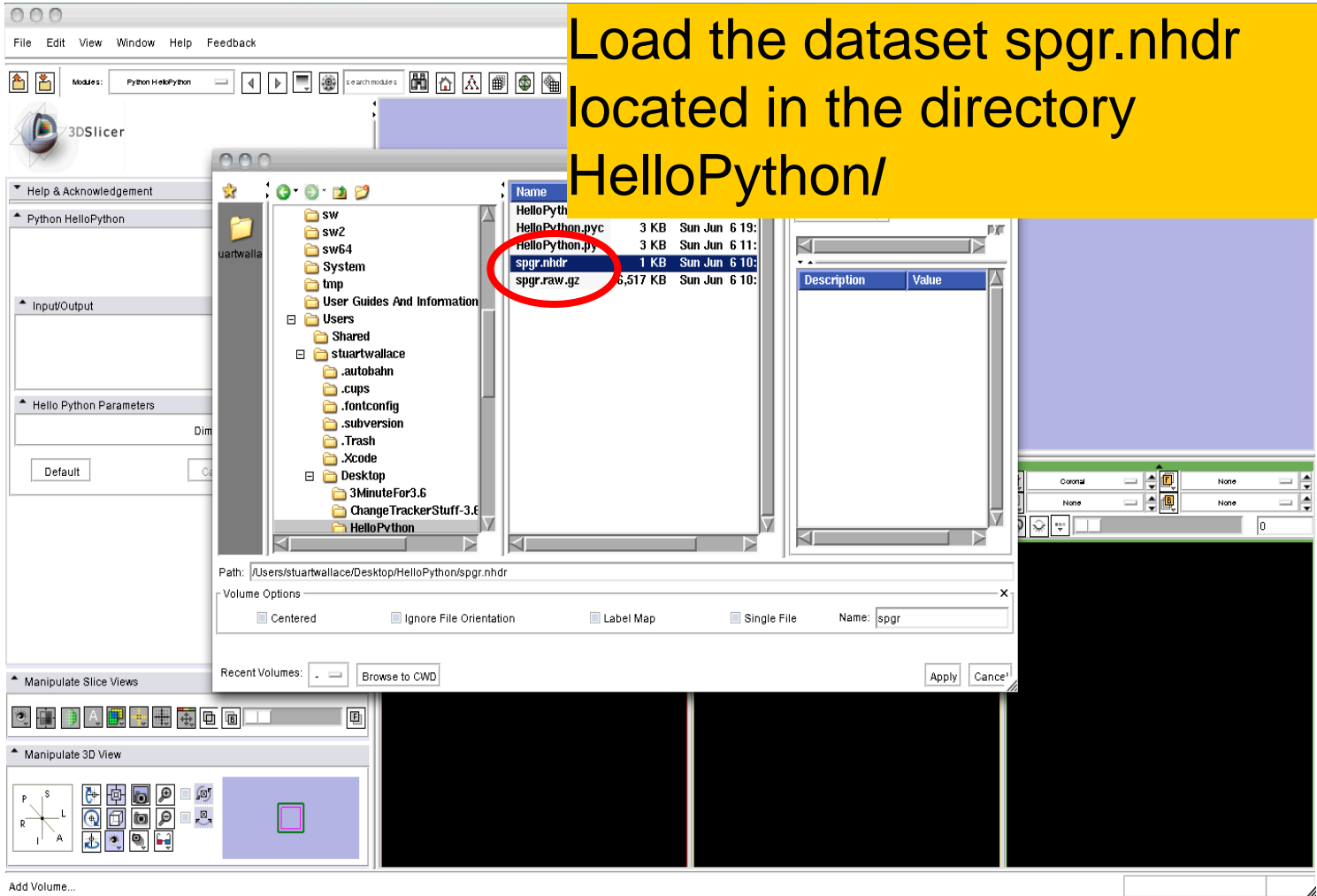
Manipulate Slice Views

Manipulate 3D View

None RAS: (87.3, 1.0, -125.0)

Python Console

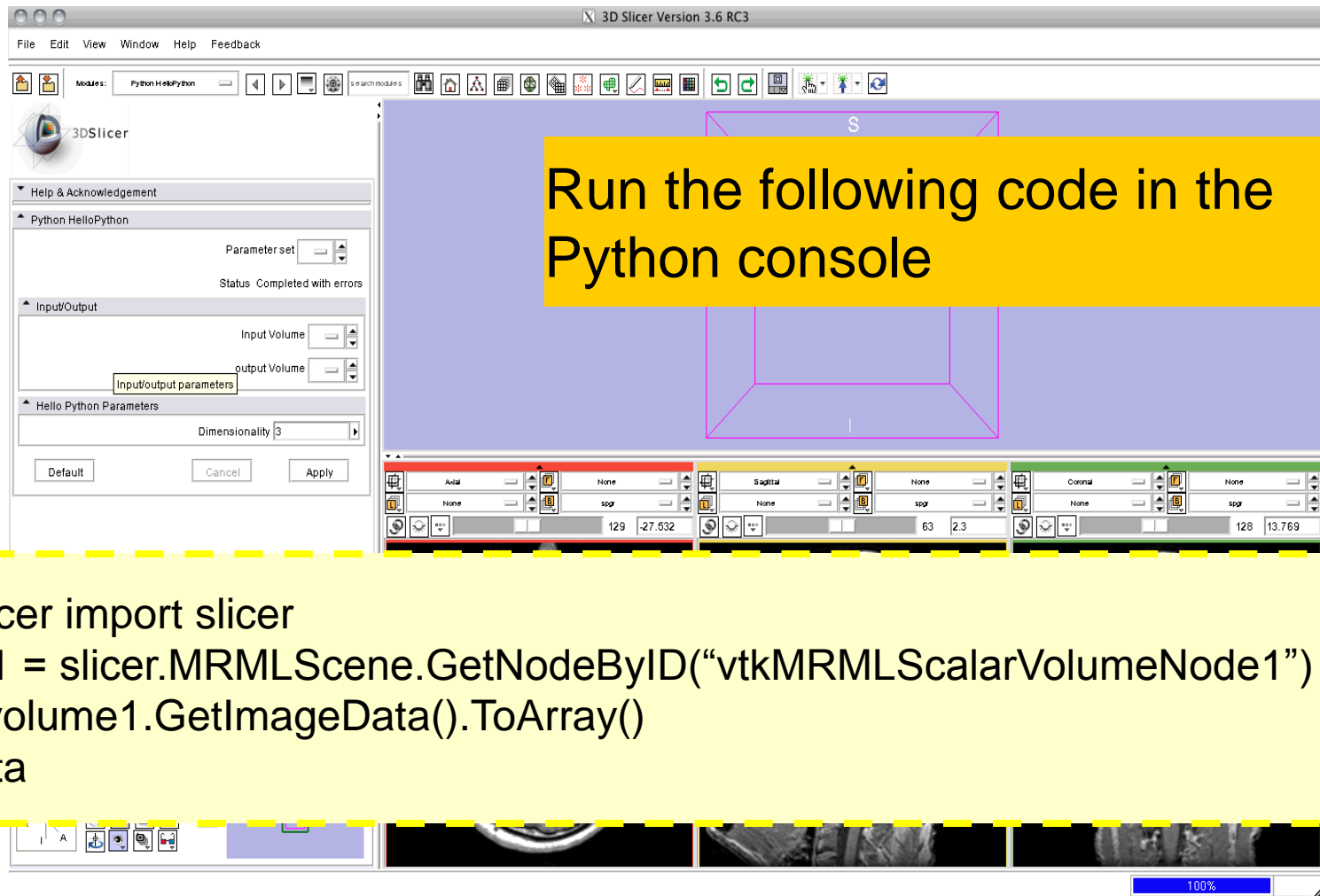
Load the dataset spgr.nhdr located in the directory HelloPython/



The screenshot shows the 3DSlicer interface with a file browser window open. The file browser displays a directory structure with the following files listed:

| Name | Size | Date | Time |
|-----------------|----------|---------------|------|
| HelloPyth | | | |
| HelloPython.pyc | 3 KB | Sun Jun 6 19: | |
| HelloPython.py | 3 KB | Sun Jun 6 11: | |
| spgr.nhdr | 1 KB | Sun Jun 6 10: | |
| spgr.raw.gz | 6,517 KB | Sun Jun 6 10: | |

The file 'spgr.nhdr' is circled in red. The file browser window also shows the path: /Users/stuartwallace/Desktop/HelloPython/spgr.nhdr and the file name 'spgr' in the 'Name' field.

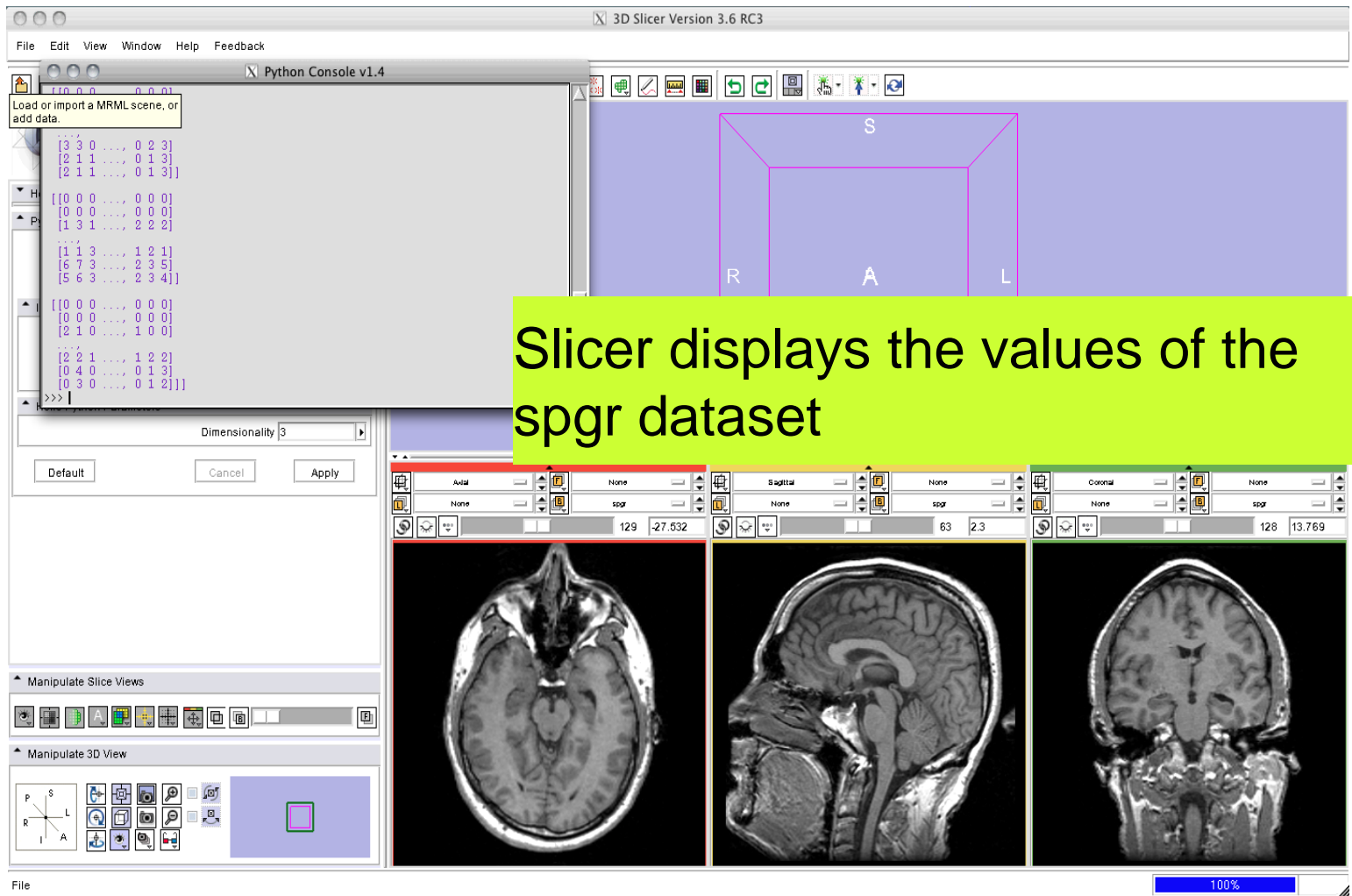


The screenshot shows the 3D Slicer 3.6 RC3 interface. A yellow text box is overlaid on the main 3D view, containing the instruction: "Run the following code in the Python console". Below this, a yellow dashed box contains the following Python code:

```
from Slicer import slicer
volume1 = slicer.MRMLScene.GetNodeByID("vtkMRMLScalarVolumeNode1")
data = volume1.GetImageData().ToArray()
print data
```


The interface also shows a parameter set dialog for "Python HelloPython" on the left, with fields for "Input Volume", "Output Volume", and "Dimensionality" (set to 3). The bottom of the interface shows the Slicer console and a 3D view of a volume with a purple bounding box.

Python Console



The screenshot displays the 3D Slicer software interface. At the top, the title bar reads "3D Slicer Version 3.6 RC3". Below it is a menu bar with "File", "Edit", "View", "Window", "Help", and "Feedback". A "Python Console v1.4" window is open, showing a list of matrices. A yellow text box is overlaid on the console area with the text "Slicer displays the values of the spgr dataset". The main 3D view area shows a purple wireframe box with axes labeled "S" (Superior), "R" (Right), "A" (Anterior), and "L" (Left). Below the 3D view are three slice view windows: Axial, Sagittal, and Coronal. The Axial view shows a brain slice with coordinates 129, -27.532. The Sagittal view shows a brain slice with coordinates 63, 2.3. The Coronal view shows a brain slice with coordinates 128, 13.769. At the bottom, there are panels for "Manipulate Slice Views" and "Manipulate 3D View". The status bar at the bottom right shows "100%" zoom.

HelloPython.py



Module Description

Module Parameters

Execute function

```
#!/usr/bin/env python
XML = """<?xml version="1.0" encoding="utf-8"?>
<executable>
  <category>Demonstration </category>
  <title>Python HelloPython</title>
  <description> Slicer developer course in Python
</description>
  <version>1.0</version>

  <license></license>
  <contributor> This module was developed by Sonia Pujol, Ph.D., Harvard University. </contributor>
  <documentation-url> http://www.slicer.org/slicerWiki/index.php/Slicer3.6:Training </documentation-url>
  <acknowledgements>
    This work is part of the National Alliance for Medical Image Computing (NA-MIC), funded by the National Institutes of
    Health through the NIH Roadmap for Medical Research, Grant U54 EB005149. </acknowledgements>

  <parameters>
    <label>Input/output</label>
    <description>Input/output parameters</description>
    <image>
      <name>helloPythonInputVolume</name>
      <label>Input Volume</label>
      <channel>input</channel>
      <index>0</index>
      <description>input volume</description>
    </image>
    <image>
      <name>helloPythonOutputVolume</name>
      <label>output Volume</label>
      <channel>output</channel>
      <index>1</index>
      <description>output volume</description>
    </image>
  </parameters>
  <parameters>
    <label>Hello Python Parameters</label>
    <description>Parameters of the Python Hello Python module </description>
    <integer>
      <name>dimensionality</name>
      <longflag>dimensionality</longflag>
      <description>Dimensionality of the Laplace operator</description>
      <label>Dimensionality</label>
      <default>3</default>
      <constraints>
        <minimum>2</minimum>
        <maximum>3</maximum>
      </constraints>
    </integer>
  </parameters>

</executable>
"""

def Execute ():
    slicer = __import__("Slicer")
    slicer = slicer.slicer
    scene = slicer.HFRMLScene

    return
```



Module Description

```
#!/usr/bin/env python
XML = """<?xml version="1.0" encoding="utf-8"?>
<executable>
<category>Demonstration </category>
<title>Python HelloPython</title>
<description> Slicer developer course in Python </description>
<version>1.0</version>
<license></license>
<contributor> This module was developed by Sonia Pujol, Ph.D., Harvard University. </contributor>
<documentation-url> http://www.slicer.org/slicerWiki/index.php/Slicer3.6:Training </documentation-url>
<acknowledgements>
    This work is part of the National Alliance for Medical Image Computing (NA-MIC),
    funded by the National Institutes of Health through the NIH Roadmap for Medical Research,
    Grant U54 EB005149.
</acknowledgements>
```

Module Parameters

```
<parameters>  
  <label>Input/Output</label>  
  <description>Input/output parameters</description>
```

Input
Volume

```
<image>  
  <name>HelloPythonInputVolume</name>  
  <label>Input Volume</label>  
  <channel>input</channel>  
  <index>0</index>  
  <description>input volume</description>  
</image>
```

A file that
specifies
the image

Output
Volume

```
<image>  
  <name>HelloPythonOutputVolume</name>  
  <label>Output Volume</label>  
  <channel>output</channel>  
  <index>1</index>  
  <description>output volume</description>  
</image>  
</parameters>
```



Execute Function

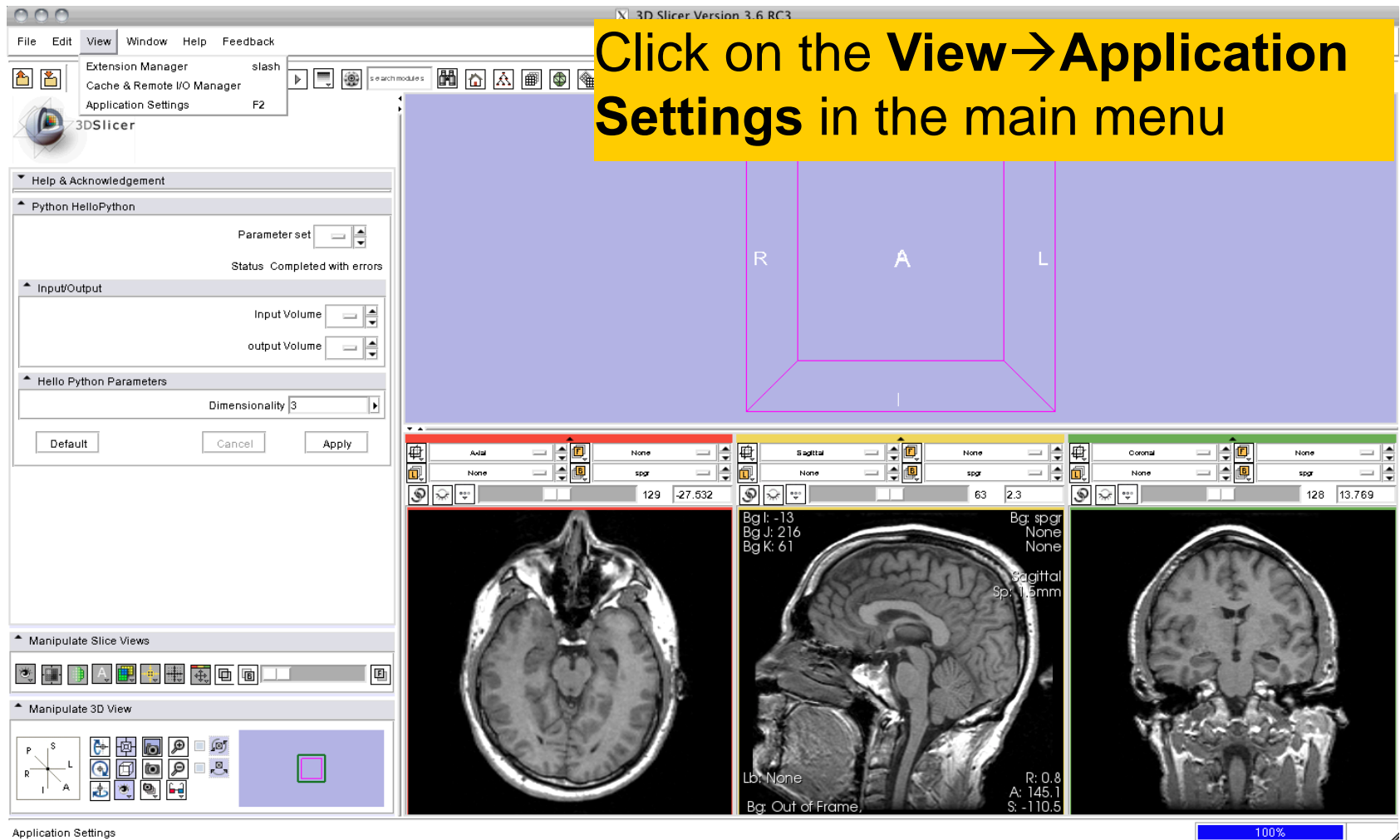
```
def Execute ():  
  
    Slicer = __import__("Slicer")  
    slicer = Slicer.slicer  
    scene = slicer.MRMLScene  
  
    return
```

The Slicer object will be the main interface to Slicer as a whole. s

Integrating HelloPython to Slicer3

3D Slicer Version 3.6 RC3

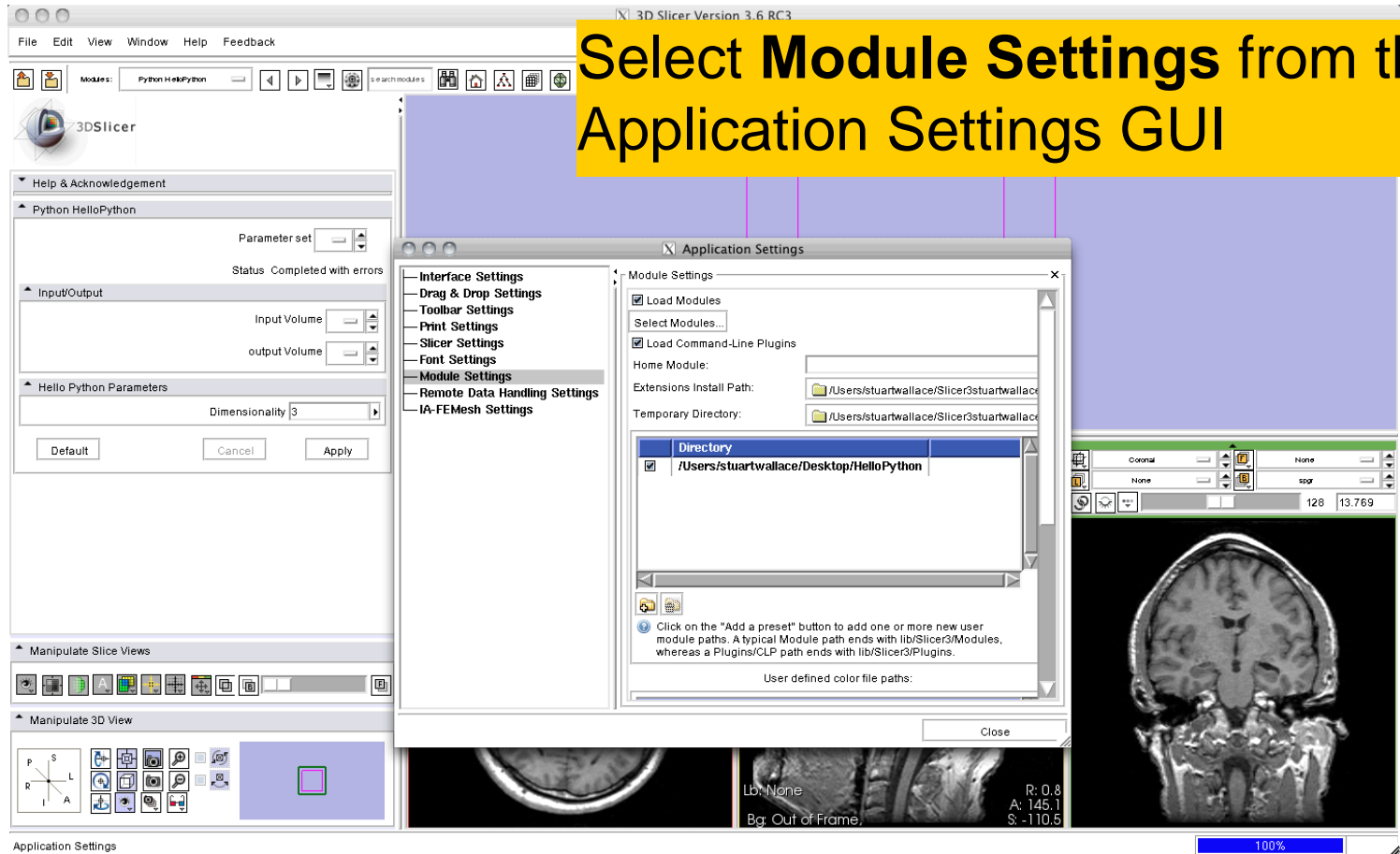
Click on the **View → **Application Settings** in the main menu**



Application Settings

100%

Integrating HelloPython to Slicer3



Select Module Settings from the Application Settings GUI

3D Slicer Version 3.6 RC3

Application Settings

Module Settings

- Load Modules
- Load Command-Line Plugins

Home Module:

Extensions Install Path: /Users/stuartwallace/Slicer3stuartwallace

Temporary Directory: /Users/stuartwallace/Slicer3stuartwallace

| Directory |
|--|
| <input checked="" type="checkbox"/> /Users/stuartwallace/Desktop/HelloPython |

Click on the "Add a preset" button to add one or more new user module paths. A typical Module path ends with lib/Slicer3/Modules, whereas a Plugins/CLP path ends with lib/Slicer3/Plugins.

User defined color file paths:

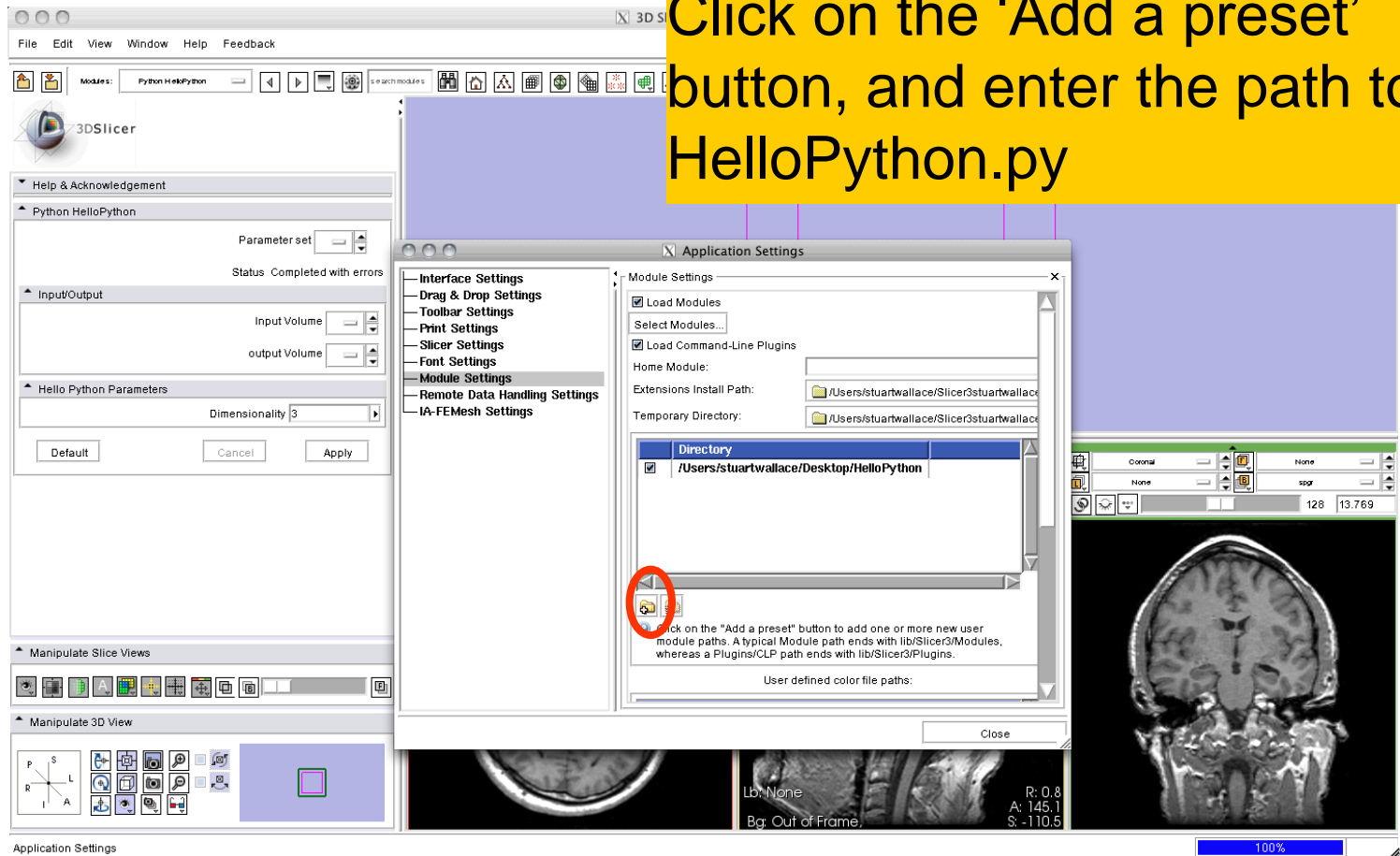
Close

Application Settings

100%

Integrating HelloPython to Slicer3

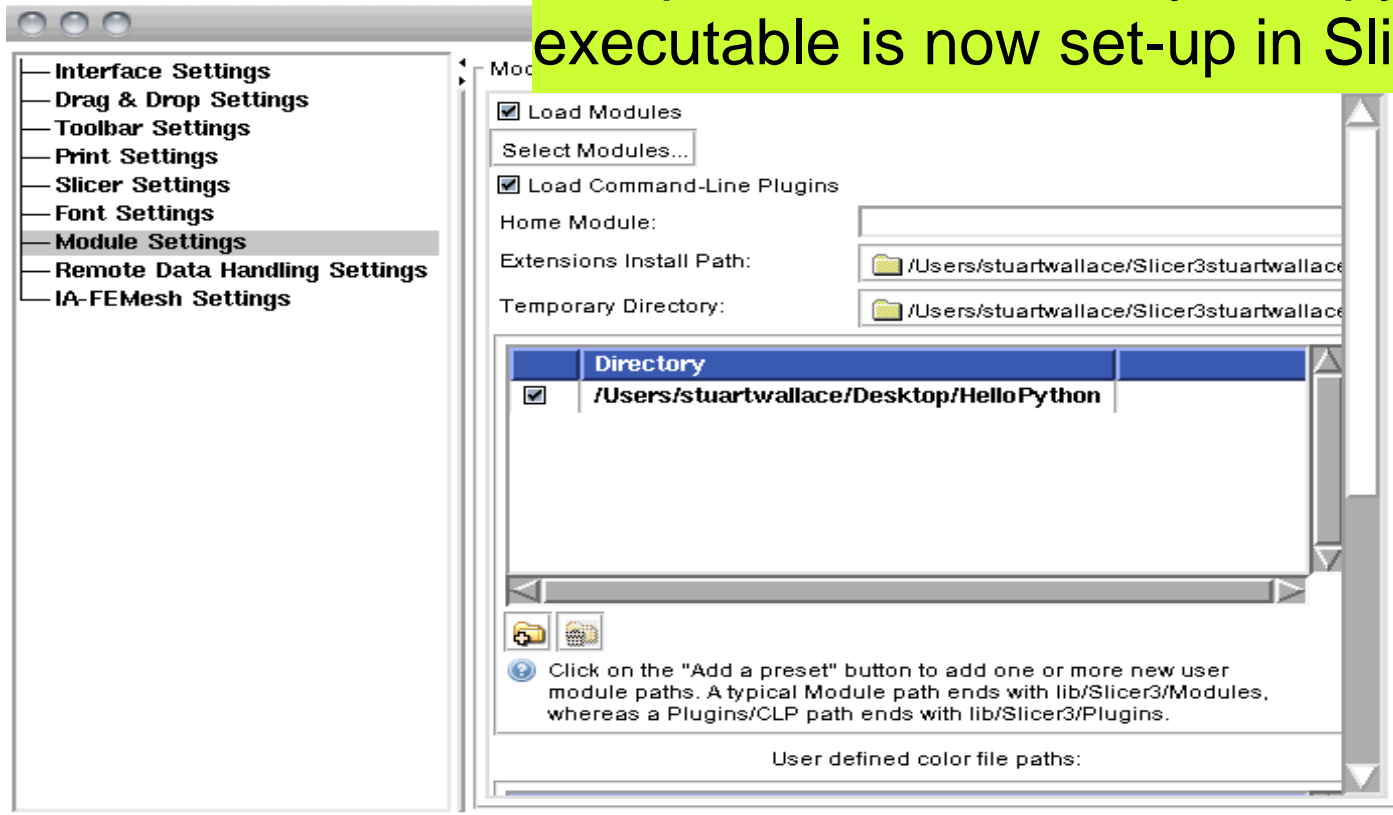
Click on the 'Add a preset' button, and enter the path to HelloPython.py



The screenshot shows the 3DSlicer application window with the 'Application Settings' dialog box open. The 'Module Settings' tab is selected, and the 'Add a preset' button is circled in red. The 'Directory' list shows the path `/Users/stuartwallace/Desktop/HelloPython`. The 'Add a preset' button is located at the bottom left of the dialog box. The main window shows the 'Python HelloPython' module settings and a 3D view of a brain MRI slice.

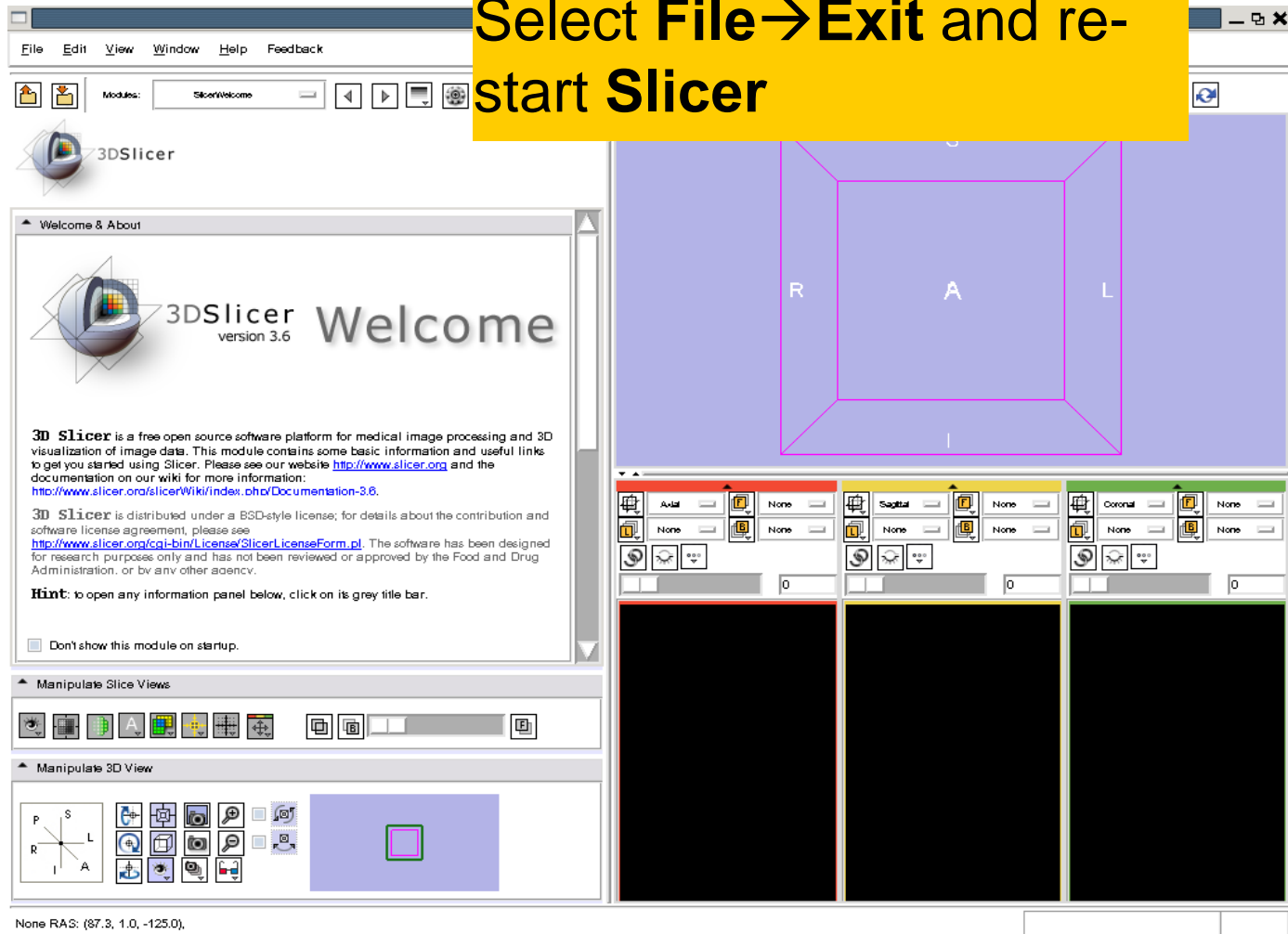
Integrating HelloPython to Slicer3

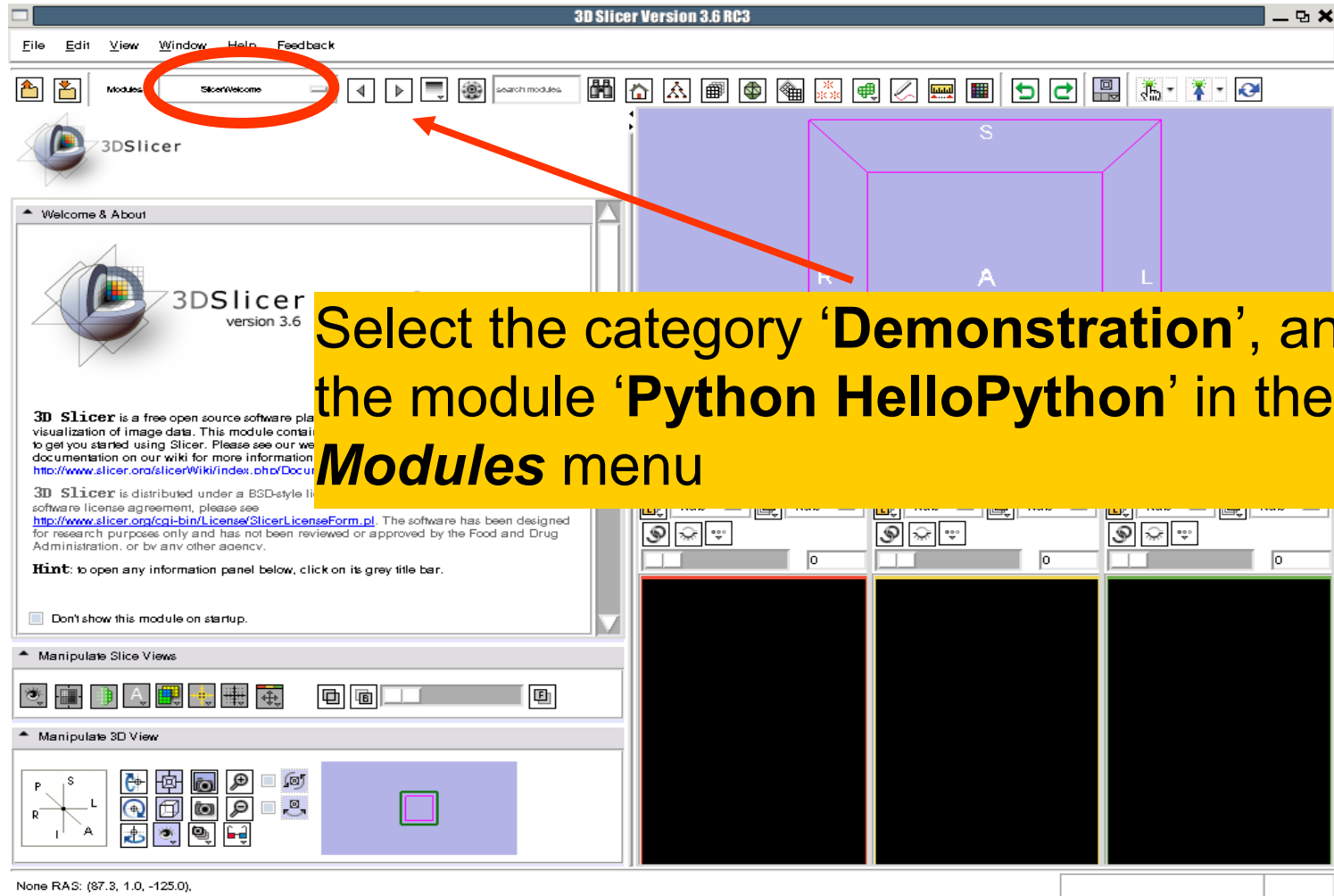
The path to the HelloPython.py executable is now set-up in Slicer3.



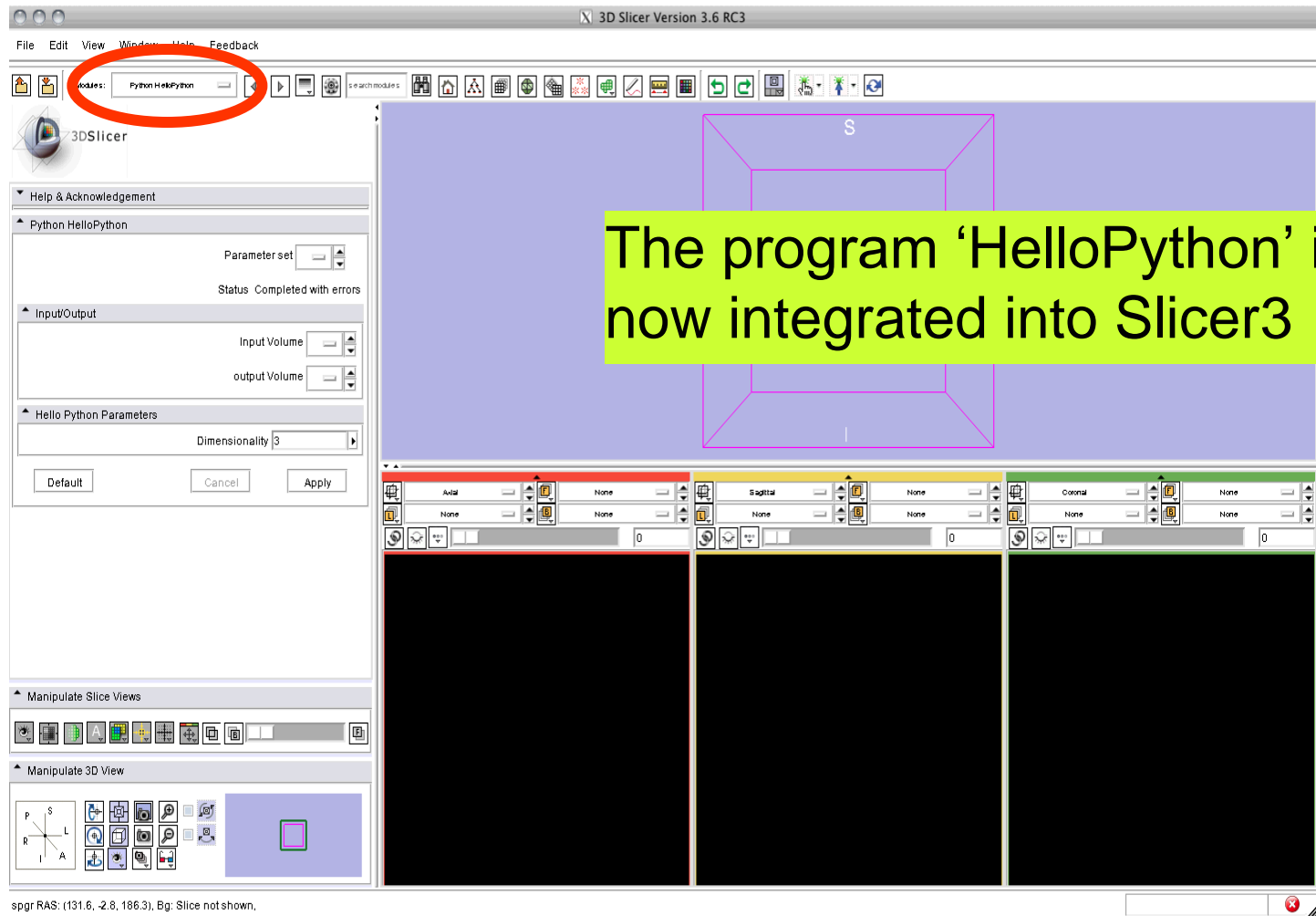
Click on the **Close** to exit the Application Settings window.

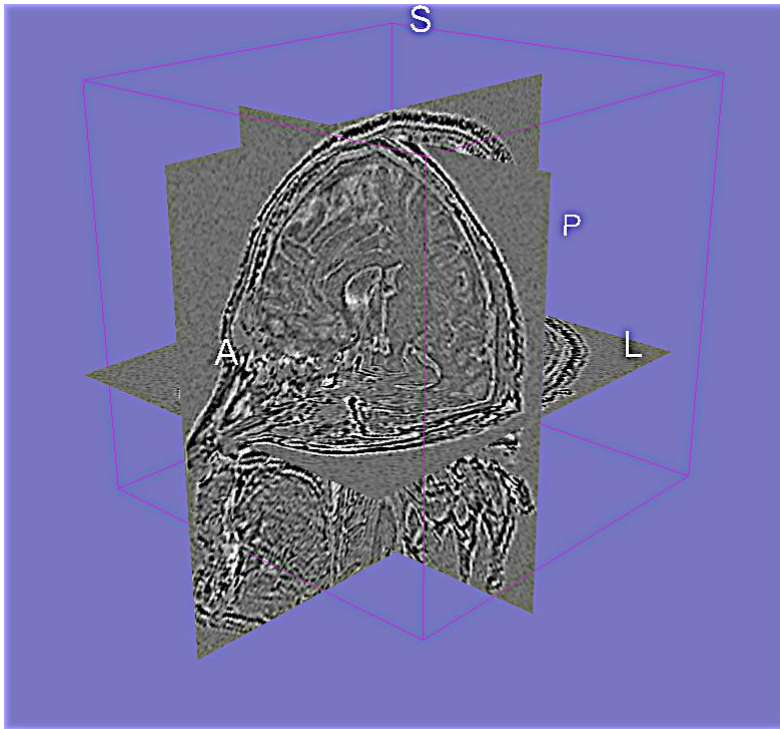
Integrating HelloPython to Slicer3





HelloPython Module





Part B: Implementing the Laplace* Operator

*named after Pierre-Simon, Marquis de Laplace (1749-1827)



Execute Function

```
def Execute (HelloPythonInputVolume, HelloPythonOutputVolume):
```

Add the I/O code

```
Slicer = __import__("Slicer")
```

```
slicer = Slicer.slicer
```

```
scene = slicer.MRMLScene
```

```
inputVolume = scene.GetNodeByID(HelloPythonInputVolume)
```

```
outputVolume = scene.GetNodeByID(HelloPythonOutputVolume)
```

```
return
```




Laplace Operator

```
def Execute (HelloPythonInputVolume, HelloPythonOutputVolume):
```

```
    Slicer = __import__("Slicer")
```

```
    slicer = Slicer.slicer
```

```
    scene = slicer.MRMLScene
```

```
    inputVolume = scene.GetNodeByID(HelloPythonInputVolume)
```

```
    outputVolume = scene.GetNodeByID(HelloPythonOutputVolume)
```

```
    laplacian = slicer.vtkImageLaplacian()
```

```
    laplacian.SetInput(inputVolume.GetImageData())
```

```
    return
```

Add the Laplace operator

Laplace Operator

```
<parameters>
  <label>Input/Output</label>
  <description>Input/output parameters</description>
  <image>
    <name>HelloPythonInputVolume</name>
    <label>Input Volume</label>
    <channel>input</channel>
    <index>0</index>
    <description>input volume</description>
  </image>
  <image>
    <name>HelloPythonOutputVolume</name>
    <label>Output Volume</label>
    <channel>output</channel>
    <index>1</index>
    <description>output volume</description>
  </image>
</parameters>
```

```
<parameters>
  <label>Hello Python Parameters</label>
  <description> Parameters of the Python Hello Python module </description>
</parameters>
```

Add a new parameter group for the Laplace operator



Laplace Operator

```
<parameters>
  <label>Hello Python Parameters</label>
  <description>Parameters of the Python Hello Python module </description>
  <integer>
    <name>dimensionality</name>
    <longflag>dimensionality</longflag>
    <description>Dimensionality of the Laplace operator</description>
    <label>Dimensionality</label>
    <default>3</default>
    <constraints>
      <minimum>2</minimum>
      <maximum>3</maximum>
    </constraints>
  </integer>
</parameters>
```

Add the Laplace
operator's
dimensionality

Laplace Operator

```
def Execute (HelloPythonInputVolume, HelloPythonOutputVolume,
```

```
dimensionality=3):
```

```
Slicer = __import__("Slicer")
```

```
slicer = Slicer.slicer
```

```
scene = slicer.MRMLScene
```

```
inputVolume = scene.GetNodeByID(HelloPythonInputVolume)
```

```
outputVolume = scene.GetNodeByID(HelloPythonOutputVolume)
```

```
laplacian = slicer.vtkImageLaplacian()
```

```
laplacian.SetInput(inputVolume.GetImageData())
```

```
laplacian.SetDimensionality(dimensionality)
```

```
return
```

Set-up the corresponding dimensionality parameter in the Python code



Laplace Operator

```
def Execute (HelloPythonInputVolume, HelloPythonOutputVolume,  
dimensionality=3):
```

```
Slicer = __import__("Slicer")
```

```
slicer = Slicer.slicer
```

```
scene = slicer.MRMLScene
```

```
inputVolume = scene.GetNodeByID(HelloPythonInputVolume)
```

```
outputVolume = scene.GetNodeByID(HelloPythonOutputVolume)
```

```
laplacian = slicer.vtkImageLaplacian()
```

```
laplacian.SetInput(inputVolume.GetImageData())
```

```
laplacian.SetDimensionality(dimensionality)
```

```
laplacian.Update()
```

```
outputVolume.SetAndObserveImageData(laplacian.GetOutput())
```

```
return
```

Add code to get the output
of the Laplace operator



Laplace Operator

```
def Execute (HelloPythonInputVolume, HelloPythonOutputVolume,  
dimensionality=3):
```

```
Slicer = __import__("Slicer")
```

```
slicer = Slicer.slicer
```

```
scene = slicer.MRMLScene
```

```
inputVolume = scene.GetNodeByID(HelloPythonInputVolume)
```

```
outputVolume = scene.GetNodeByID(HelloPythonOutputVolume)
```

```
laplacian = slicer.vtkImageLaplacian()
```

```
laplacian.SetInput(inputVolume.GetImageData())
```

```
laplacian.SetDimensionality(dimensionality)
```

```
laplacian.Update()
```

```
outputVolume.SetAndObserveImageData(laplacian.GetOutput())
```

```
matrix = slicer.vtkMatrix4x4()
```

```
inputVolume.GetIJKToRASMatrix(matrix)
```

```
outputVolume.SetIJKToRASMatrix(matrix)
```

```
return
```

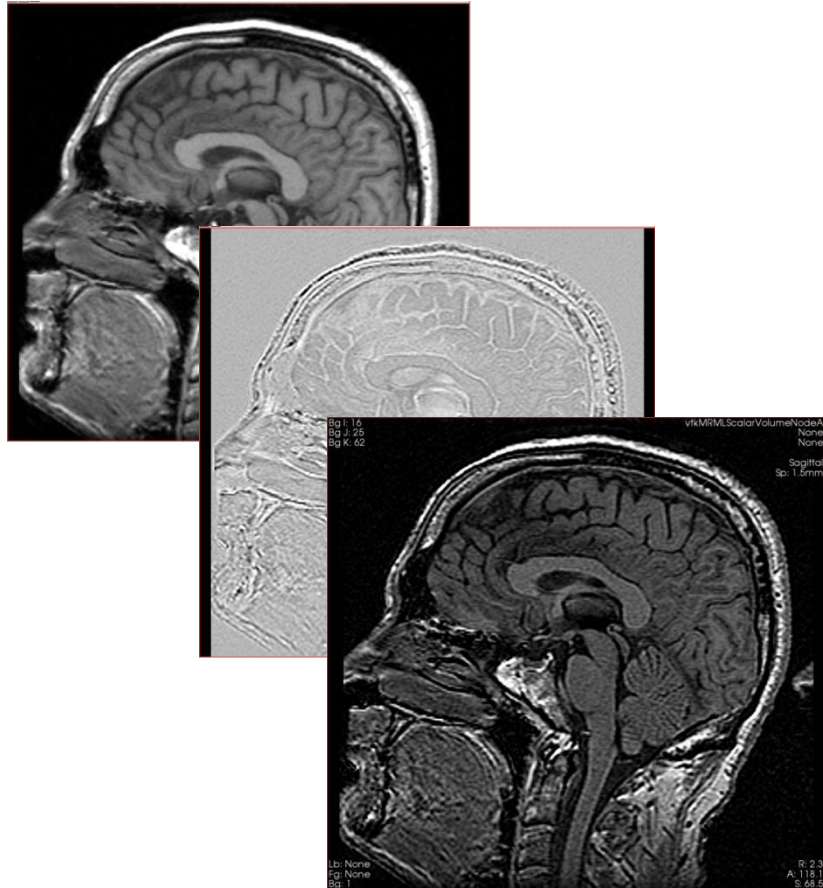
Place back the Laplacian of the image in the RAS reference system.



Integrating HelloPython to Slicer3

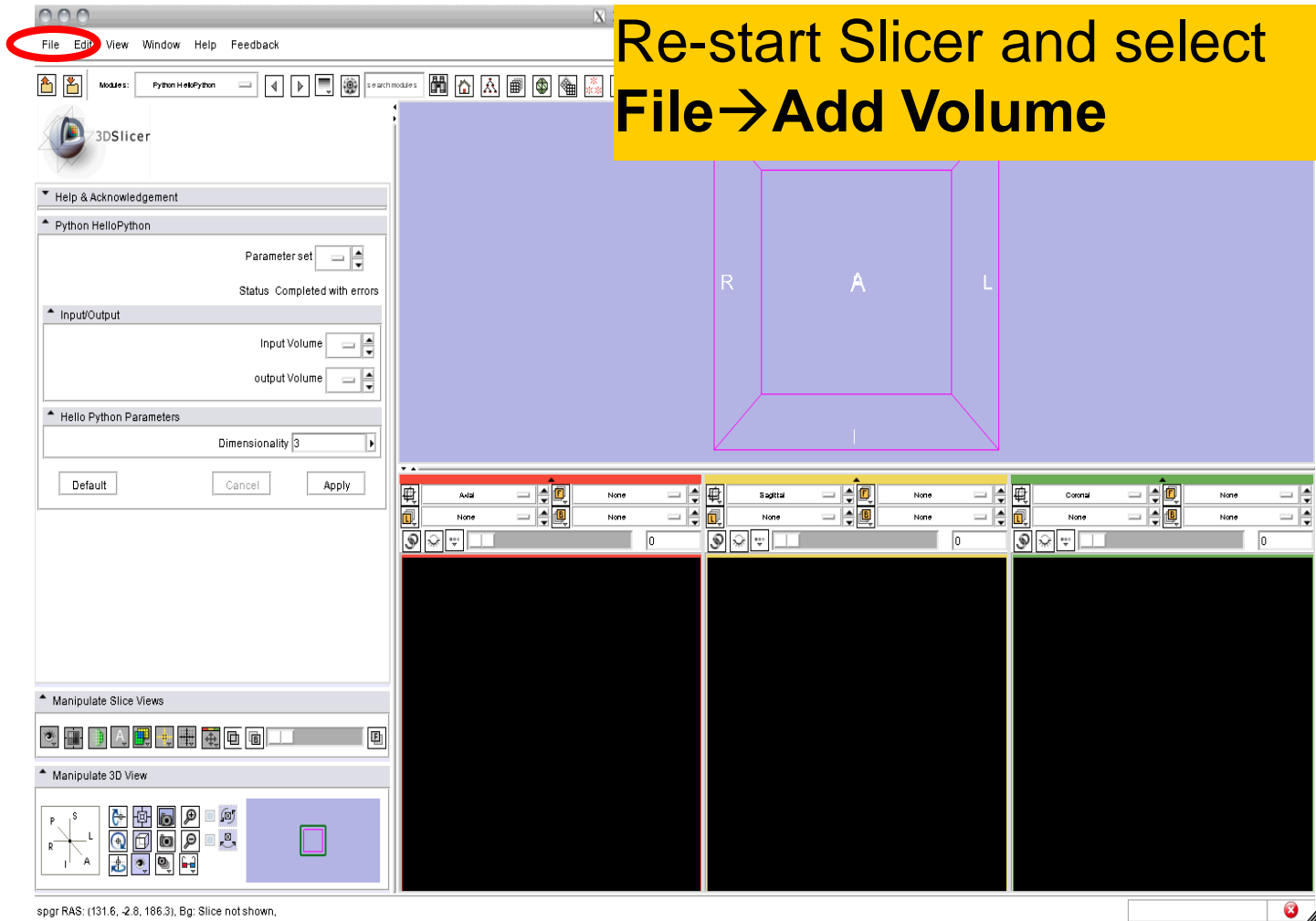
```
def Execute (HelloPythonInputVolume, HelloPythonOutputVolume,
dimensionality=3):
    Slicer = __import__("Slicer")
    slicer = Slicer.slicer
    scene = slicer.MRMLScene
    inputVolume = scene.GetNodeByID(HelloPythonInputVolume)
    outputVolume = scene.GetNodeByID(HelloPythonOutputVolume)
    laplacian = slicer.vtkImageLaplacian()
    laplacian.SetInput(inputVolume.GetImageData())
    laplacian.SetDimensionality(dimensionality)
    laplacian.Update()
    outputVolume.SetAndObserveImageData(laplacian.GetOutput())
    matrix = slicer.vtkMatrix4x4()
    inputVolume.GetIJKToRASMatrix(matrix)
    outputVolume.SetIJKToRASMatrix(matrix)
    return
```

Save the HelloPython.py file and exit Slicer.



Part C: Image Sharpening with the Laplace Operator

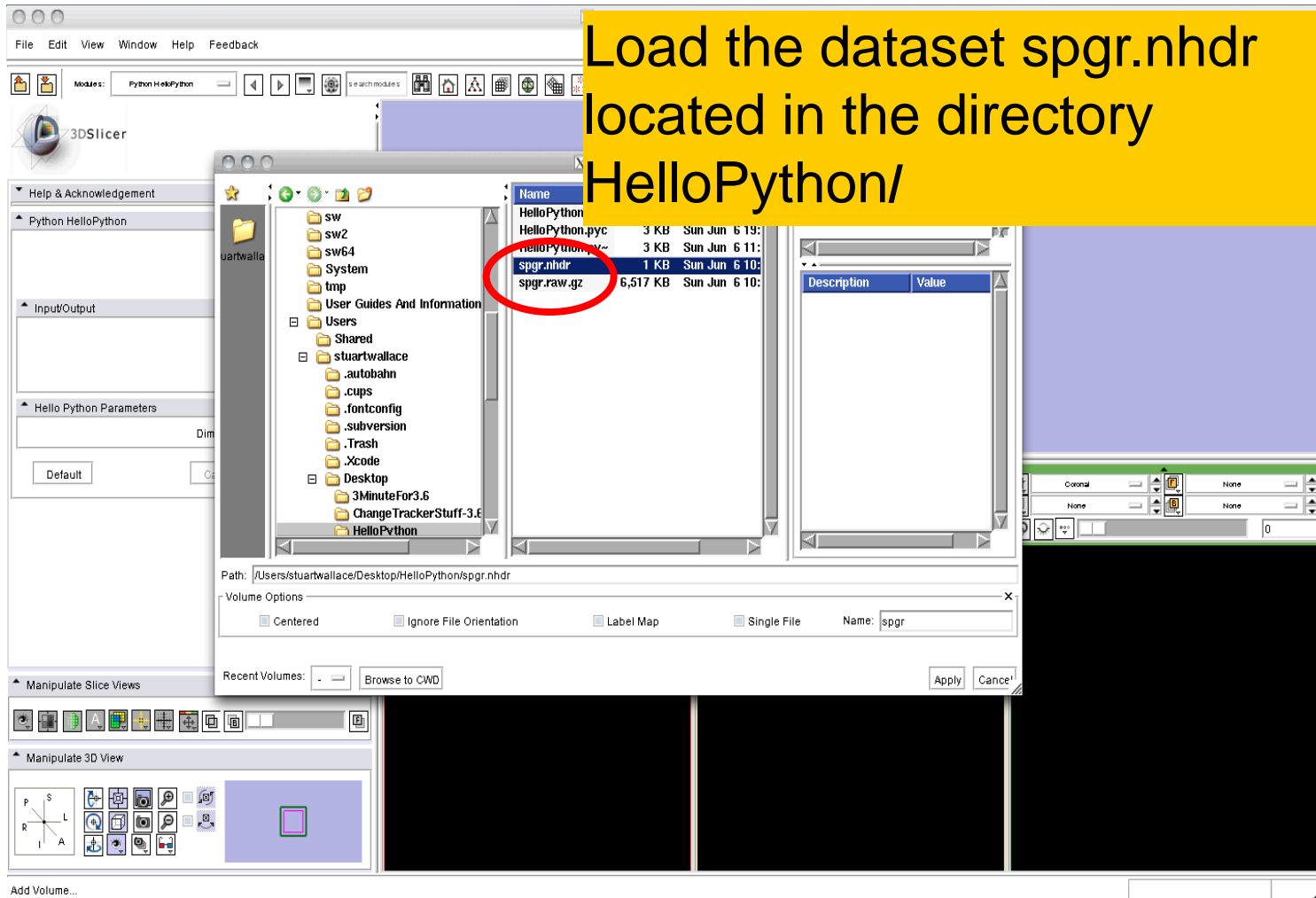
Running the Laplace Operator



The screenshot shows the 3DSlicer application window. The menu bar at the top includes 'File', 'Edit', 'View', 'Window', 'Help', and 'Feedback'. The 'File' menu is circled in red. A yellow callout box with black text reads: 'Re-start Slicer and select File → Add Volume'. The main 3D view area is a light purple color with a white wireframe box in the center, labeled with 'R', 'A', 'L', and 'I' for Right, Anterior, Left, and Inferior. Below the 3D view are three slice view panels: Axial (red header), Sagittal (yellow header), and Coronal (green header). Each panel has a 'None' selection and a '0' value. The left sidebar contains several panels: 'Help & Acknowledgement', 'Python HelloPython' (with 'Parameter set' and 'Status Completed with errors'), 'Input/Output' (with 'Input Volume' and 'output Volume'), 'Hello Python Parameters' (with 'Dimensionality 3'), 'Manipulate Slice Views', and 'Manipulate 3D View' (with a 3D orientation diagram and a small 3D view). At the bottom left, the status bar reads 'spgr RAS: (131.6, -2.8, 186.3), Bg: Slice not shown.'

Running the Laplace Operator

Load the dataset spgr.nhdr located in the directory HelloPython/



| Name | Size | Modified |
|-----------------|----------|---------------|
| HelloPython | | |
| HelloPython.pyc | 3 KB | Sun Jun 6 19: |
| HelloPython.pyw | 3 KB | Sun Jun 6 11: |
| spgr.nhdr | 1 KB | Sun Jun 6 10: |
| spgr.raw.gz | 6,517 KB | Sun Jun 6 10: |

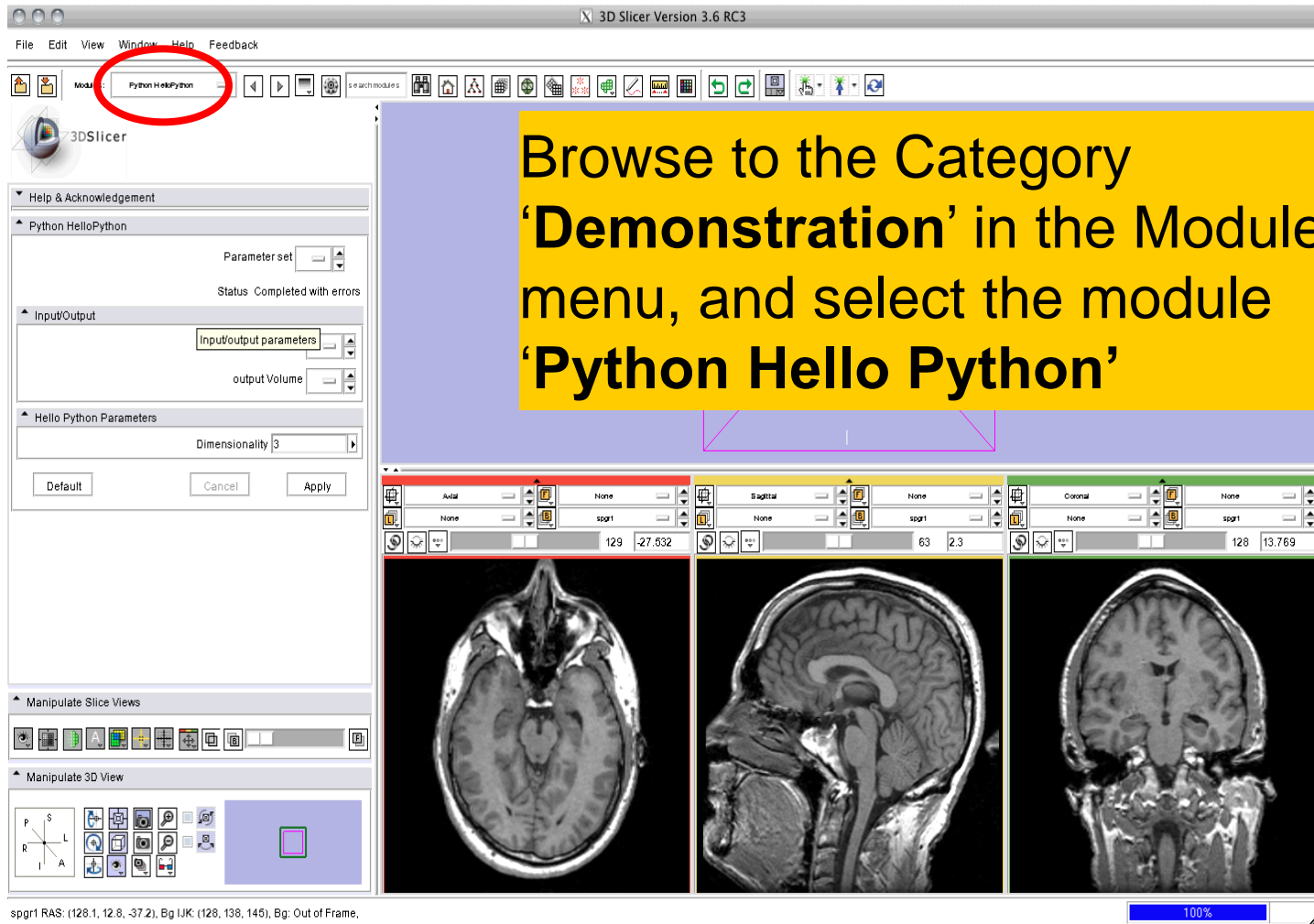
Path: /Users/stuartwallace/Desktop/HelloPython/spgr.nhdr

Volume Options

Centered Ignore File Orientation Label Map Single File Name: spgr

Recent Volumes:

Running the Laplace Operator

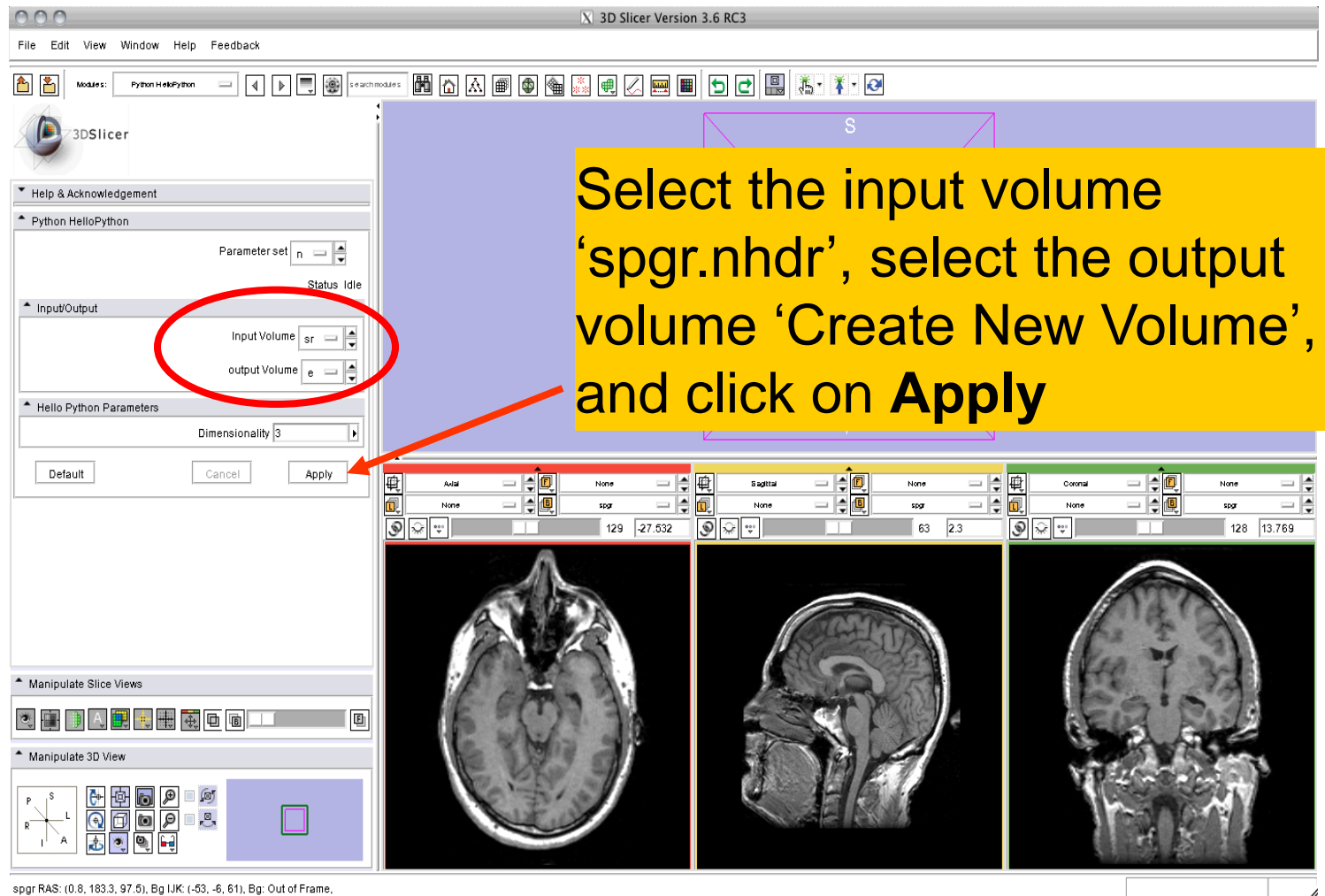


The screenshot shows the 3D Slicer 3.6 RC3 interface. The 'Modules' menu is open, and 'Python Hello Python' is selected, highlighted with a red circle. A yellow text box with black text is overlaid on the right side of the interface, containing the following instructions:

Browse to the Category **'Demonstration'** in the Modules menu, and select the module **'Python Hello Python'**

The interface also shows the 'Python Hello Python' module's parameter set, status, and input/output fields. The main 3D view displays three orthogonal MRI slices: Axial, Sagittal, and Coronal. The status bar at the bottom indicates the RAS coordinates for the selected slice: sgr1 RAS: (128.1, 12.8, -37.2), Bg IJK: (128, 138, 145), Bg: Out of Frame.

Running the Laplace Operator



3D Slicer Version 3.6 RC3

File Edit View Window Help Feedback

Modules: Python HelloPython

3DSlicer

Help & Acknowledgement

Python HelloPython

Parameter set: n

Status: Idle

Input/Output

Input Volume: **sr**

Output Volume: e

Hello Python Parameters

Dimensionality: 3

Default Cancel Apply

Manipulate Slice Views

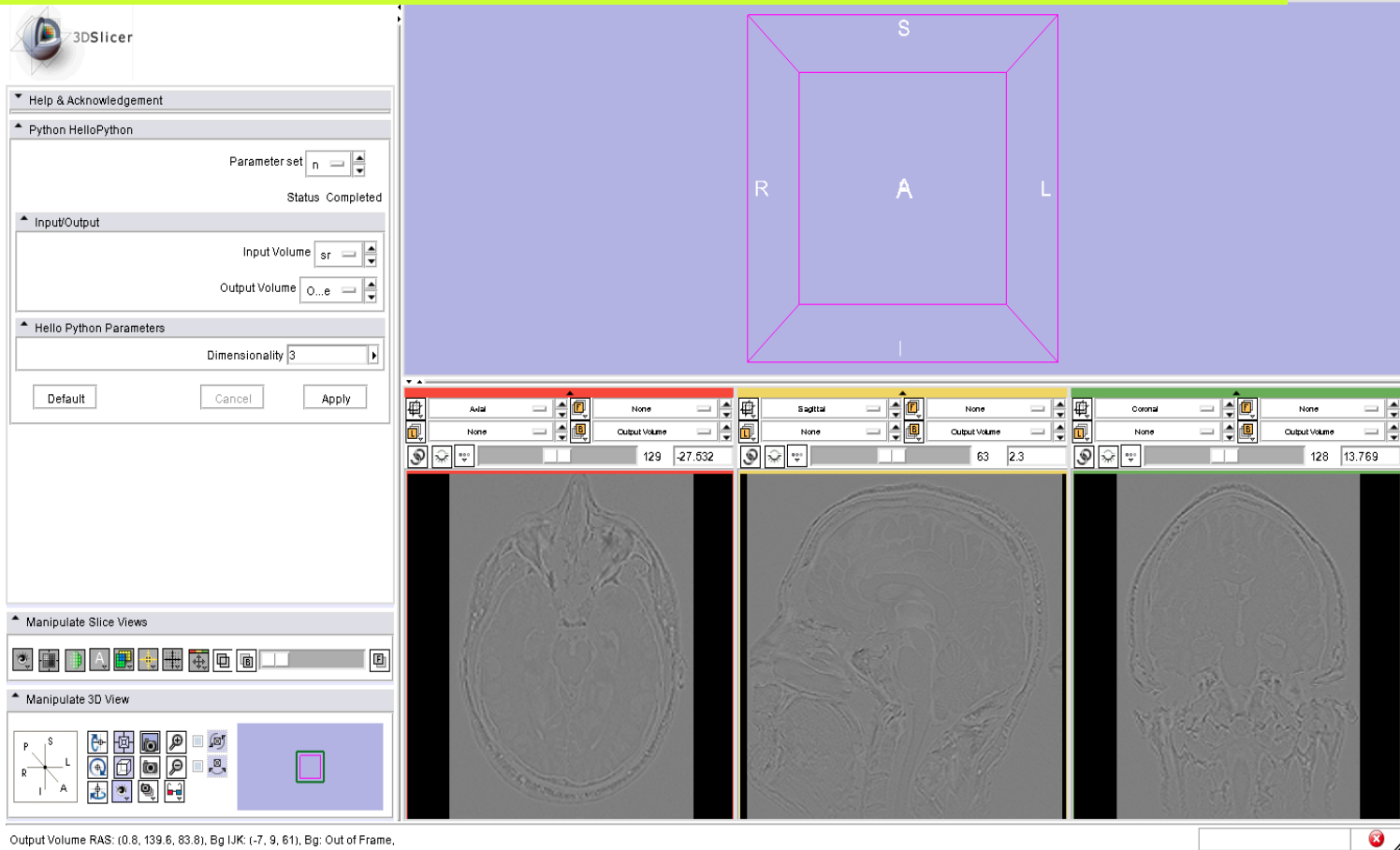
Manipulate 3D View

spgr RAS: (0.8, 183.3, 97.5), Bg IJK: (-63, -6, 61), Bg: Out of Frame.

Select the input volume 'spgr.nhdr', select the output volume 'Create New Volume', and click on **Apply**

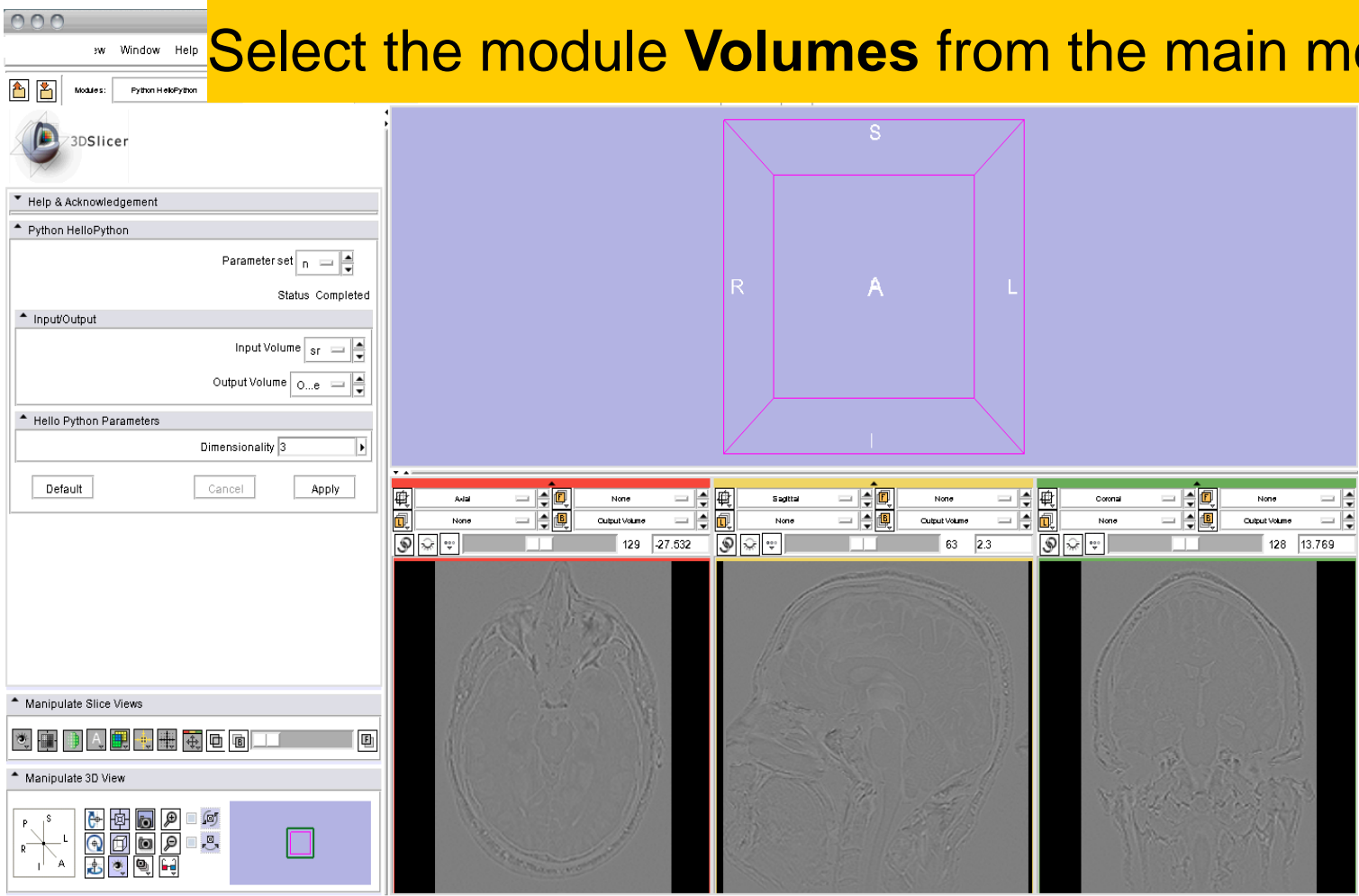
Running the Laplace Operator

Slicer displays the Laplacian of the spgr image.



Laplacian of the image

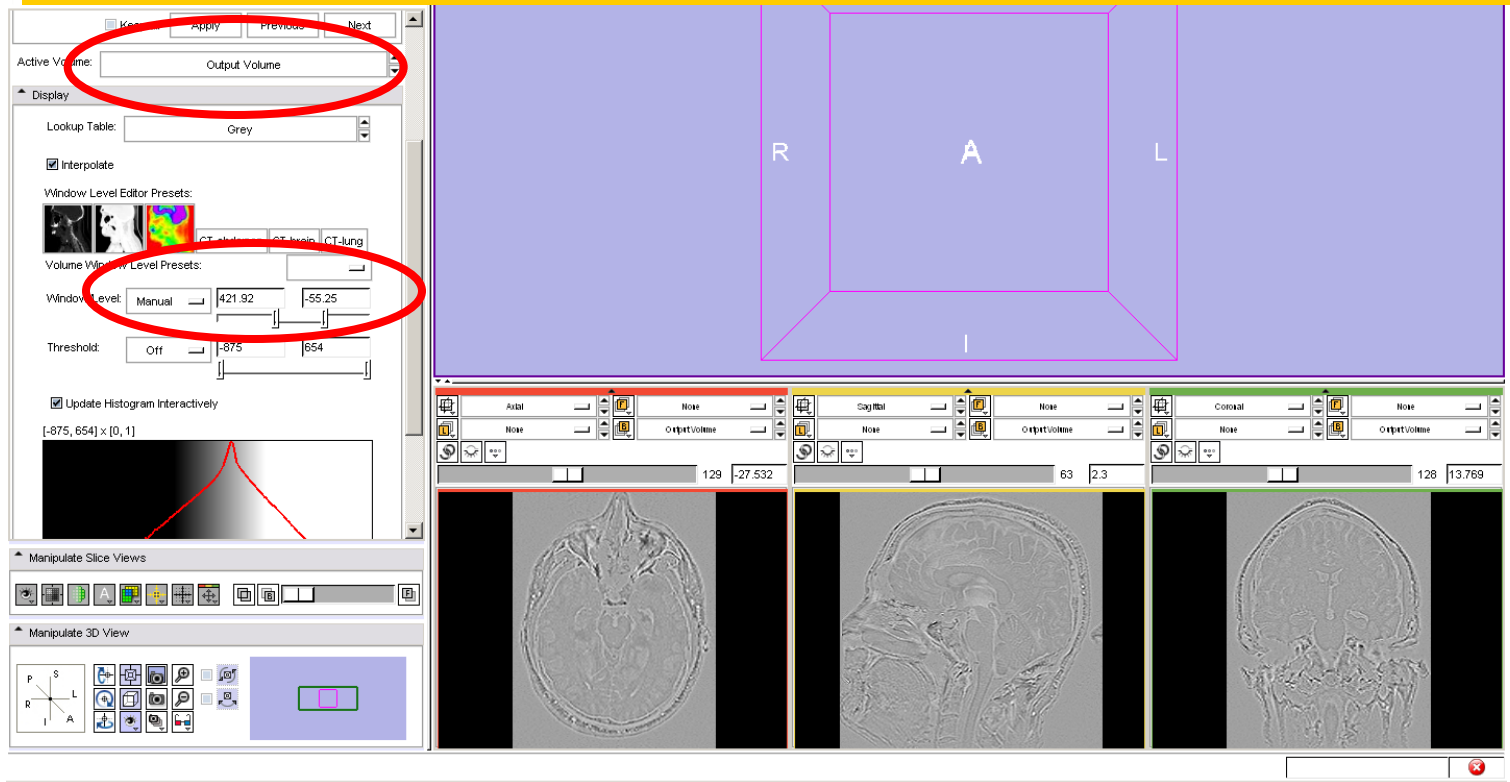
Select the module **Volumes** from the main menu



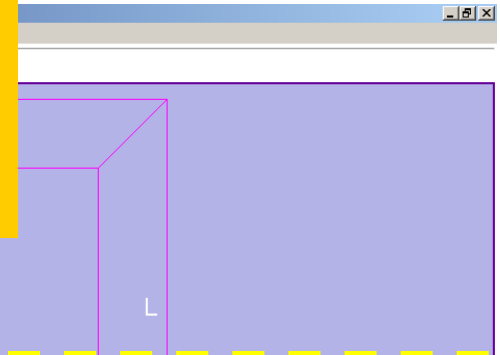
The screenshot shows the 3DSlicer application window. The main menu is open, and the 'Volumes' module is selected. The interface displays three orthogonal slice views: Axial, Sagittal, and Coronal. Each view shows a grayscale image of a brain slice with a purple rectangular region of interest (ROI) overlaid. The ROI is labeled with 'S' (Superior), 'I' (Inferior), 'R' (Right), and 'L' (Left). The 'Volumes' module parameters are visible on the left, showing 'Input Volume' set to 'sr' and 'Output Volume' set to 'O...e'. The status bar at the bottom indicates 'Output Volume RAS: (0.8, 139.6, 83.8), Bg IJK: (-7, 9, 61), Bg: Out of Frame.'

Laplacian of the image

Set the Active Volume to **Output Volume** and adjust the Window/Level parameters



Run the following code in the Python console to subtract the Laplacian of the image to the original image



```
import Slicer
volume1 = Slicer.slicer.MRMLScene.GetNodeByID("vtkMRMLScalarVolumeNode1")
volume2 = Slicer.slicer.MRMLScene.GetNodeByID("vtkMRMLScalarVolumeNode2")
plugin = Slicer.Plugin("Subtract Images")
plugin.Execute(volume1,volume2)
```

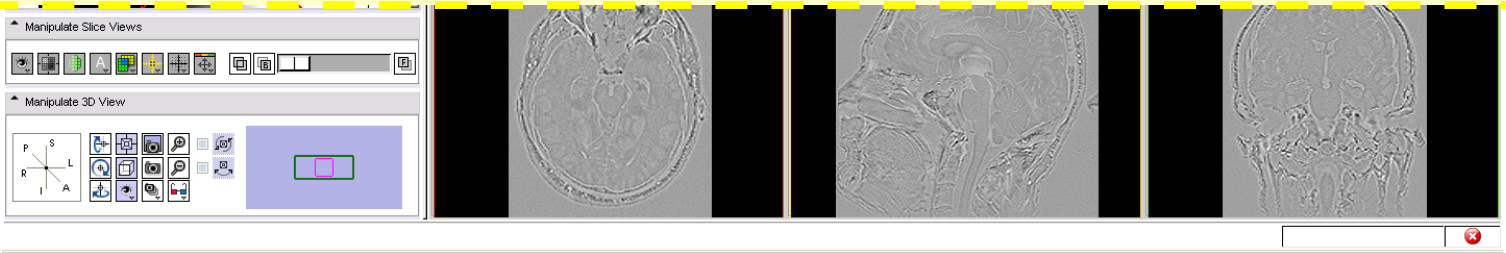
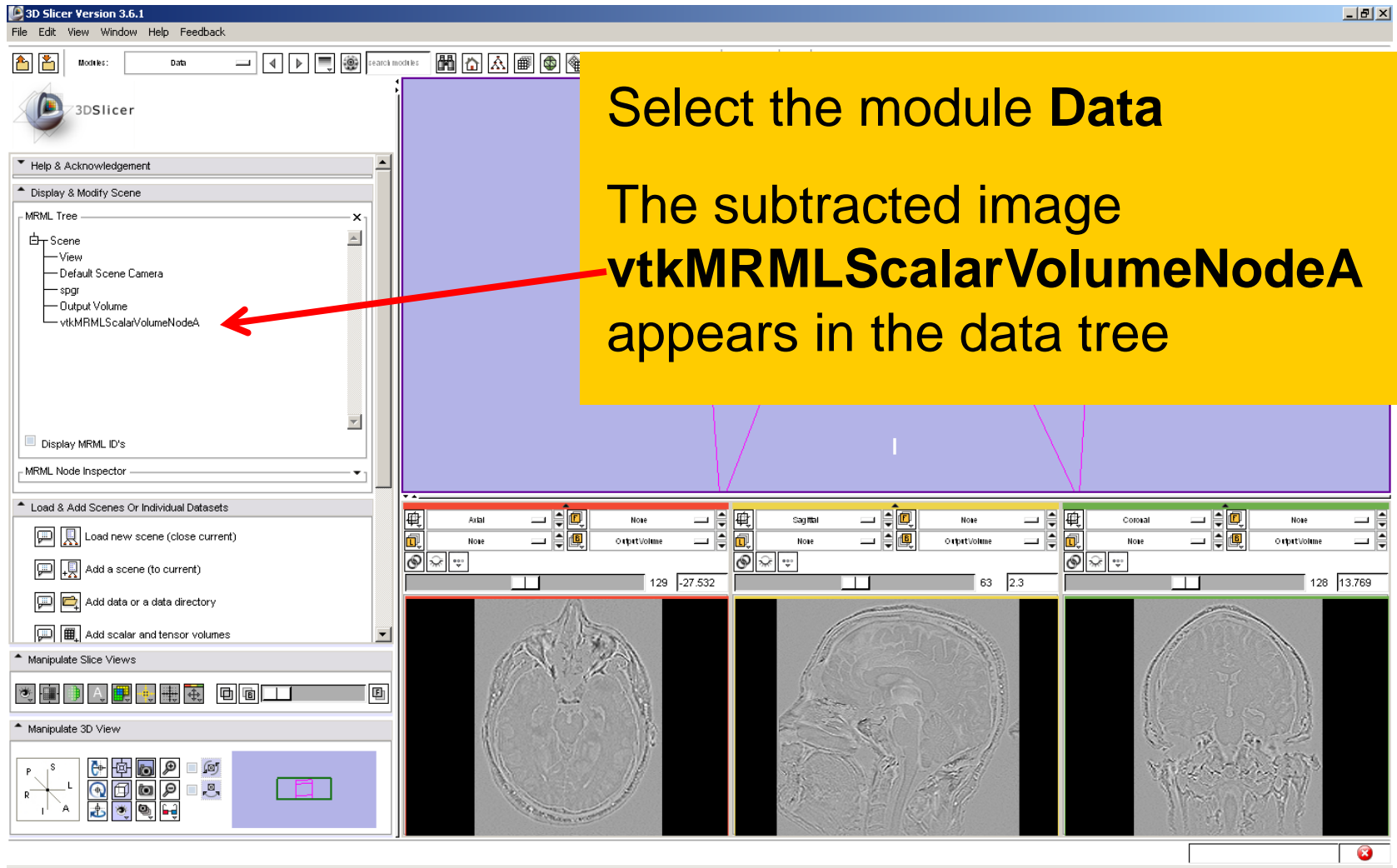


Image Sharpening



The screenshot shows the 3D Slicer 3.6.1 interface. The top menu bar includes File, Edit, View, Window, Help, and Feedback. Below the menu is a toolbar with various icons. The left sidebar contains several panels: 'Help & Acknowledgement', 'Display & Modify Scene', 'Load & Add Scenes Or Individual Datasets', 'Manipulate Slice Views', and 'Manipulate 3D View'. The 'Display & Modify Scene' panel shows the MRML Tree with the following structure:

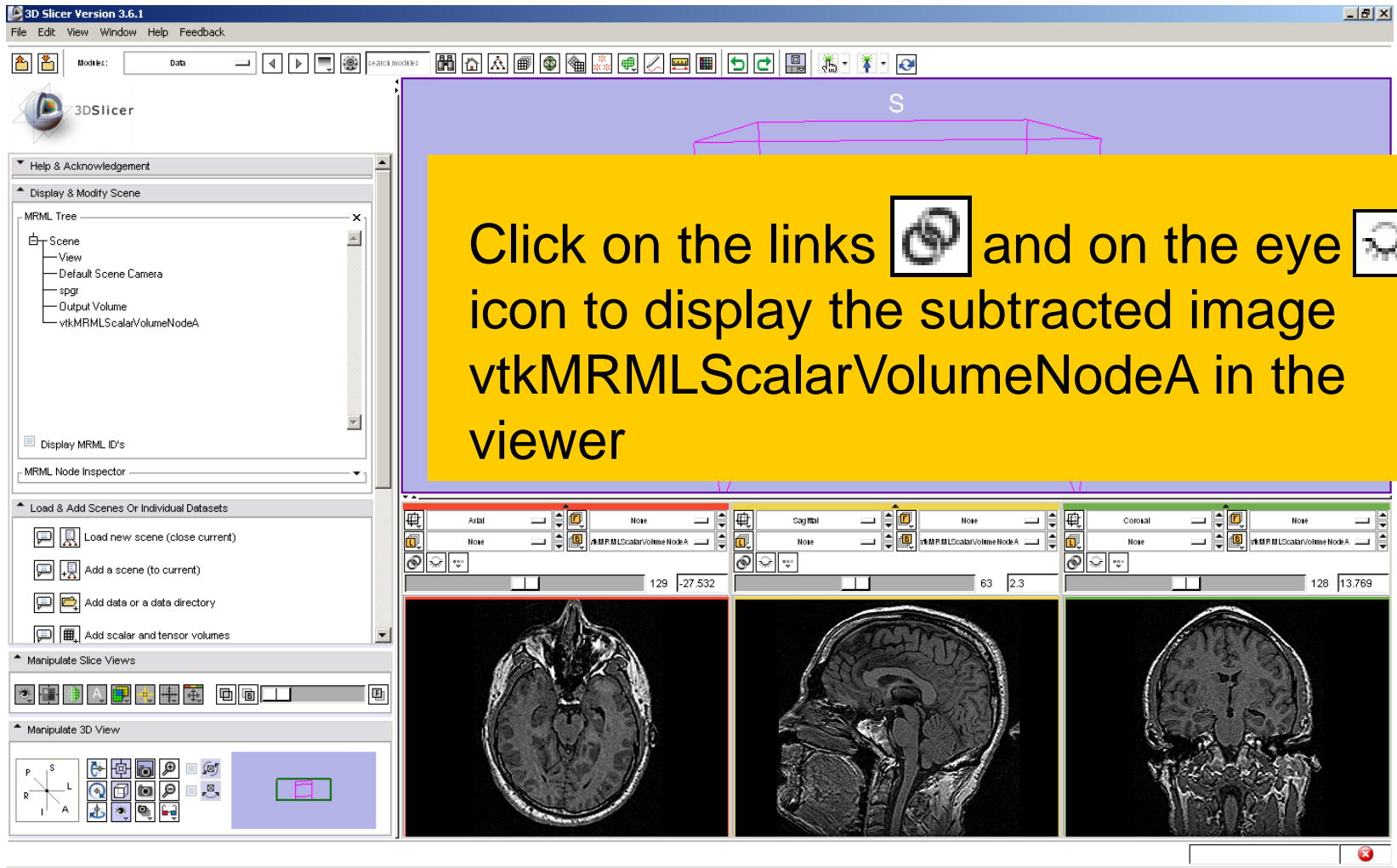
- Scene
 - View
 - Default Scene Camera
 - spgr
 - Output Volume
 - vtkMRMLScalarVolumeNodeA

A red arrow points to the 'vtkMRMLScalarVolumeNodeA' node in the tree. The 'Load & Add Scenes Or Individual Datasets' panel has buttons for 'Load new scene (close current)', 'Add a scene (to current)', 'Add data or a data directory', and 'Add scalar and tensor volumes'. The 'Manipulate Slice Views' panel has icons for 'None', 'Output Volume', and 'None' for Axial, Sagittal, and Coronal views. The 'Manipulate 3D View' panel has a 3D view icon and a small 3D view window. The main 3D view area shows three orthogonal slices: Axial, Sagittal, and Coronal. The Axial slice shows a brain cross-section with a red bounding box. The Sagittal slice shows a brain sagittal section with a yellow bounding box. The Coronal slice shows a brain coronal section with a green bounding box. The bottom status bar shows the coordinates for each slice: Axial (129, -27.532), Sagittal (63, 2.3), and Coronal (126, 13.769).

Select the module **Data**

The subtracted image **vtkMRMLScalarVolumeNodeA** appears in the data tree

Image Sharpening





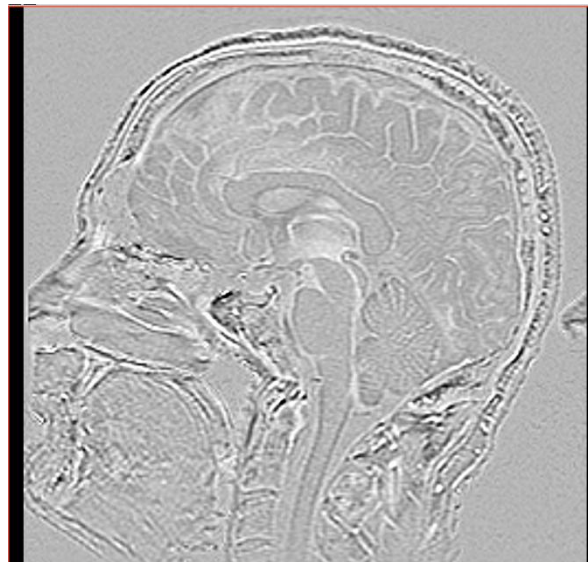
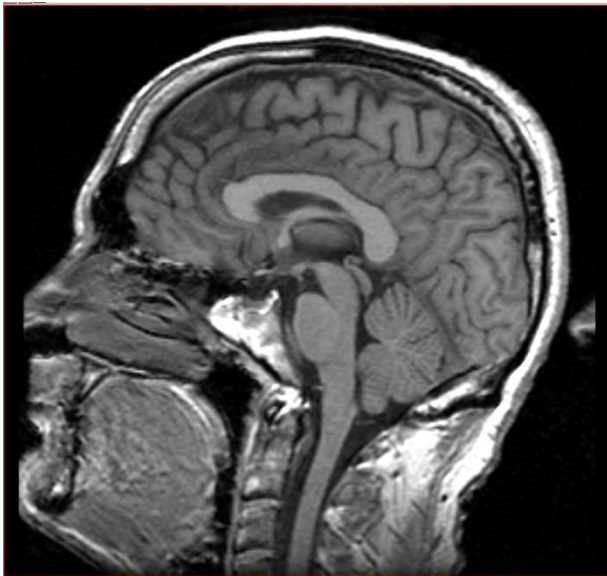
The screenshot shows the 3D Slicer 3.6.1 interface. On the left, the MRML Tree panel is visible, showing a hierarchy: Scene -> View -> Default Scene Camera -> spgr -> Output Volume -> vtkMRMLScalarVolumeNodeA. Below this, the 'Load & Add Scenes Or Individual Datasets' and 'Manipulate Slice Views' panels are partially visible. The main viewer area displays three orthogonal MRI slices: Axial, Sagittal, and Coronal. A yellow box is overlaid on the viewer area, containing the text: 'Click on the links  and on the eye  icon to display the subtracted image vtkMRMLScalarVolumeNodeA in the viewer'. A purple bracket labeled 'S' is positioned above the yellow box, indicating the slice position.

Image Sharpening

original

Laplacian

Laplacian filtered





Conclusion

- This course demonstrates how to integrate an external program in Python within Slicer3
- The **Execution Model** of Slicer3 provides a simple mechanism for incorporating command line programs as Slicer modules in Python.





Acknowledgments



National Alliance for Medical Image Computing

NIH U54EB005149



Neuroimage Analysis Center

NIH P41RR013218