



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

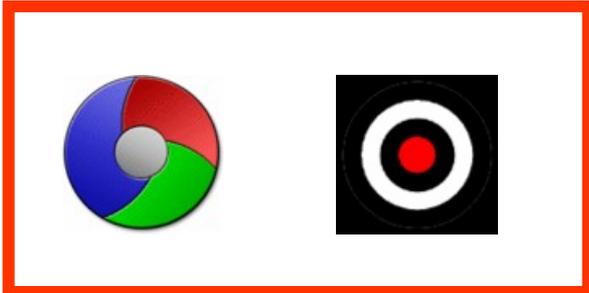
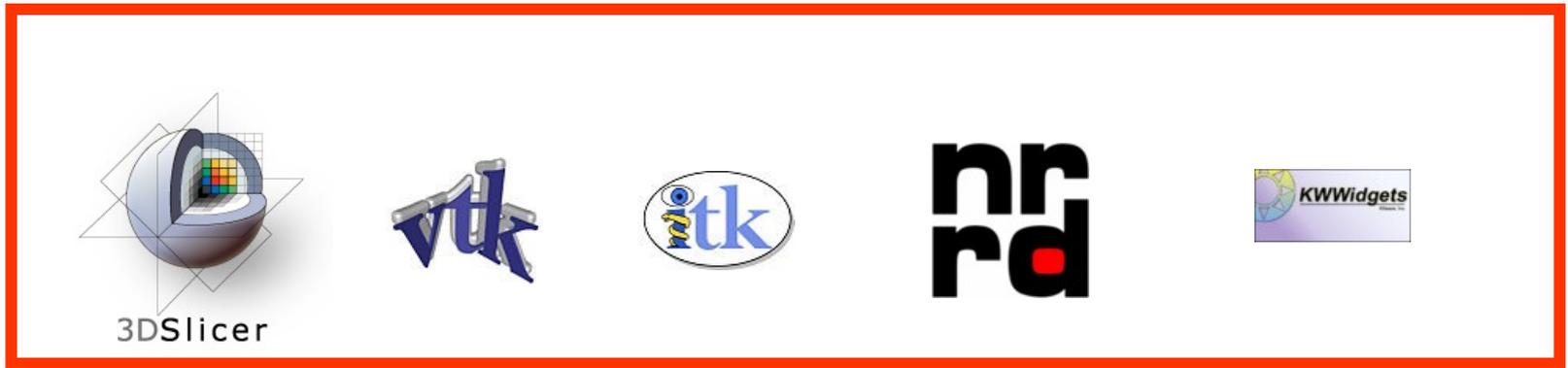
Programming into
Slicer3:

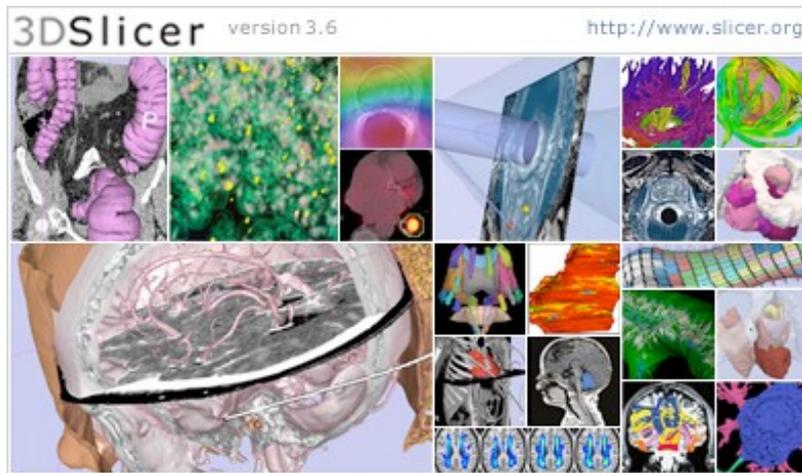
**The “Hello Python”
Tutorial**

Sonia Pujol, Ph.D.

Surgical Planning Laboratory
Harvard University

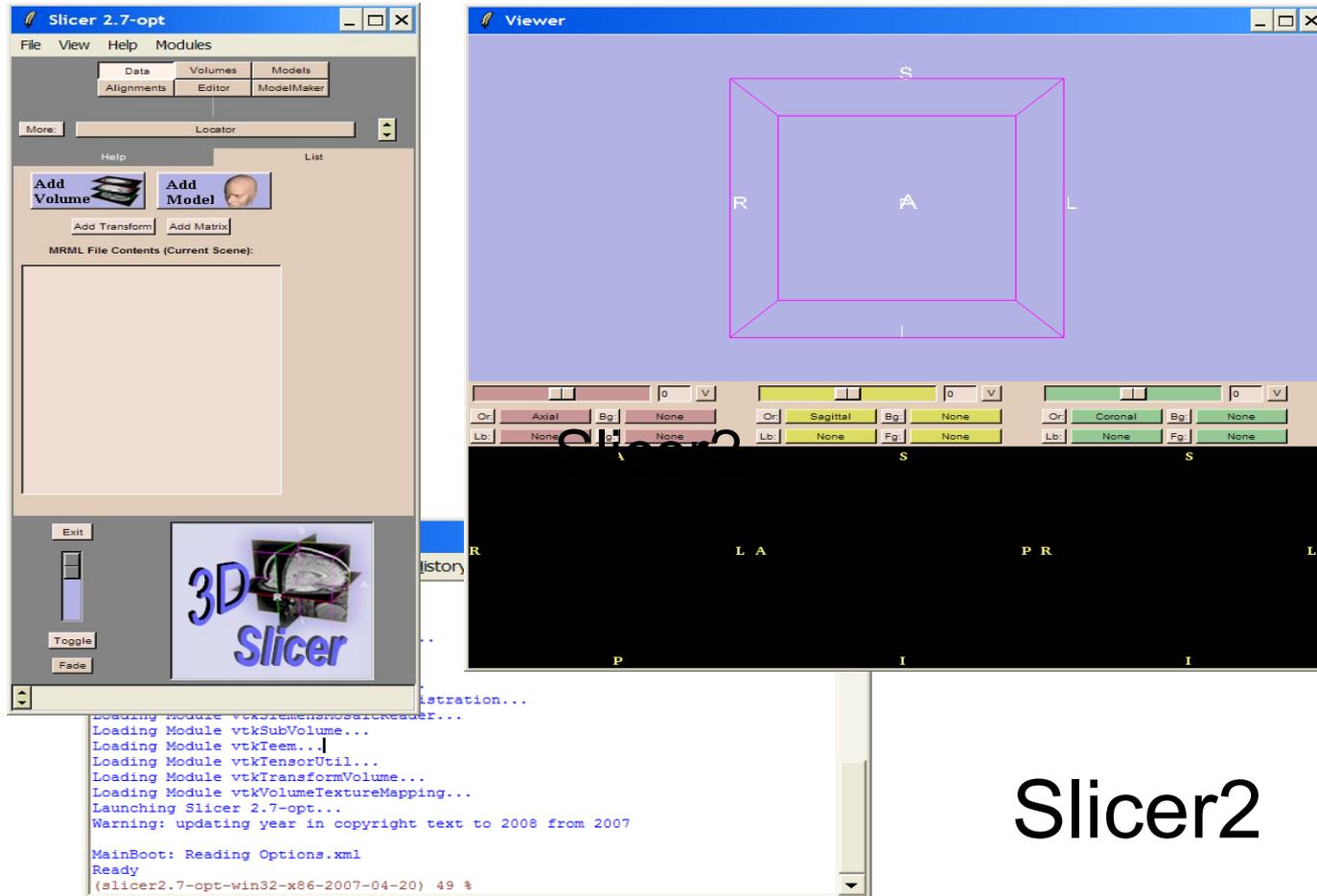
The NA-MIC Kit





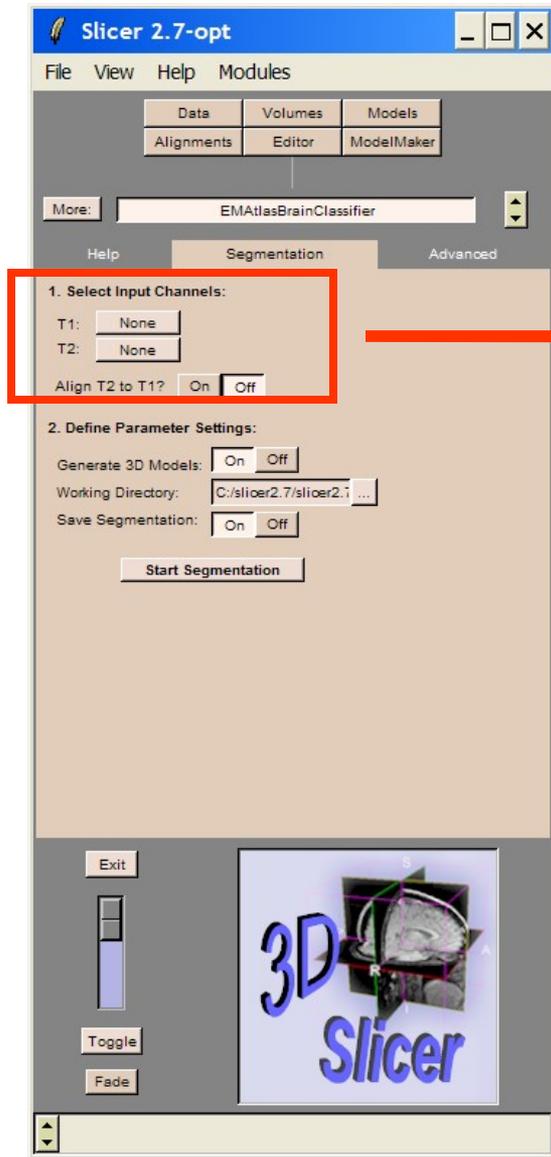
- An **end-user application** for image analysis
- An **open-source environment** for software development
- A software platform that is both **easy to use** for clinical researchers and **easy to extend** for programmers

Before Slicer3



Slicer2

Programming into Slicer2

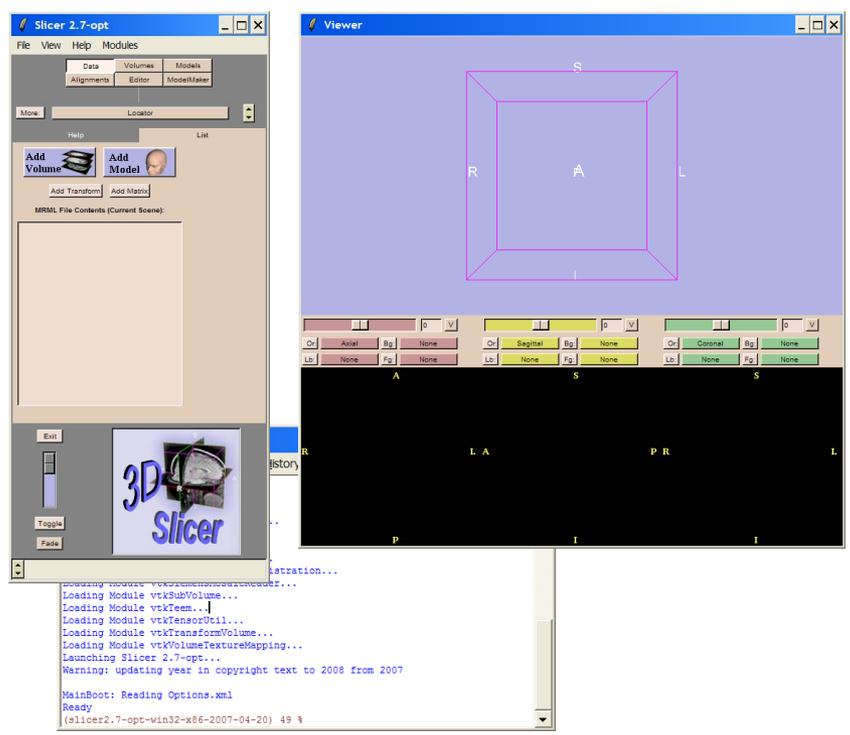


```

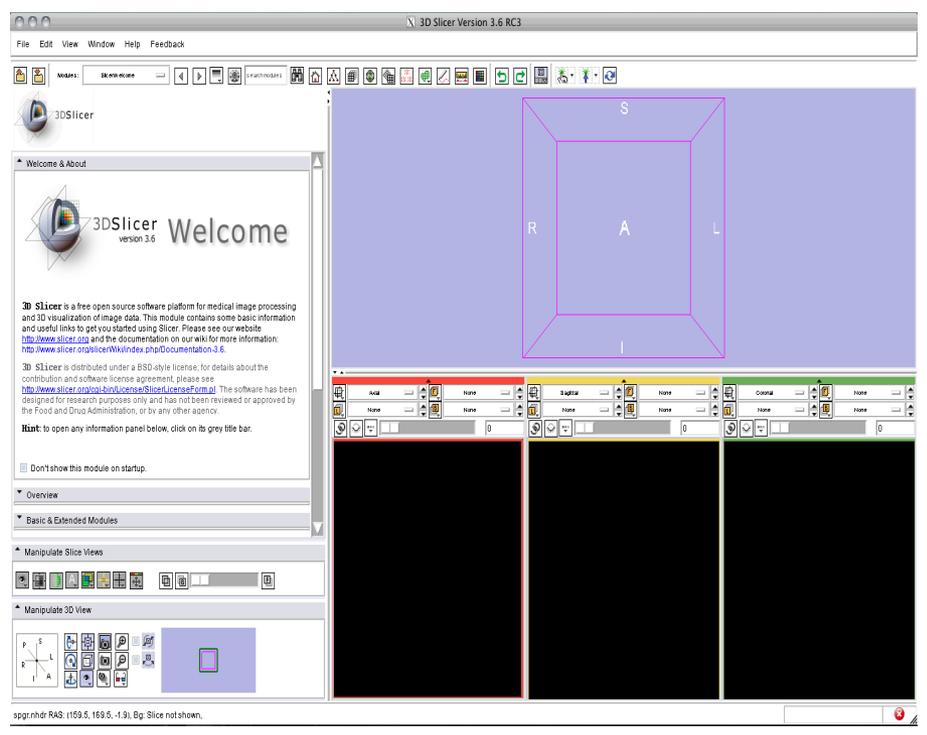
#-----
# 1. Step
#-----
set f $fSeg.fStep1
DevAddLabel $f.ITitle "1. Select Input Channels: " WTA
pack $f.ITitle -side top -padx $Gui(pad) -pady 1 -anchor w
frame $f.fInput -bg $Gui(activeWorkspace)
pack $f.fInput -side top -padx 0 -pady 0 -anchor w
foreach frame "Left Right" {
    frame $f.fInput.f$frame -bg $Gui(activeWorkspace)
    pack $f.fInput.f$frame -side left -padx 0 -pady $Gui(pad) }
foreach LABEL "T1 T2" Input "SPGR T2W" {
    DevAddLabel $f.fInput.fLeft.I$Input " ${LABEL}:"
    pack $f.fInput.fLeft.I$Input -side top -padx $Gui(pad) -pady 1 -anchor w
    set menubutton $f.fInput.fRight.m${Input}Select
    set menu $f.fInput.fRight.m${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.fAlign -bg $Gui(activeWorkspace)
TooltipAdd $f.fAlign "If the input T1 and T2 are not aligned with each other set flag here"
pack $f.fAlign -side top -padx 0 -pady 2 -padx $Gui(pad) -anchor w
DevAddLabel $f.fAlign.IAlign "Align T2 to T1? "
pack $f.fAlign.IAlign -side left -padx $Gui(pad) -pady 1 -anchor w
foreach value "1 0" text "On Off" width "4 4" {
    eval {radiobutton $f.fAlign.r$value -width $width -indicatoron 0
        -text "$text" -value "$value" -variable EMAtlasBrainClassifier(AlignInput) } $Gui(WCA)
    pack $f.fAlign.r$value -side left -padx 0 -pady 0 }
  
```

From Slicer2 to Slicer3



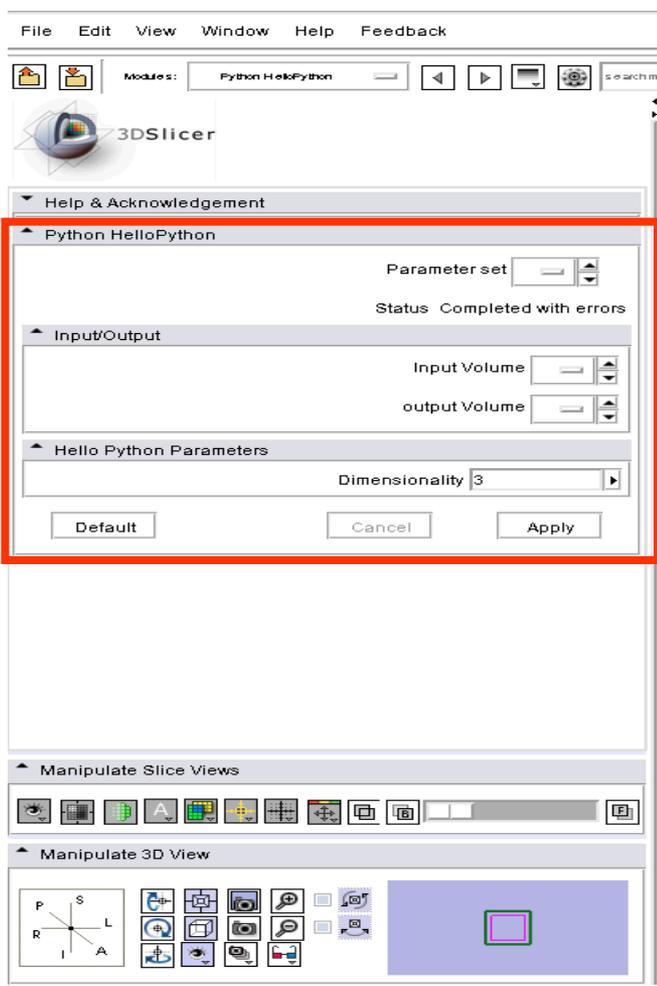
Slicer2



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>

```



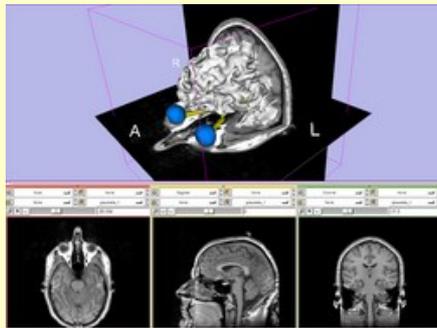
Slicer3 Execution Model

- 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.



Pre-requisite

- This course supposes that you have taken the following tutorial:



Slicer3 Data Loading and Visualization, Sonia Pujol Ph.D.

- The tutorial is available on the Slicer3.6 101 compendium:
http://www.slicer.org/slicerWiki/index.php/Slicer3.6:Training#Software_tutorials



Material

This course requires the following material

- Slicer3-3.6.1 release version

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

- HelloPython.zip

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

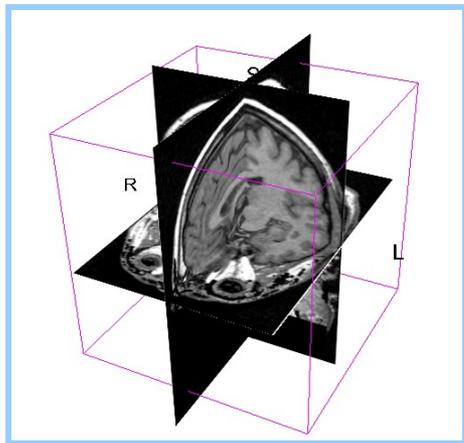
Disclaimer

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.



HelloPython Course Material

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>
</executable>
"""
```

HelloPython.py

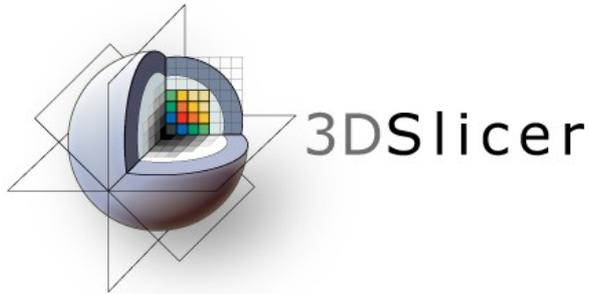


Overview

- 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



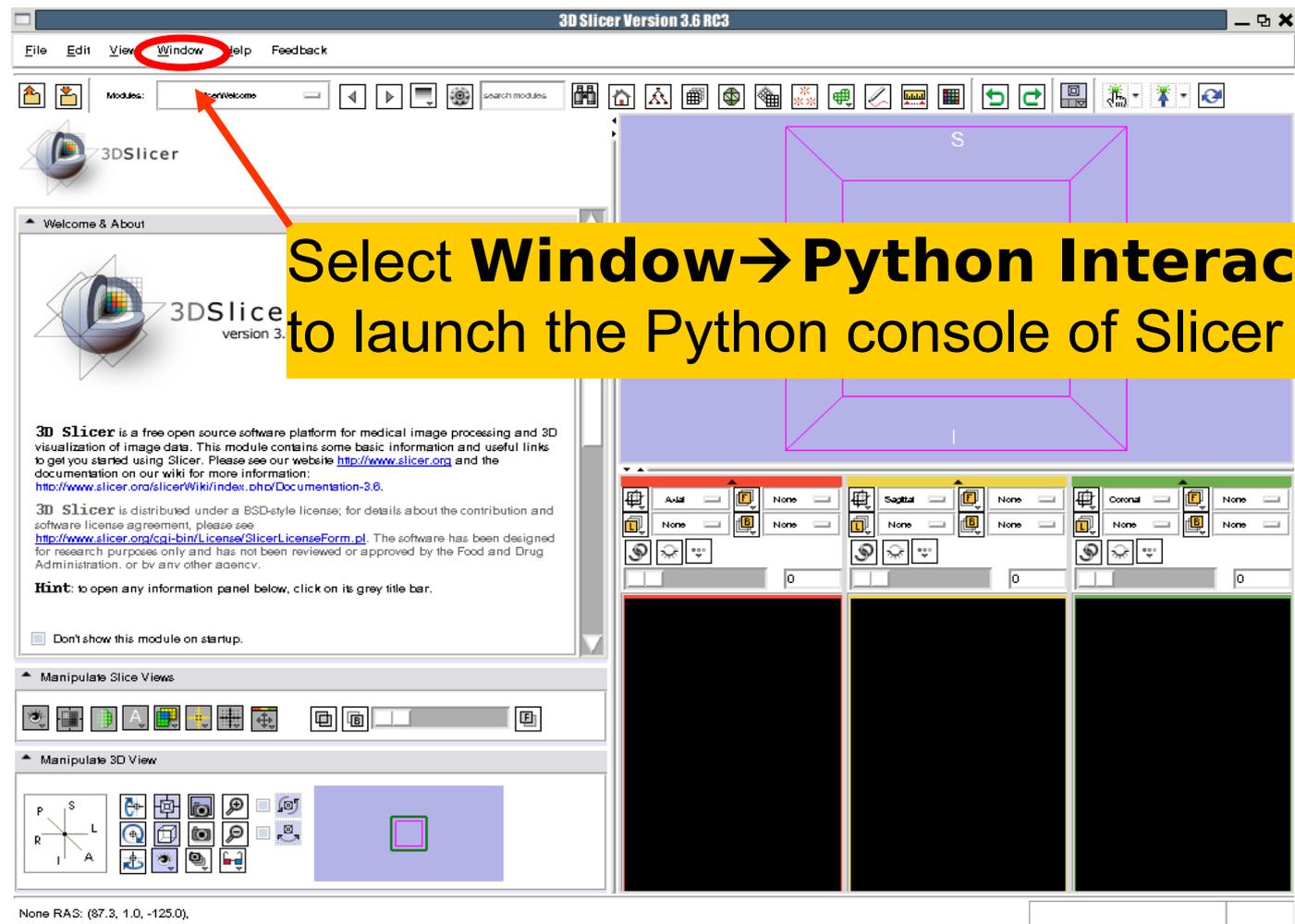
```
#!/usr/bin/env python
XML = """<?xml version="1.0" encoding="utf-8">
<executable>
  <category>Demonstration </category>
  <title>Python HelloPythons</title>
  <description> Slicer developer course in Python
  </description>
  <version>1.0</version>

  <license></license>
  <contributors> 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>

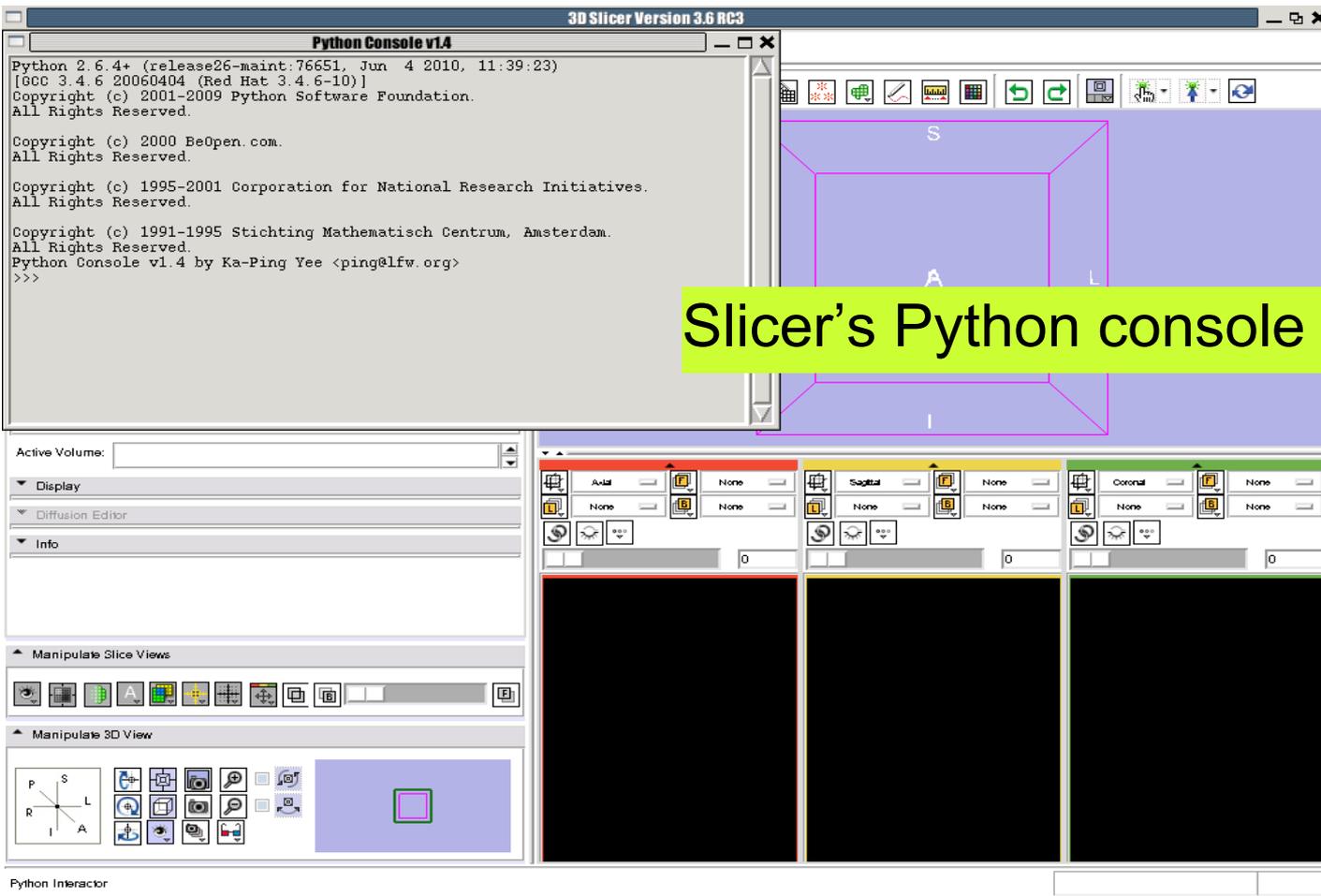
  <parameters>
    <label>Input/Output</label>
    <description>Input/output parameters</description>
    <image>
      <name>helloPythonInputVolume</name>
      <label>Input Volume</label>
      <channel>input</channel>
      <index>0</index>
      <description>input volume</description>
    </image>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

Part A: Integrating HelloPython into Slicer3

Python Console

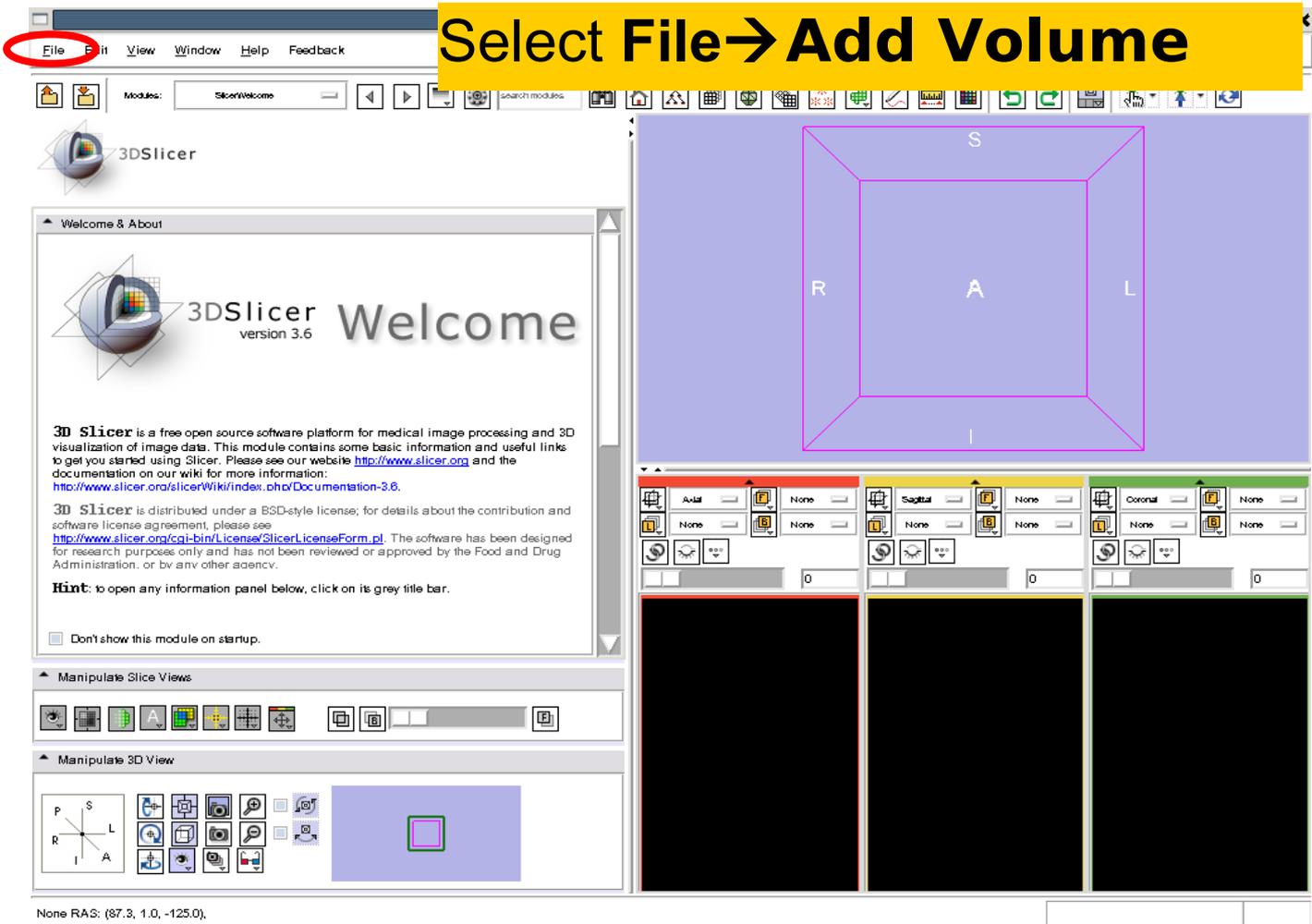


Python Console



Python Console

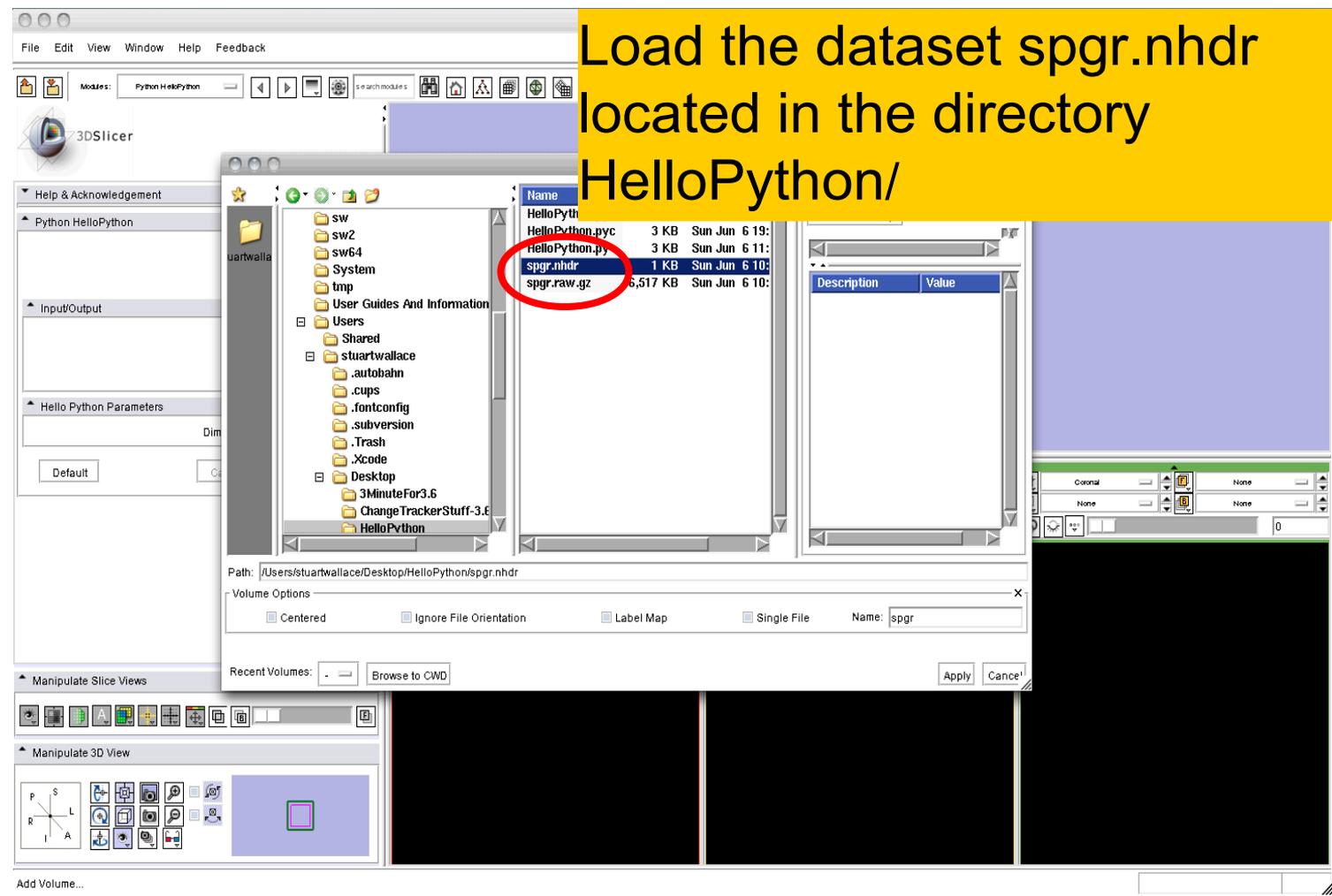
Select File → Add Volume



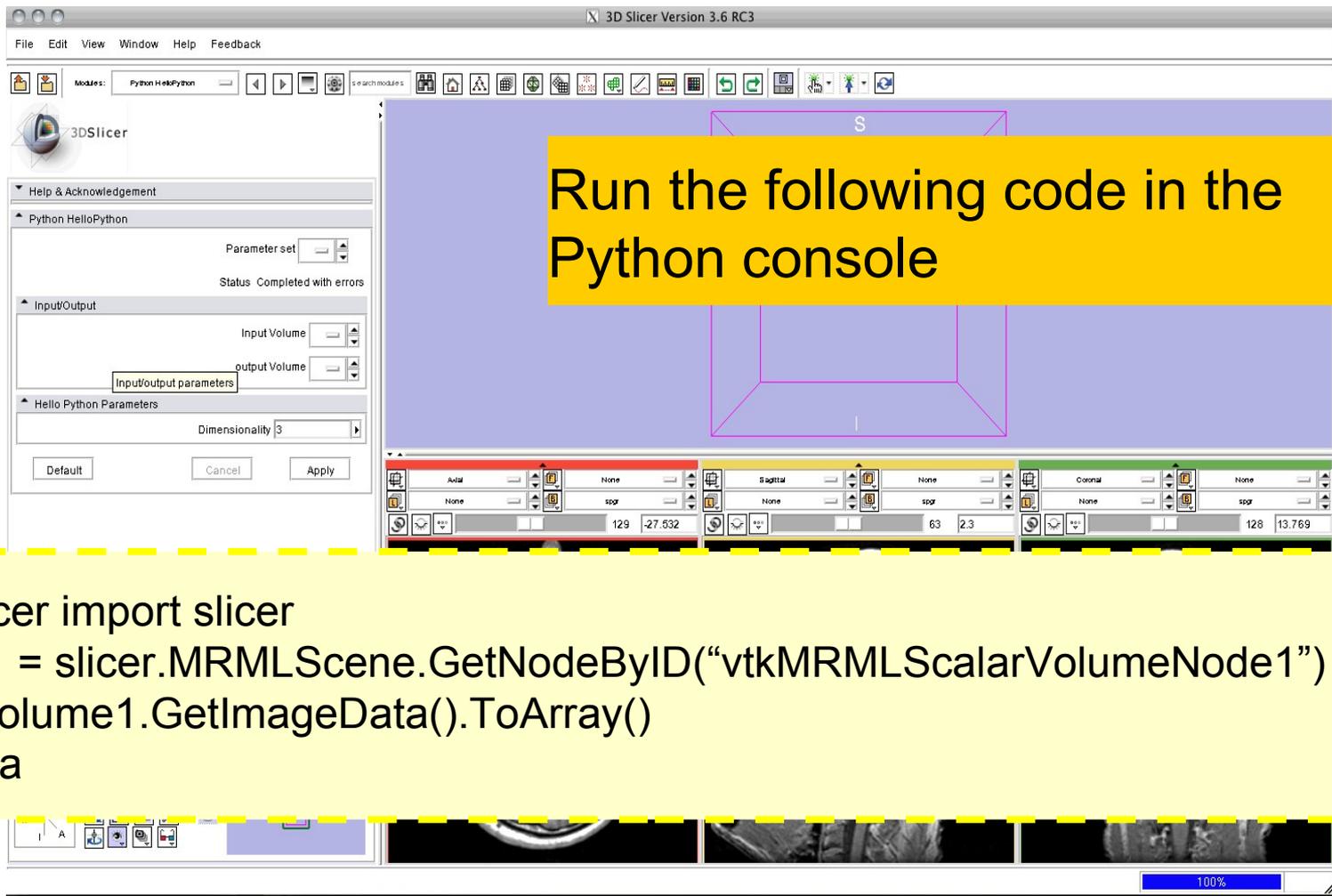
The screenshot shows the 3DSlicer v3.6 interface. The 'File' menu is highlighted with a red circle. The 'Add Volume' button (represented by a folder icon) is highlighted in the top toolbar. The main 3D view area is currently empty, showing a purple grid with axes labeled R (Right), L (Left), S (Superior), and I (Inferior). The 'Welcome & About' panel is open on the left, displaying the 3DSlicer logo and version information. Below it are sections for 'Manipulate Slice Views' and 'Manipulate 3D View'.

Python Console

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

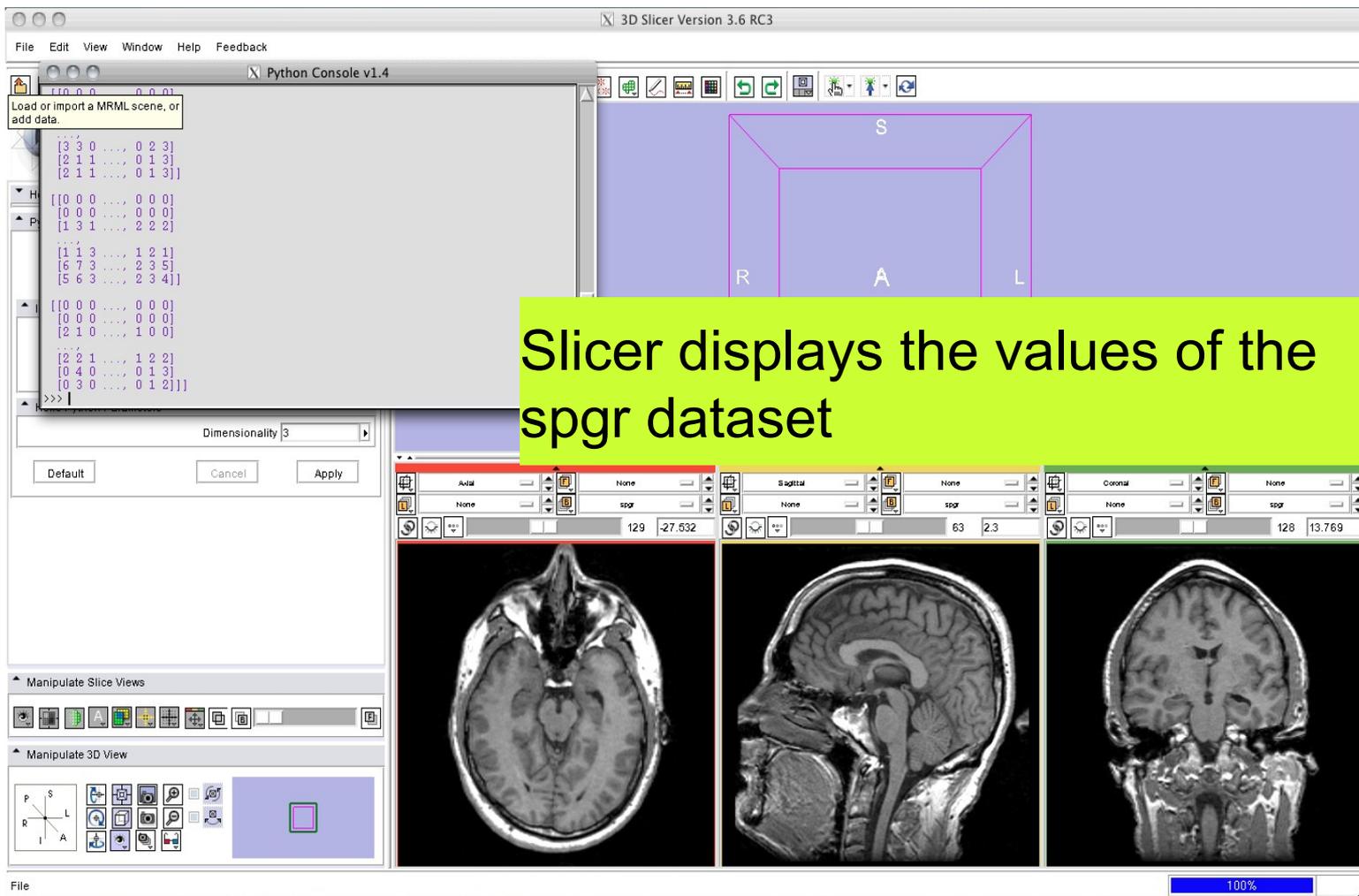


Python Console

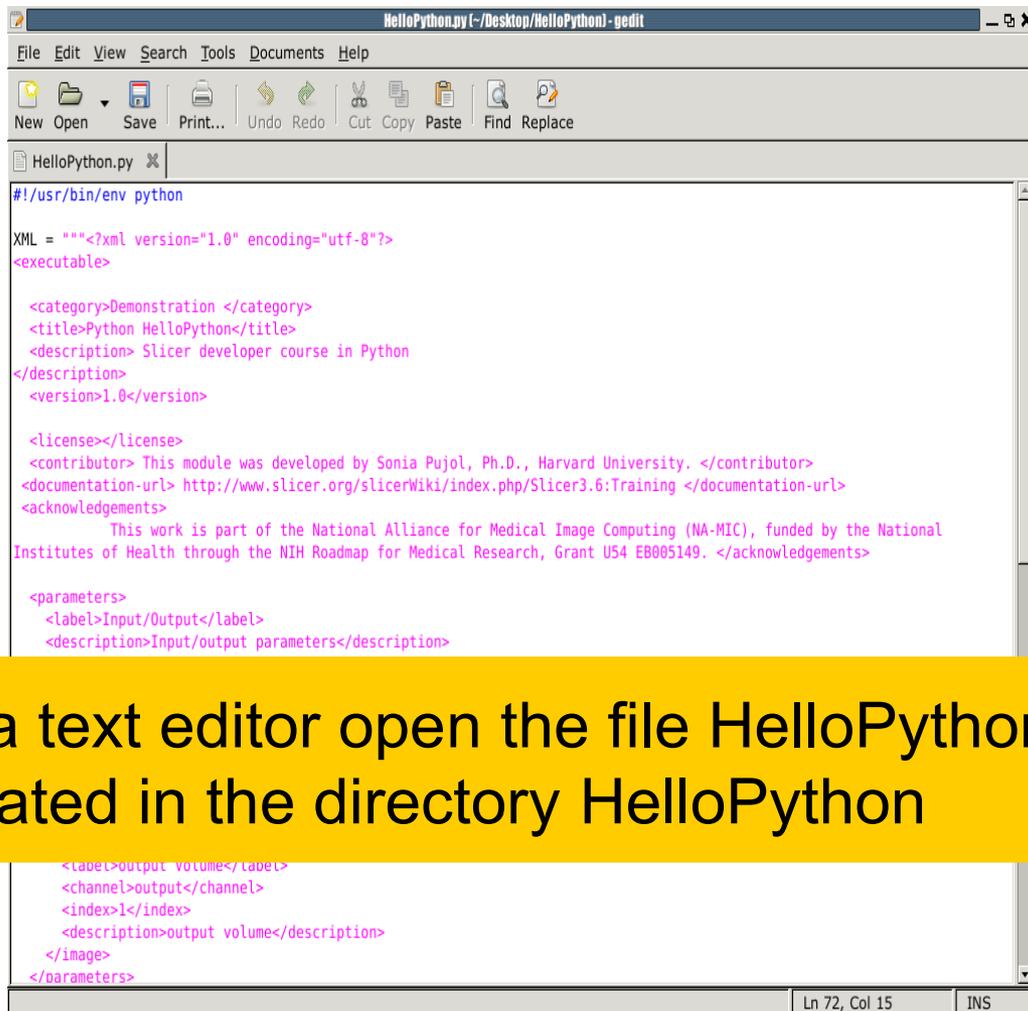


```
from Slicer import slicer
volume1 = slicer.MRMLScene.GetNodeByID("vtkMRMLScalarVolumeNode1")
data = volume1.GetImageData().ToArray()
print data
```

Python Console



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 U54 EB005149. </acknowledgements>

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

    <label>output volume</label>
    <channel>output</channel>
    <index>1</index>
    <description>output volume</description>
  </image>
</parameters>

```

In a text editor open the file HelloPython.py located in the directory HelloPython

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>
</executable>

<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.MRMLScene

    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
```



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

The screenshot shows the 3D Slicer 3.6 RC3 interface. A yellow callout box with the text "Select Module Settings from the Application Settings GUI" is overlaid on the top right. The "Application Settings" dialog is open, showing the "Module Settings" tab. Under "Module Settings", the "Select Modules..." button is visible. Below it, a list of directories is shown, with the following table:

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

The "Python HelloPython" module settings are also visible in the background, showing "Input/Output" and "Hello Python Parameters" sections. The main 3D view shows a coronal MRI slice of a brain.



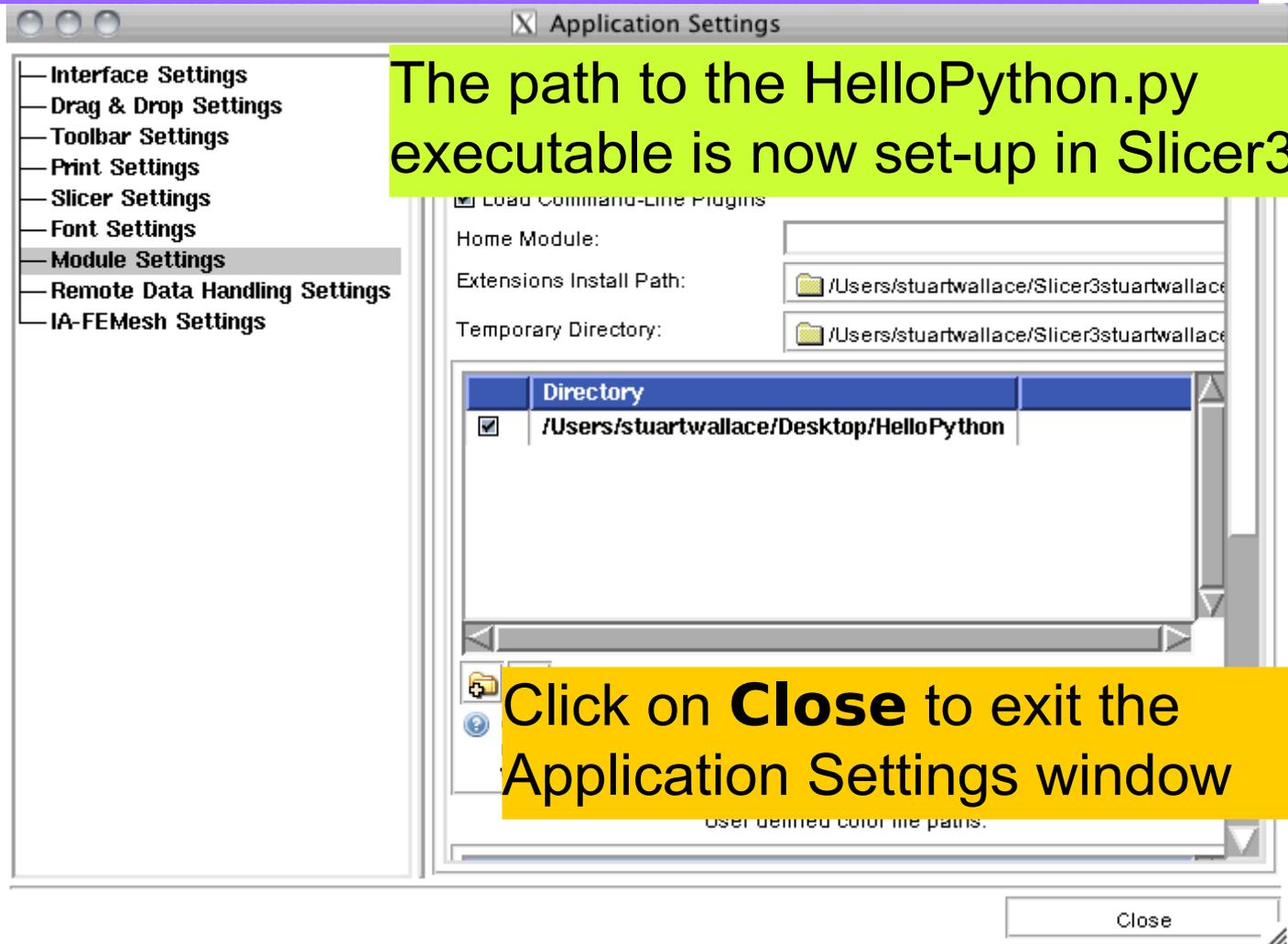
Integrating HelloPython to Slicer3

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

The screenshot shows the Slicer3 application window with the 'Application Settings' dialog box open. The 'Module Settings' tab is selected, and the 'Add a preset' button (represented by a folder icon) is circled in red. The 'Directory' list contains the path `/Users/stuartwallace/Desktop/HelloPython`. The background shows the Slicer3 interface with a 3D view of a brain scan and various toolbars.



Integrating HelloPython to Slicer3





Integrating HelloPython to Slicer3

Select **File** → **Exit** and re-start Slicer

3DSlicer version 3.6

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>.

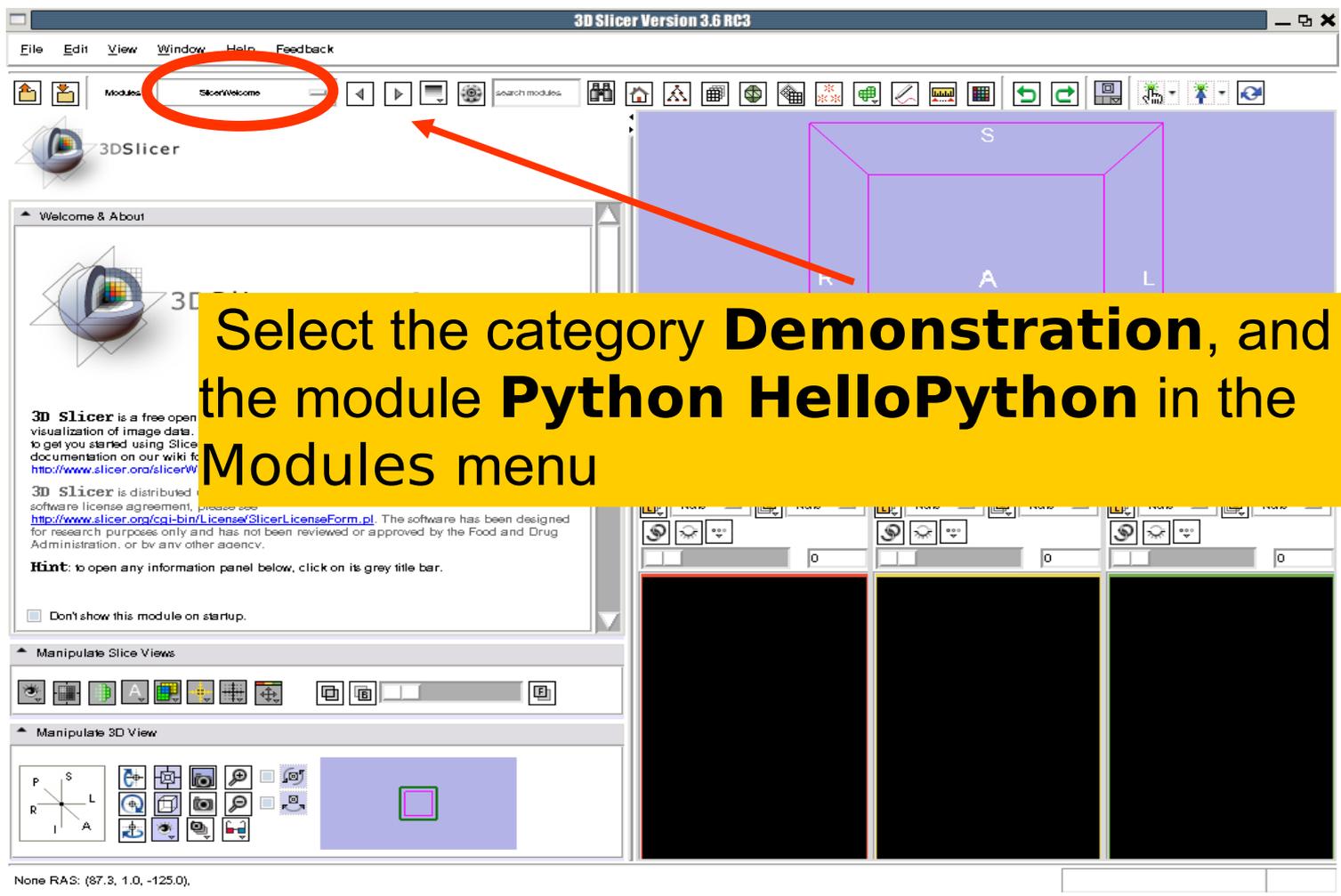
3D Slicer is distributed under a BSD-style license; for details about the contribution and software license agreement, please see <http://www.slicer.org/cgi-bin/license/SlicerLicenseForm.pl>. The software has been designed for research purposes only and has not been reviewed or approved by the Food and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

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

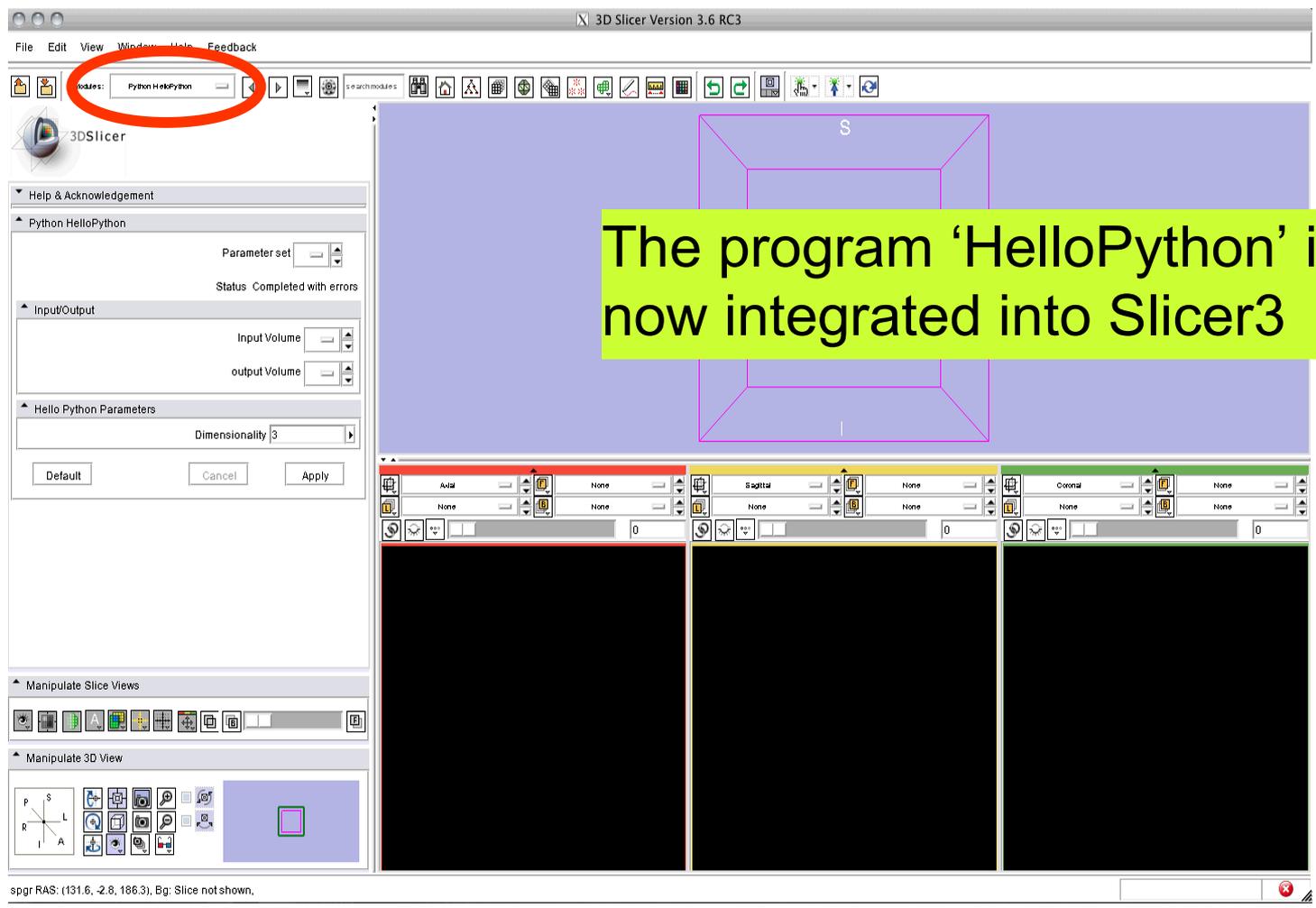
Don't show this module on startup.

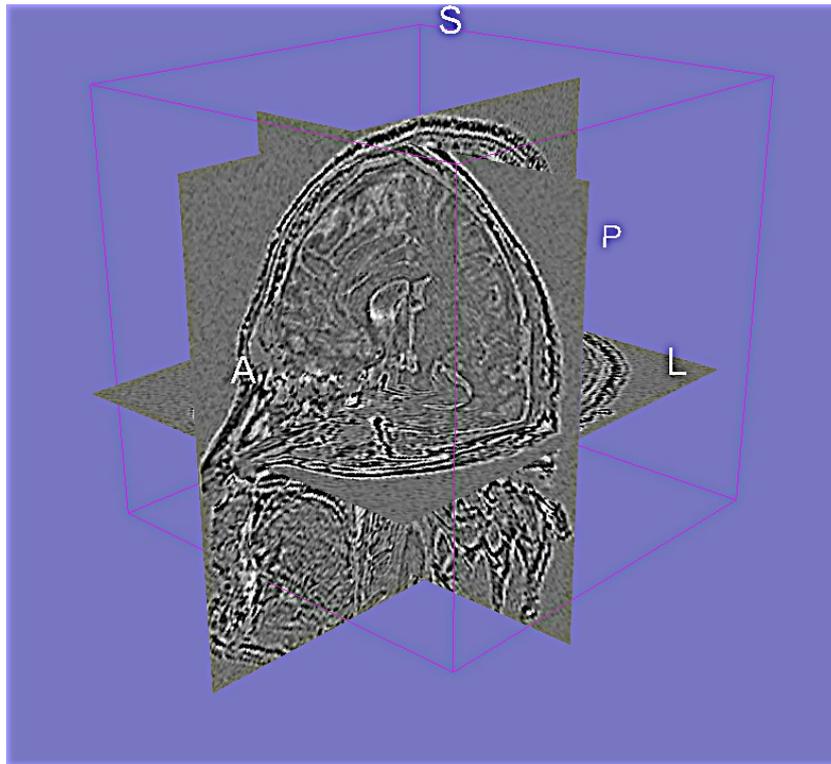
None RAS: (87.3, 1.0, -125.0)

HelloPython module



HelloPython Module





Part B: Implementing the Laplace* Operator

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



Execute Function

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

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

```
    slicer = Slicer.slicer
```

```
    scene = slicer.MRMLScene
```

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

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

```
    return
```

Add the I/O code

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
```

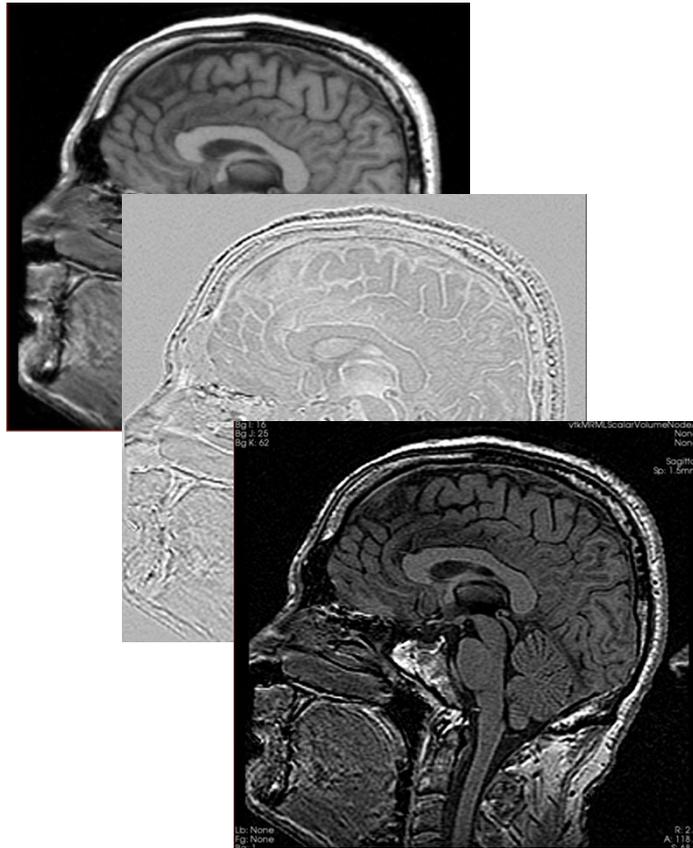
Place back the Laplacian of the image in the RAS reference system.



Integrating HelloPython to Slicer3

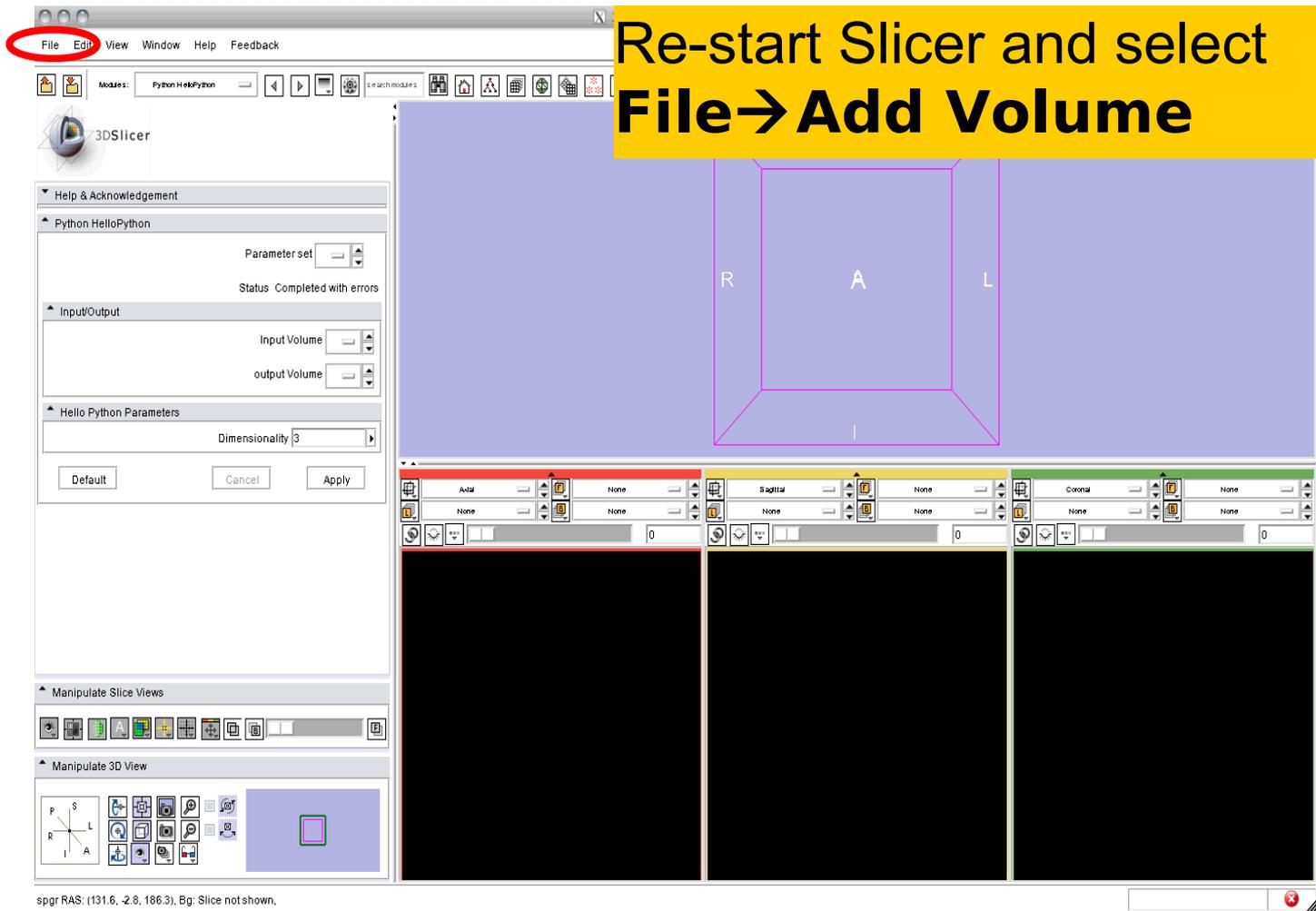
```
def Execute (HelloPythonInputVolume, HelloPythonOutputVolume,
dimensionality=3):
    Slicer = __import__("Slicer")
    slicer = Slicer.slicer
    scene = slicer.MRMLScene
    inputVolume = scene.GetNodeByID(HelloPythonInputVolume)
    outputVolume = scene.GetNodeByID(HelloPythonOutputVolume)
    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

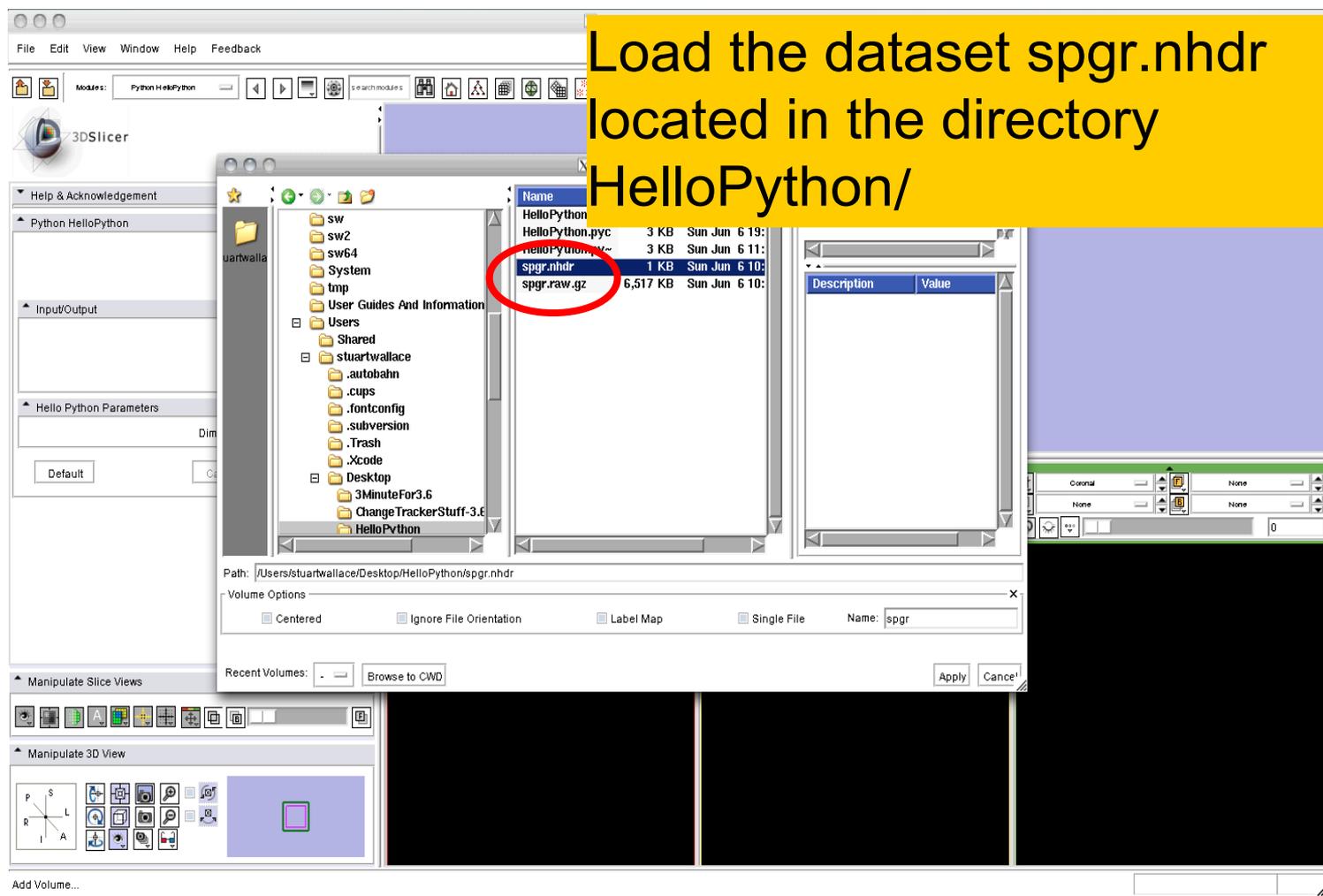
Running the Laplace Operator



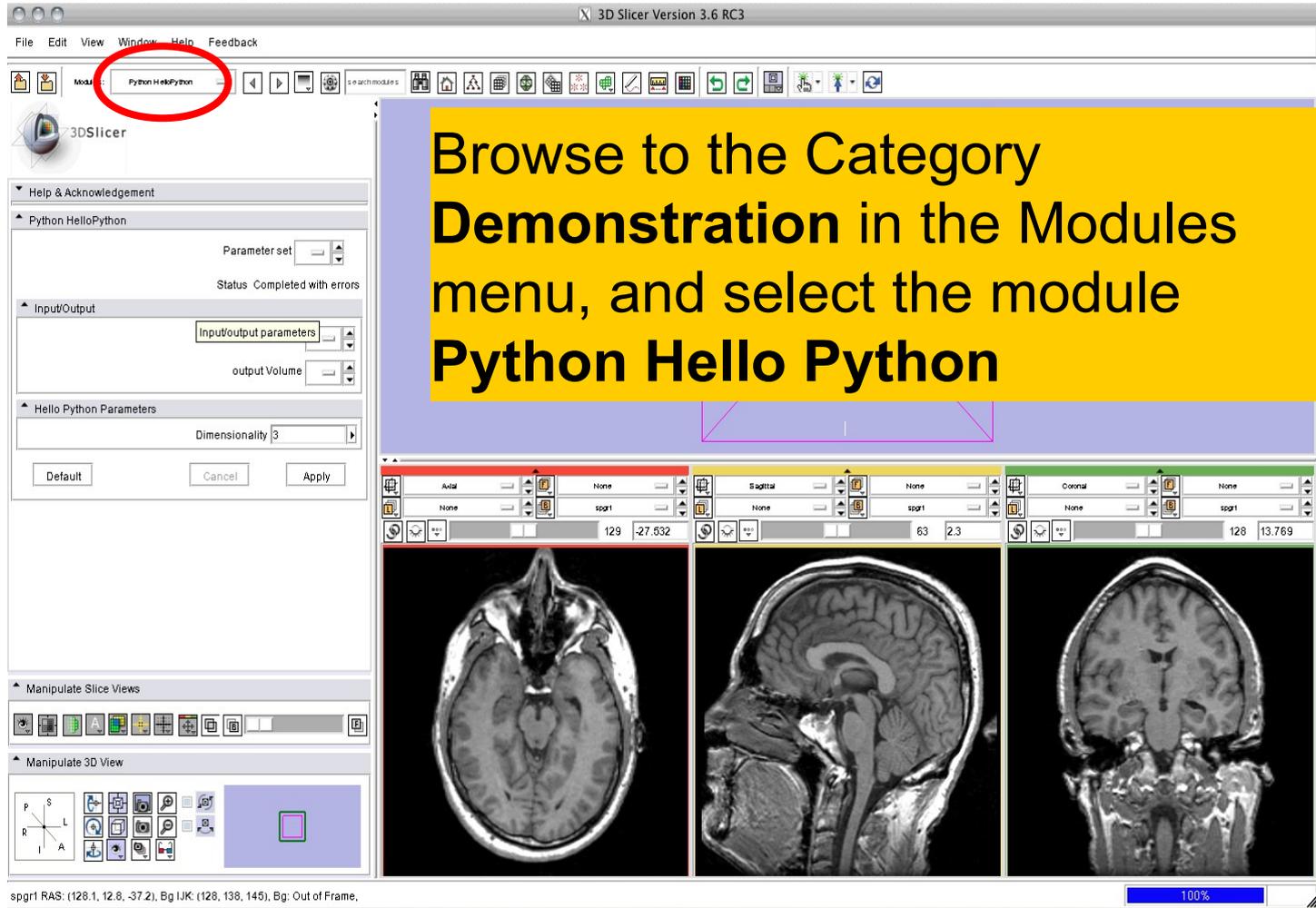
The screenshot shows the 3DSlicer application window. The 'File' menu is circled in red. A yellow callout box with black text reads: "Re-start Slicer and select **File**→**Add Volume**". The main 3D view area is a light purple color with a pink wireframe box in the center, labeled with 'R', 'A', 'L', and 'I' at its corners. Below the 3D view are three slice view panels: Axial (red header), Sagittal (yellow header), and Coronal (green header). Each panel shows a black slice view with a white crosshair and a numerical value of '0'. 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' dropdowns), 'Hello Python Parameters' (with 'Dimensionality' set to '3'), 'Manipulate Slice Views', and 'Manipulate 3D View' (with a 3D orientation diagram and a small 3D view window). The status bar at the bottom left shows '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/

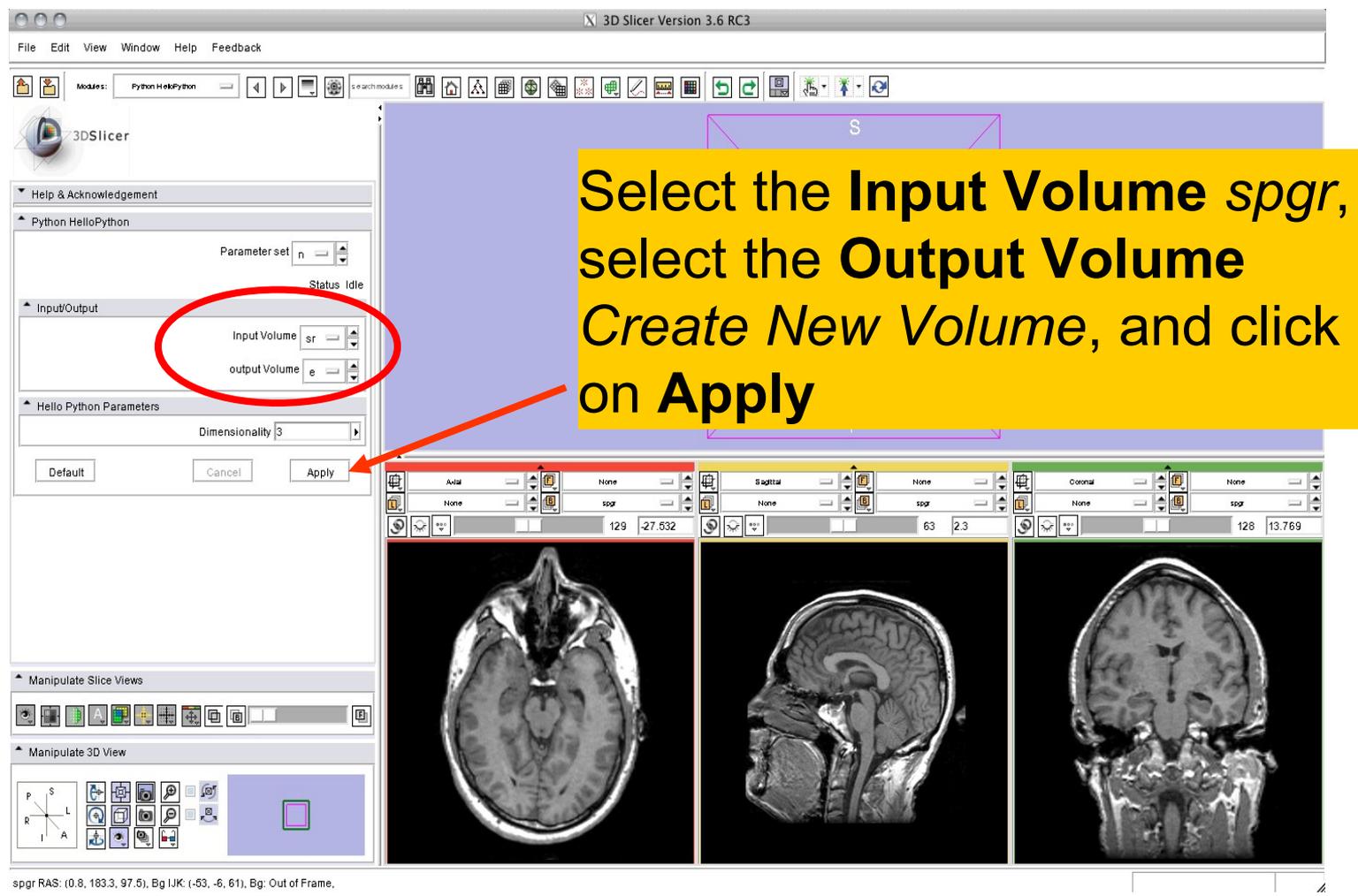


Running the Laplace Operator



The screenshot shows the 3D Slicer 3.6 RC3 interface. The 'Modules' menu is open, and 'Python Hello Python' is highlighted with a red circle. A yellow text box on the right contains the following text: **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 options. The main view displays three orthogonal MRI slices: Axial, Sagittal, and Coronal.

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

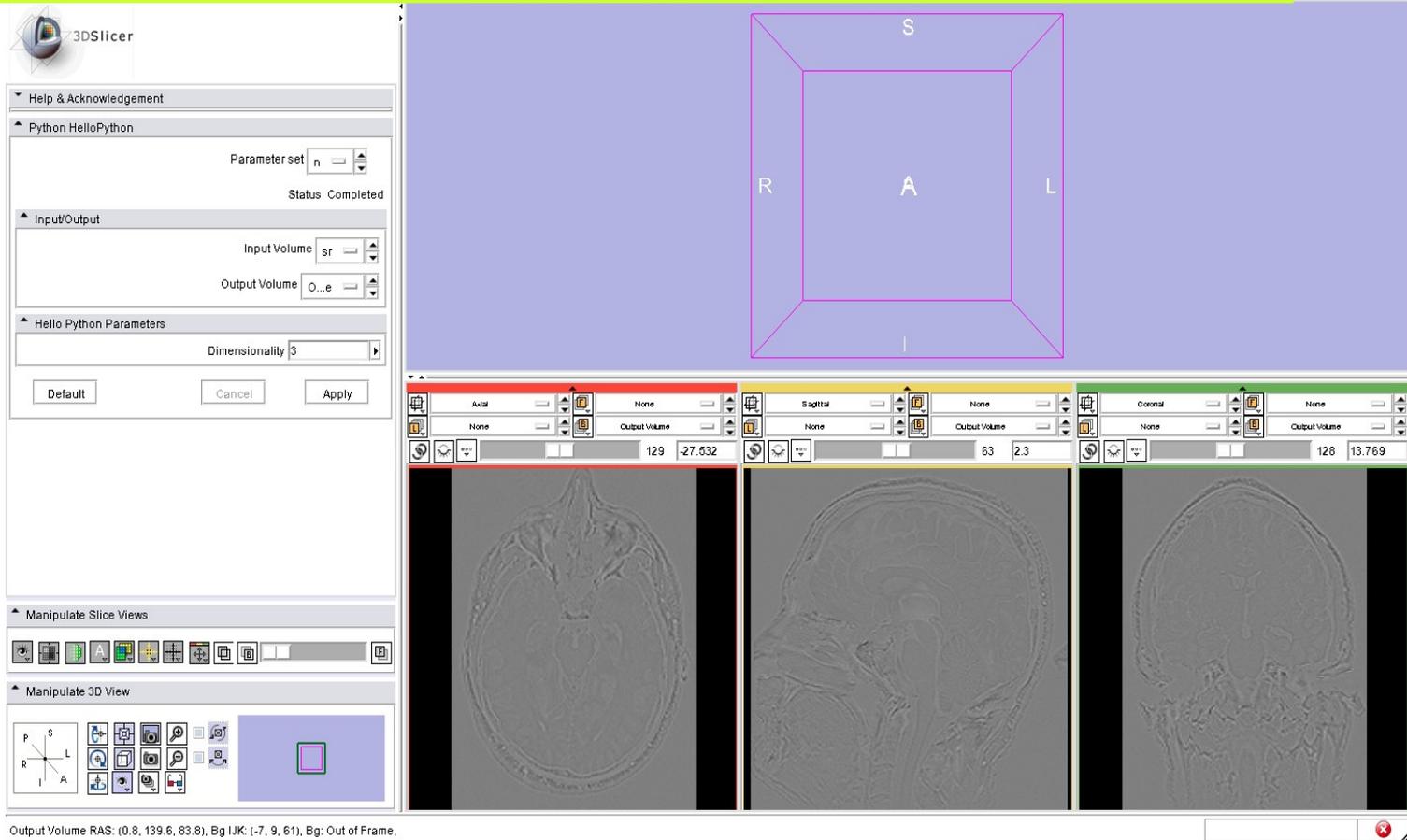
Manipulate 3D View

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

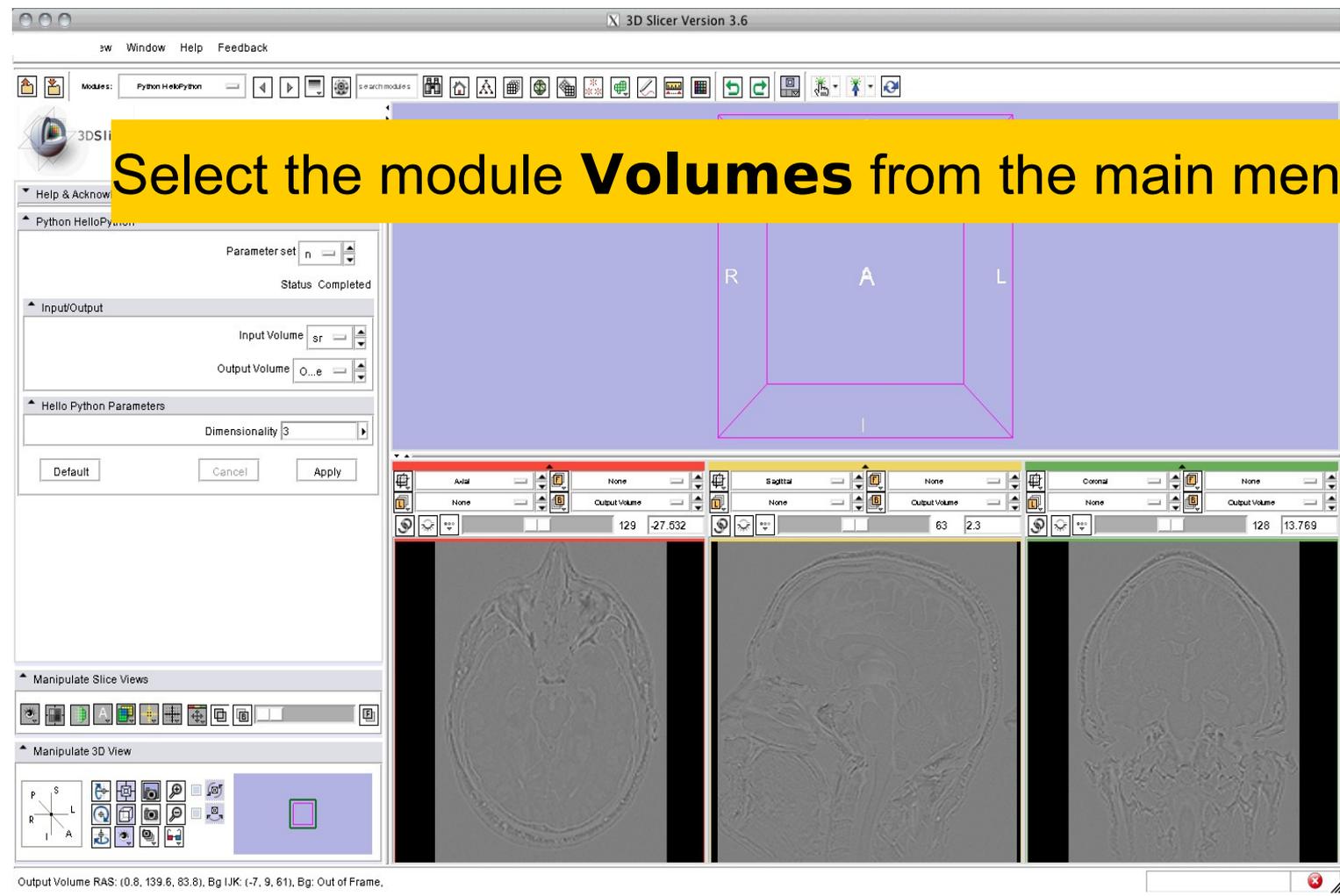
Select the Input Volume *spgr*, select the Output Volume *Create New Volume*, and click on Apply

Running the Laplace Operator

Slicer displays the Laplacian of the spgr image



Laplacian of the image



Laplacian of the image

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

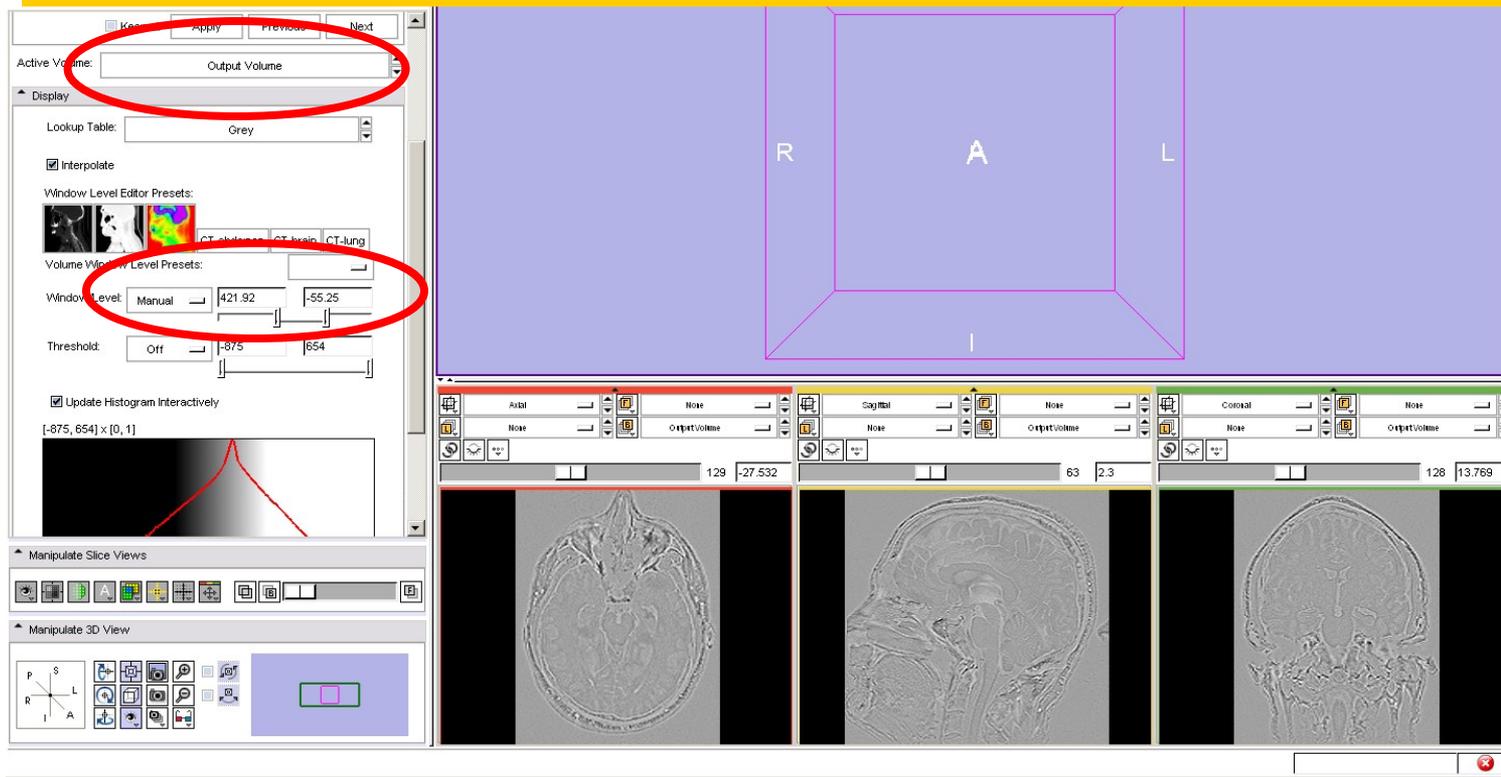
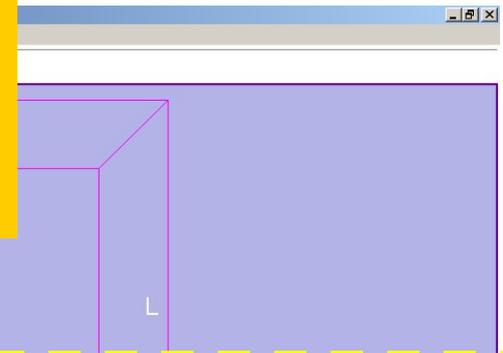


Image Sharpening

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)
```

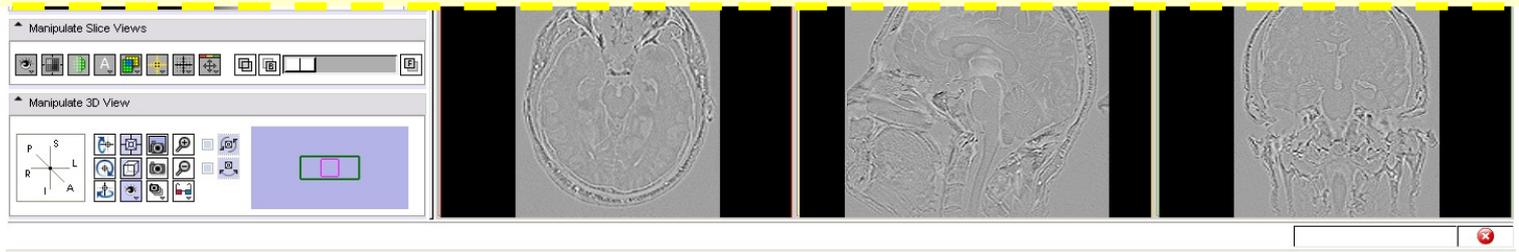
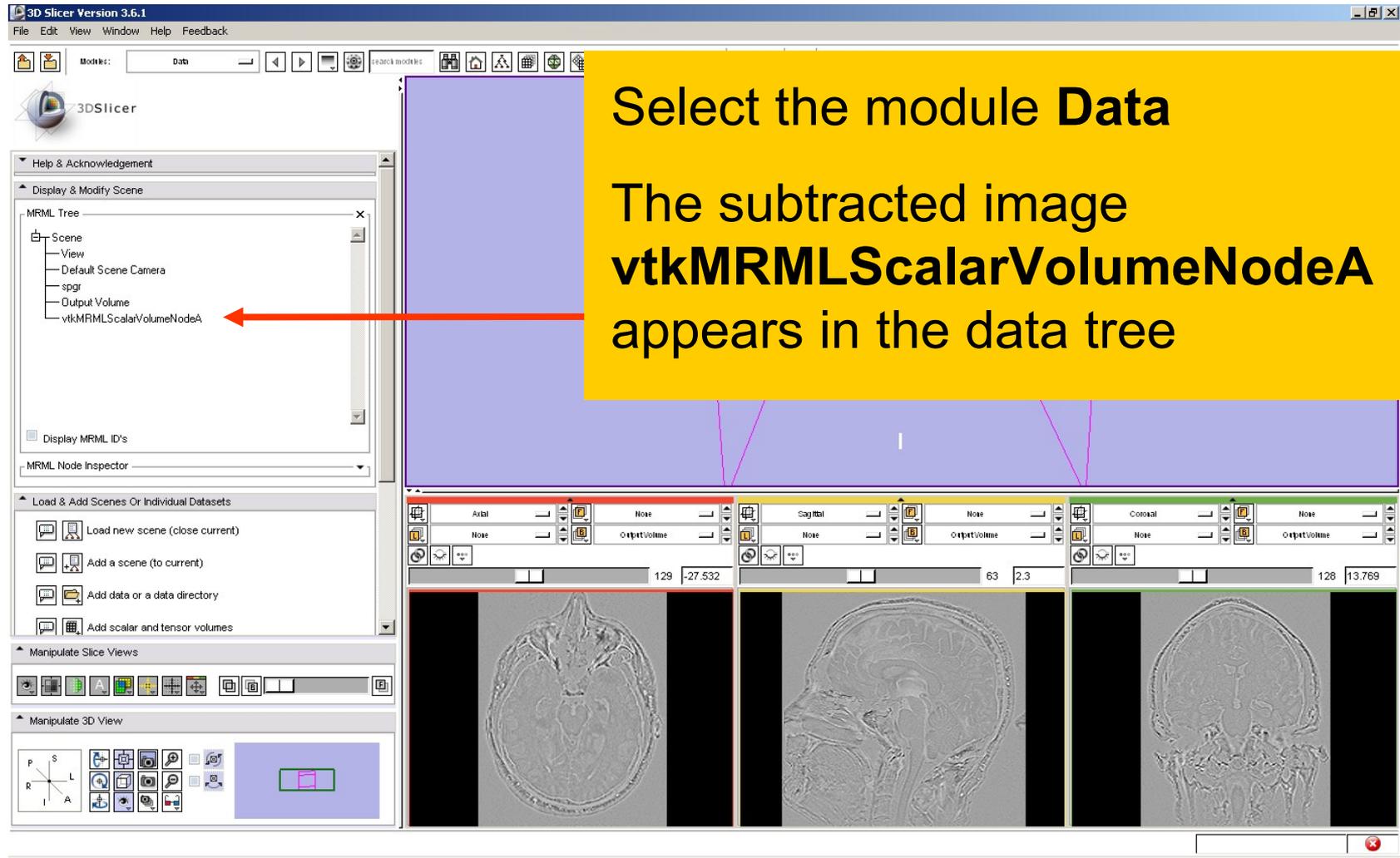


Image Sharpening

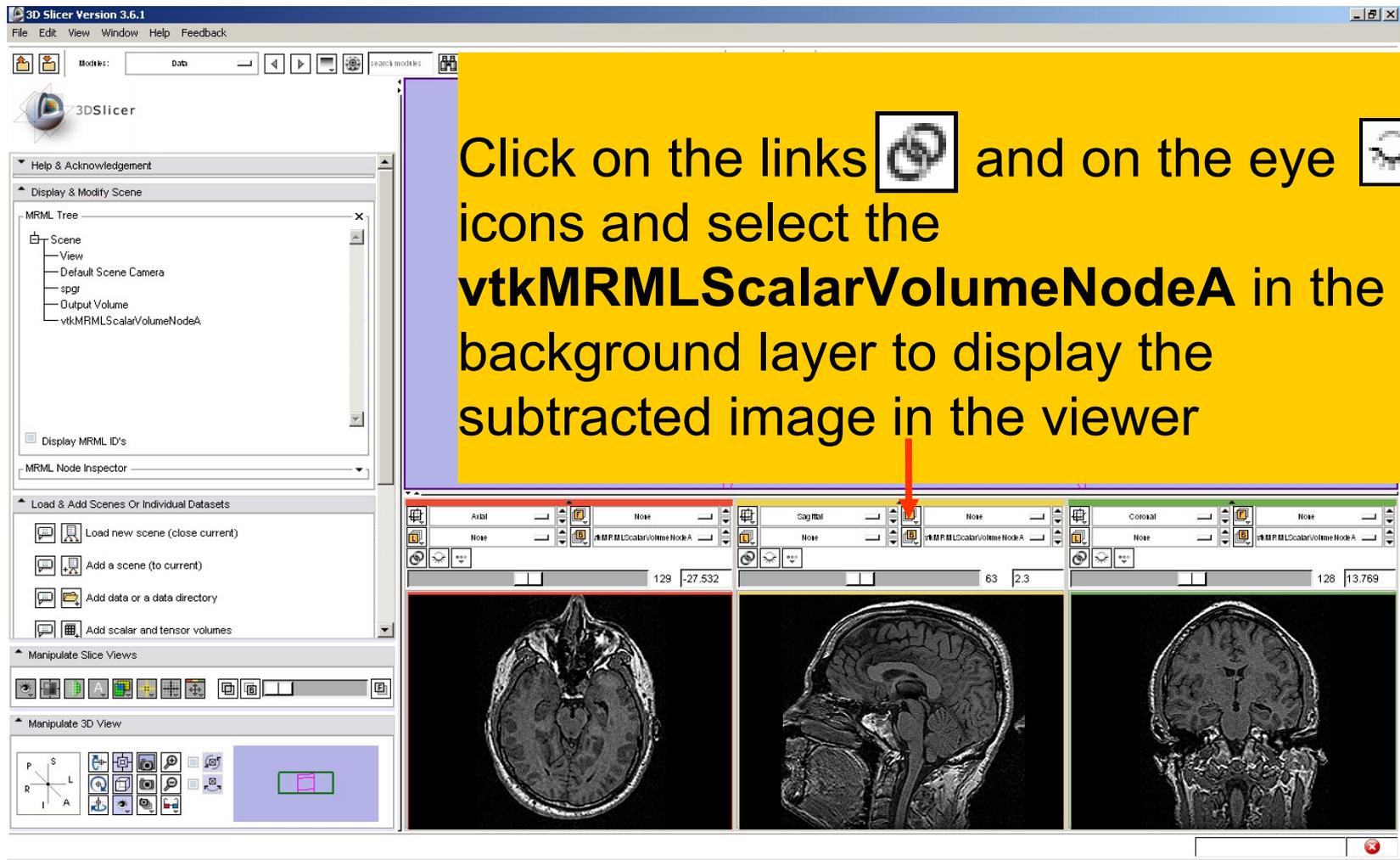


The screenshot shows the 3D Slicer 3.6.1 interface. The top menu bar includes File, Edit, View, Window, Help, and Feedback. Below the menu is a toolbar with various icons. The left sidebar contains several panels: 'Help & Acknowledgement', 'Display & Modify Scene', 'Load & Add Scenes Or Individual Datasets', 'Manipulate Slice Views', and 'Manipulate 3D View'. The 'Display & Modify Scene' panel shows the MRML Tree, which is expanded to show the 'Scene' node. Under 'Scene', there are sub-nodes for 'View', 'Default Scene Camera', 'spgr', 'Output Volume', and 'vtkMRMLScalarVolumeNodeA'. A red arrow points to 'vtkMRMLScalarVolumeNodeA'. 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 'Axial', 'Sagittal', and 'Coronal' views. The 'Manipulate 3D View' panel has a 3D view icon and a small 3D view window. The main window shows three slice views: Axial, Sagittal, and Coronal. Each slice view has a toolbar with 'None' and 'Output Volume' buttons. The Axial view shows a brain slice with a value of 129 and a range of -27.532. The Sagittal view shows a brain slice with a value of 63 and a range of 2.3. The Coronal view shows a brain slice with a value of 126 and a range of 13.769.

Select the module **Data**

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

Image Sharpening



3D Slicer Version 3.6.1

File Edit View Window Help Feedback

3DSlicer

Help & Acknowledgement

Display & Modify Scene

MRML Tree

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

Display MRML ID's

MRML Node Inspector

Load & Add Scenes Or Individual Datasets

- Load new scene (close current)
- Add a scene (to current)
- Add data or a data directory
- Add scalar and tensor volumes

Manipulate Slice Views

Manipulate 3D View

Click on the links  and on the eye  icons and select the **vtkMRMLScalarVolumeNodeA** in the background layer to display the subtracted image in the viewer

Abial None Sagittal None Coronal None

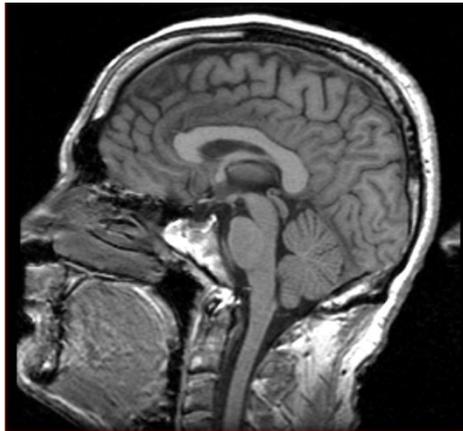
None vtkMRMLScalarVolumeNodeA None vtkMRMLScalarVolumeNodeA None

129 -27.532 63 2.3 128 13.769

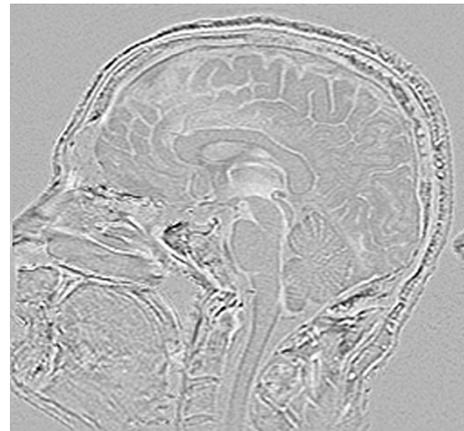
3D View: Axial, Sagittal, Coronal slices of a brain MRI scan.

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