

Anatomist: a python framework for interactive 3D visualization of neuroimaging data

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<http://brainvisa.info/doc/pyanatomist/sphinx/>





Anatomist : overview

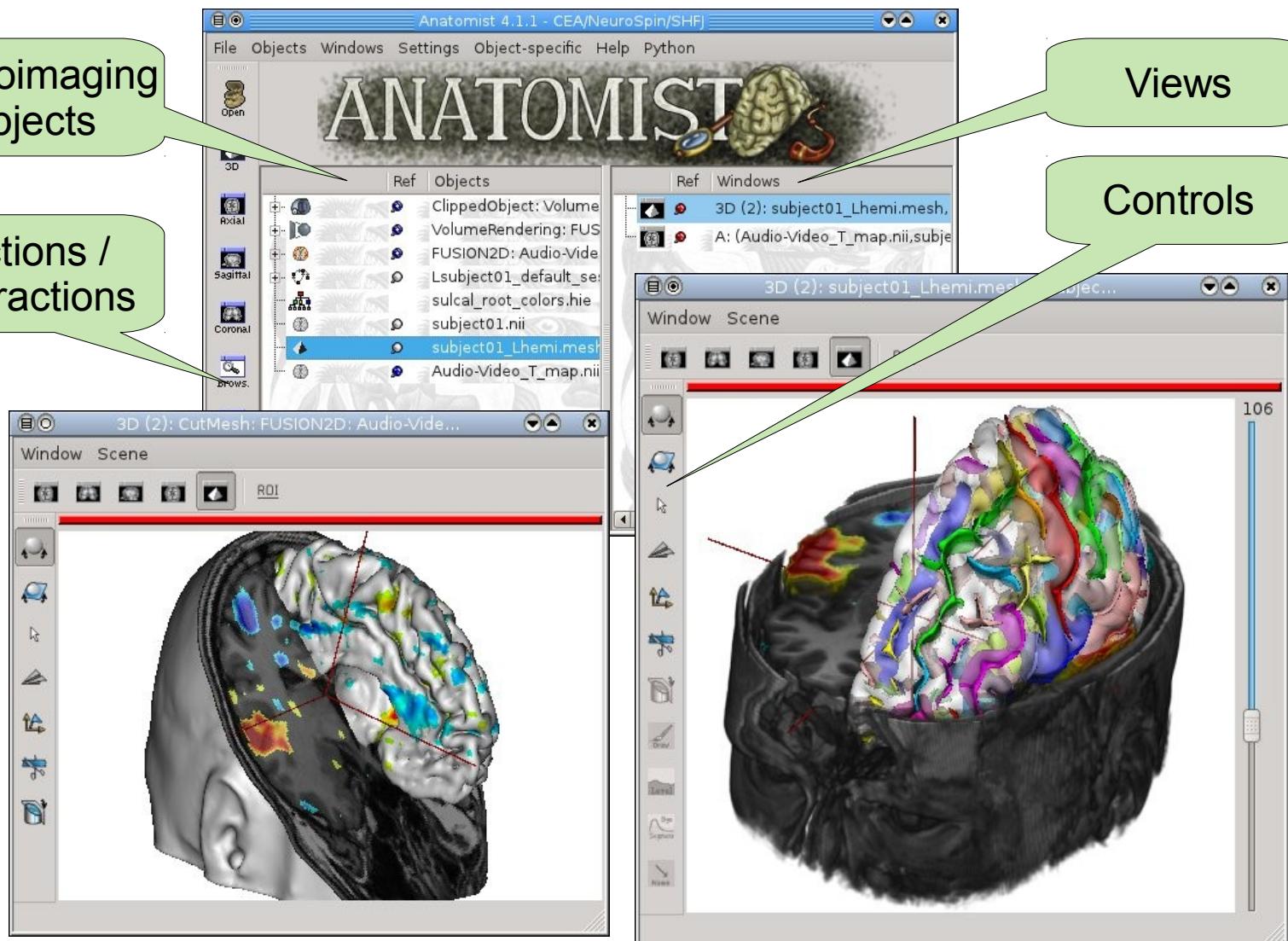
- Free software:
- Anatomist (C++ libraries) is BSD-like
 - PyAnatomist is GPL-like (due to PyQt)

Neuroimaging objects

Actions / interactions

Views

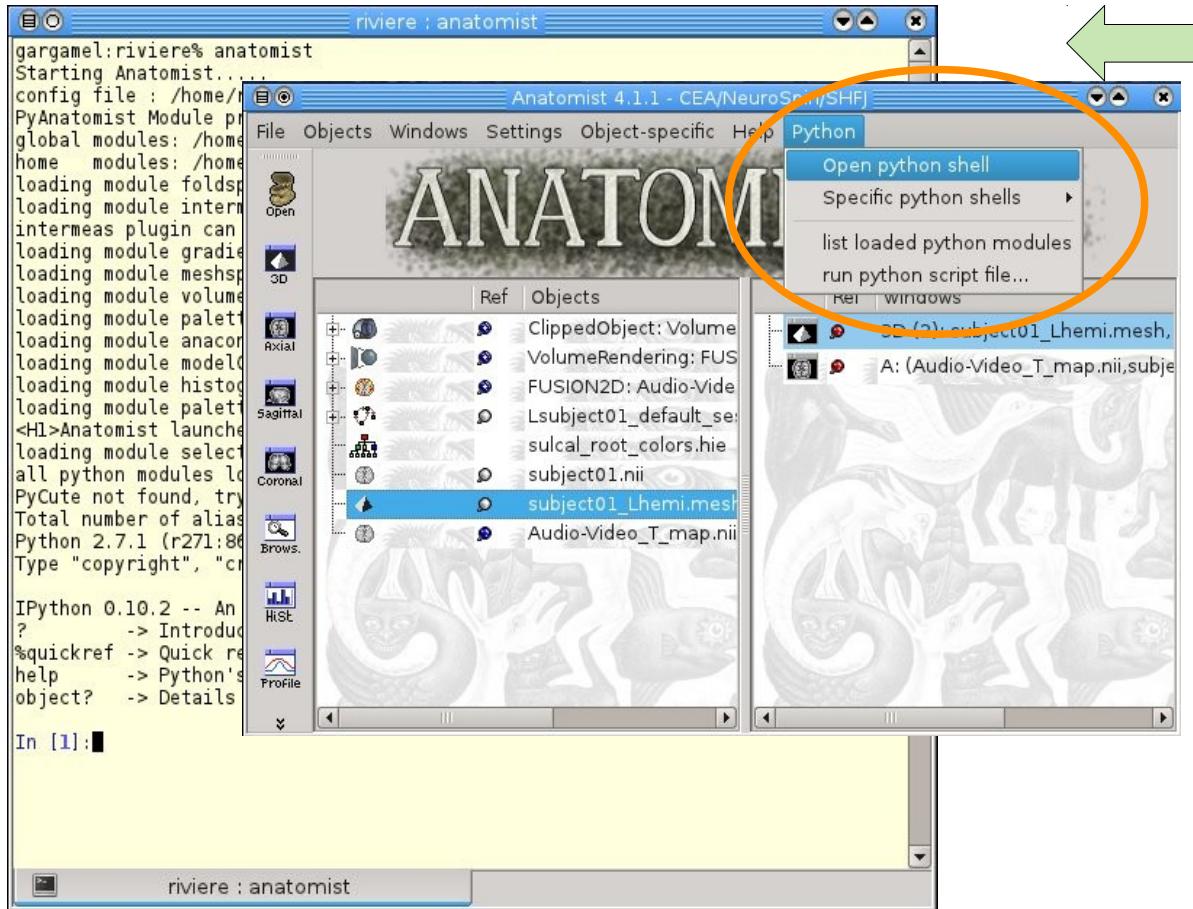
Controls





Python scripting

Several ways to enter python scripting mode



- From Anatomist: python menu
- From IPython (use -q4thread option, or --gui=qt with IPython >= 0,11):

```
% ipython -q4thread  
=> import anatomist.api as anatomist_api  
=> a = anatomist_api.Anatomist()
```

- Running a script with a Qt event loop

```
# !/usr/bin/env python  
  
import anatomist.api as anatomist_api  
import sys  
from PyQt4 import QtGui  
  
qapp = QtGui.QApplication( sys.argv )  
a = anatomist_api.Anatomist()  
QtGui.qApp.exec_()
```

- In a Python plugin for Anatomist

Location : \$HOME/.anatomist/python_plugins
In BrainVisa distributions :
<BV_dir>/share/anatomist-<version>/python_plugins

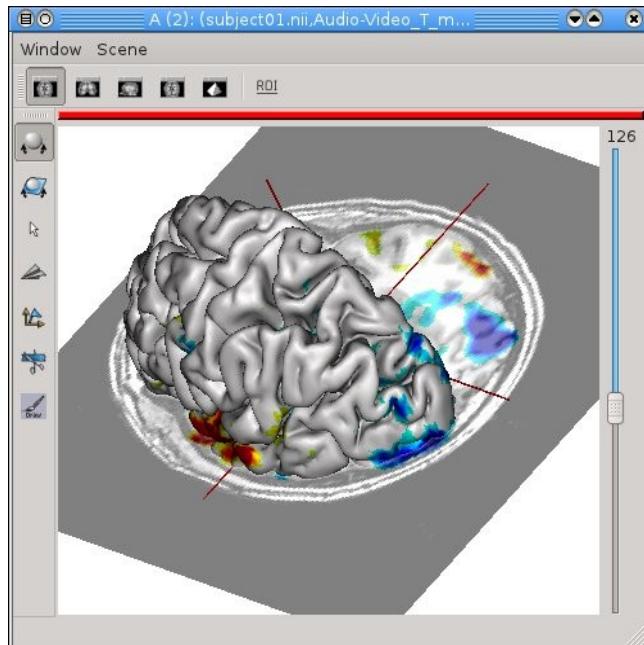
The internal commands system



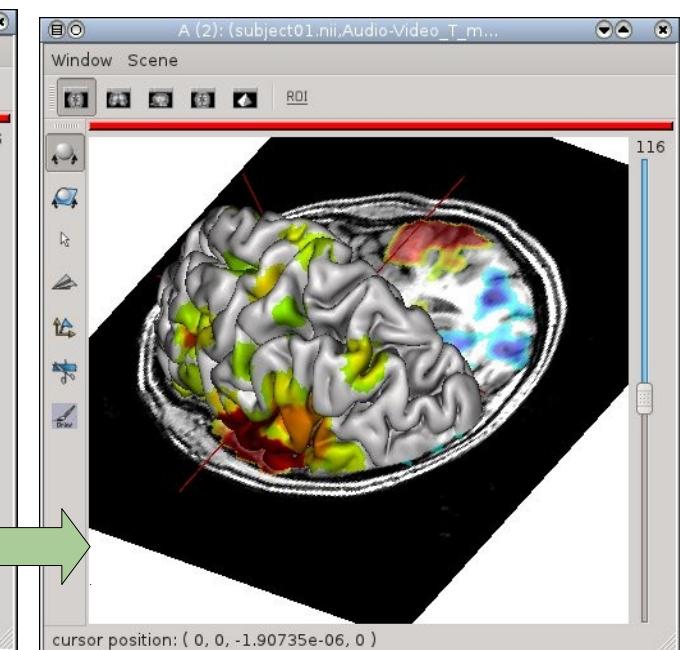
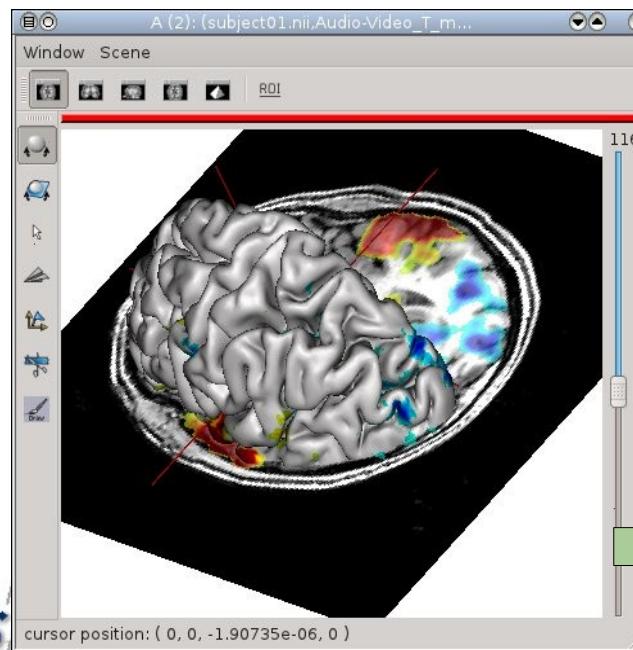
Internal commands system (this interpreter is older than the raise of Python for scientific applications),

<http://brainvisa.info/doc/anatomist/html/fr/programmation/commands.html>

Most have been ported to the newer Python API



```
>>> a.execute( 'LinkedCursor', window=win,
               position=( 0, 0, 0 ) )
>>> a.execute( 'Fusion2DParams', object=fus2d,
               mode='linear_on_defined', rate=0.5,
               reorder_objects=( vol, vol2 ) )
>>> a.execute( 'Fusion3DParams', object=fus3d,
               method='sphere', submethod='max',
               depth=8., step=0.5 )
>>> a.execute( 'Server', port=40007 )
```





Python API

A single base API for different control modes:

- Socket: control through a network connection
- Direct: access to the C++ library
- Threaded: thread-safe + direct

```
>>> import anatomist
>>> anatomist.setDefaultImplementation(anatomist.SOCKET)
>>> import anatomist.api as anatomist_api
>>> a = anatomist_api.Anatomist()
```

Or:

```
>>> import anatomist.socket.api as ana
>>> a = ana.Anatomist()
```

- The direct mode allows additional features (direct memory access)
- The socket mode allows several clients to connect to the same Anatomist, or a client program may pilot several Anatomist instances (on sevral machines)

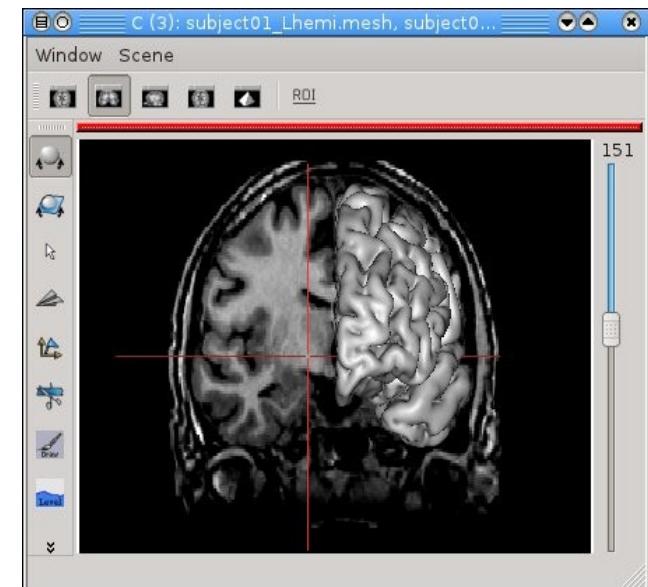
```
>>> # here we will control 2 anatomist server applications
>>> import anatomist.socket.api as anatomist_api
>>> a1 = anatomist_api.Anatomist() # default is host=localhost, port=40007
>>> a2 = anatomist_api.Anatomist( host='localhost', port=40008, forceNewInstance=True )
>>> w1 = a1.createWindow( '3D' )
>>> w2 = a2.createWindow( 'Browser' )
```



Basic operations

Loading objects, opening views

```
>>> a = anatomist_api.Anatomist()
>>> t1 = a.loadObject( 'volume.nii' )
>>> mesh = a.loadObject( 'mesh.gii' )
>>> win = a.createWindow( 'Coronal' )
>>> win.addObjects( ( t1, mesh ) )
```

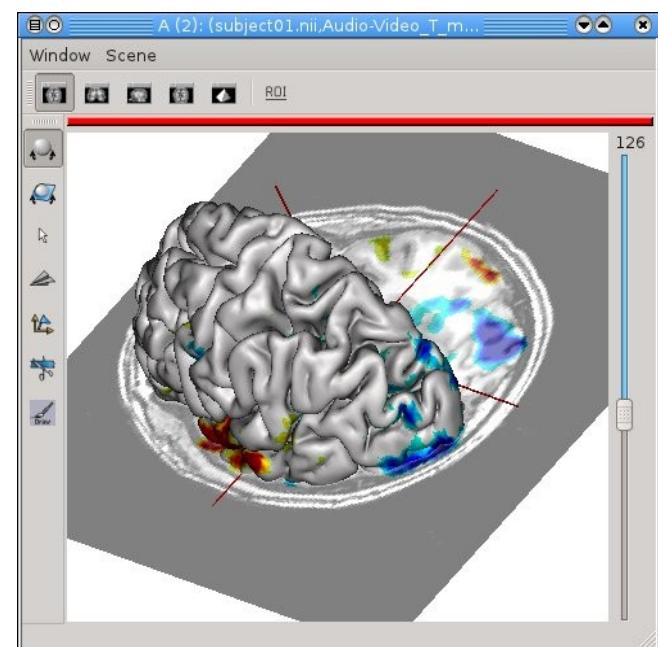


Using colors

```
>>> mesh.setMaterial( diffuse=[ 0.5, 0.5, 1., 0.8 ] )
>>> func = a.loadObject( 'Audio-Video_T_map.nii' )
>>> func.setPalette( 'tvalues100-200-100', MinVal=-4.13,
    maxVal=4.13, absoluteMode=True )
```

The “fusion” system: making new objects

```
>>> # objects may live in different coordinates systems
>>> a.applyBuiltInReferential( ( t1, func, mesh ) )
>>> fus2d = a.fusionObjects( ( t1, func ),
    method='Fusion2DMethod' )
>>> win2 = a.createWindow( 'Axial' )
>>> win2.addObjects( fus2d )
>>> # now another fusion type
>>> fus3d = a.fusionObjects( ( func, mesh ),
    method='Fusion3DMethod' )
>>> win2.addObjects( fus3d )
```



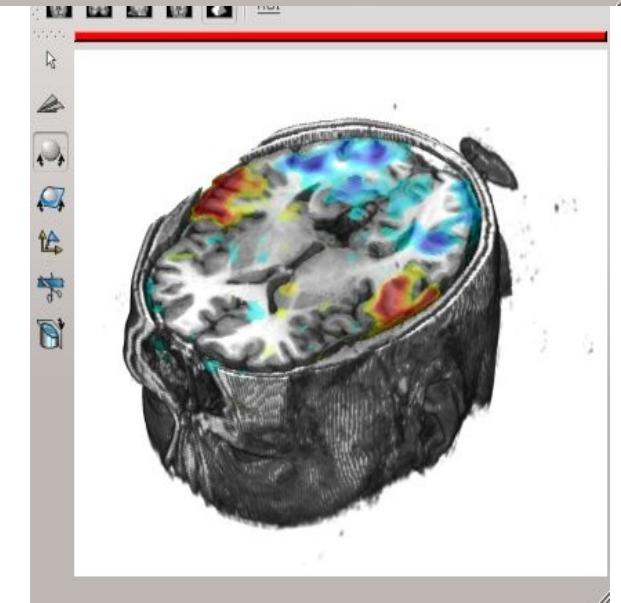
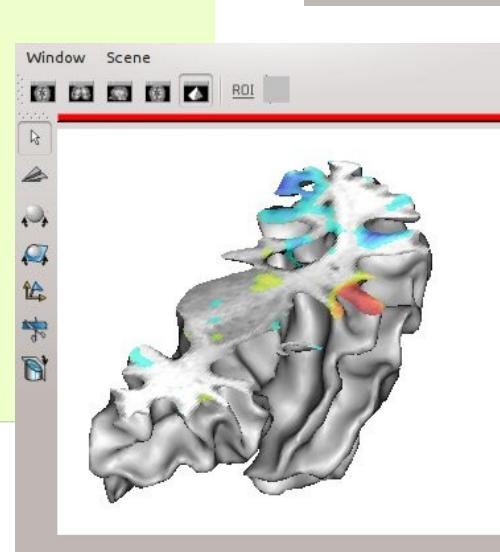
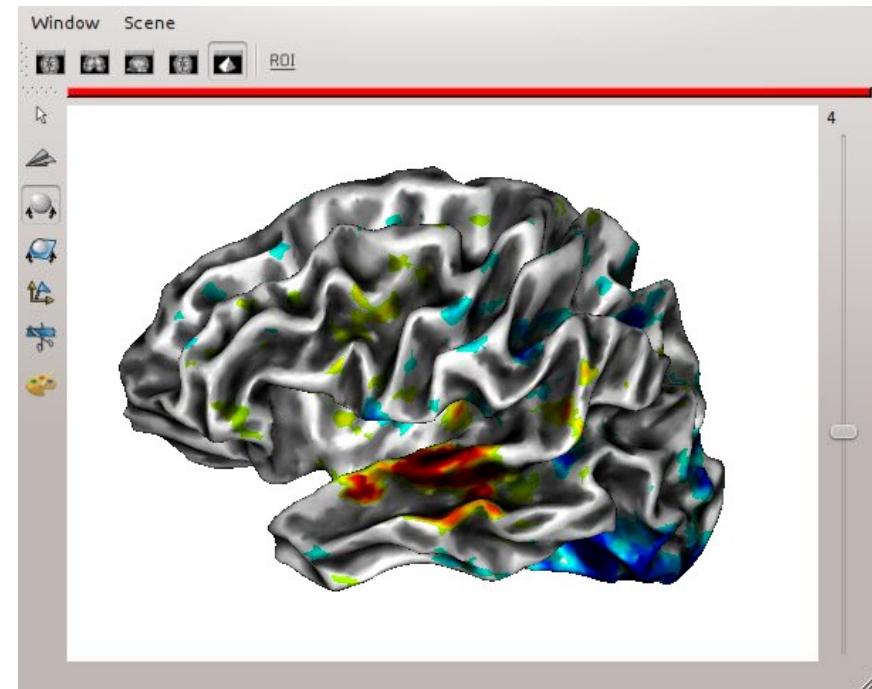
Generic “Fusion” mechanism



```
import anatomist.api as anatomist_api

a = anatomist_api.Anatomist()
t1 = a.loadObject( 'subject01.nii' )
white = a.loadObject( 'subject01_Lwhite.mesh' )
func = a.loadObject( 'Audio-Video_T_map.nii' )
inflat = a.loadObject( 'subject01_Lwhite_inflated_4d.mesh' )
curv = a.loadObject( 'subject01_Lwhite_curv.tex' )
a.applyBuiltInReferential( ( white, func ) )
func.setPalette( 'tvalues100-200-100', minVal=-4.13,
    maxVal=-4.13, absoluteMode=True )
curv.setPalette( 'B-W LINEAR', minVal=-0.691, maxVal=0.212,
    absoluteMode=True )
f3 = a.fusionObjects( ( white, func ),
    method='Fusion3DMethod' )
mtex = a.fusionObjects( ( f3, curv ),
    method='FusionMultiTextureMethod' )
tinfl = a.fusionObjects( ( inflat, mtex ),
    method='FusionTexSurfMethod' )
w = a.createWindow( '3D' )
w.addObjects( tinfl )

f2 = a.fusionObjects( ( t1, func ),
    method='Fusion2DMethod' )
cut = a.fusionObjects( ( white, f2 ),
    method='FusionCutMeshMethod' )
volr = a.fusionObjects( ( f2, ),
    method='VolumeRenderingFusionMethod' )
clip = fusionObjects( ( volr, ),
    method='FusionClipMethod' )
w2 = a.createWindow( '3D' )
w2.addObjects( cut )
w3 = a.createWindow( '3D' )
w3.addObjects( clip )
```



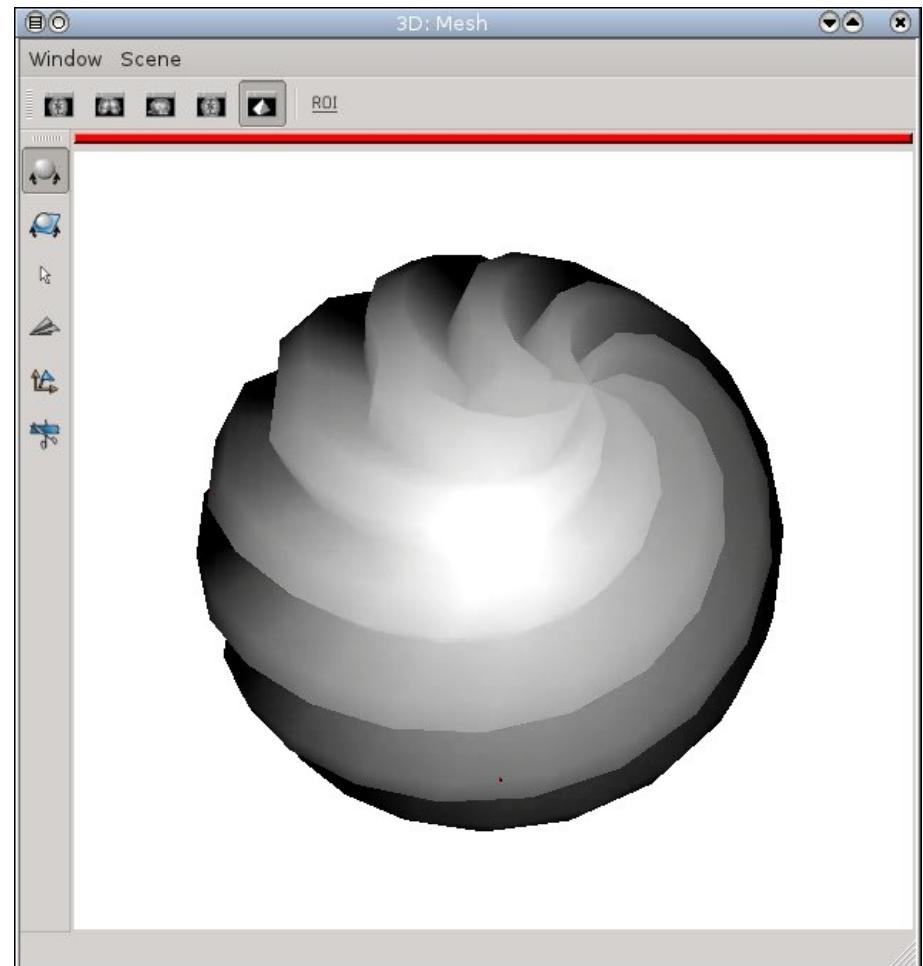
Direct mode features



```
from soma import aims
import time
import os
import anatomist.direct.api as anatomist_api
import sys
from PyQt4 import QtGui

m = aims.SurfaceGenerator.sphere( (0,0,0), 100, 500 )
a = anatomist_api.Anatomist()
# Put the mesh in anatomist
am = a.toAObject( m )
aw = a.createWindow( '3D' )
aw.addObjects( am )
coords = [ aims.Point3df(p) for p in m.vertex() ]
points = xrange( 0, len(coords), 3 )

for i in xrange( 10 ):
    # shrink
    for s in reversed(xrange(100)):
        for p in points:
            m.vertex()[p] = coords[p] * s/100.
        am.setChanged()
        am.notifyObservers()
        QtGui.qApp.processEvents()
        time.sleep( 0.01 )
    # expand
    for s in xrange(100):
        for p in points:
            m.vertex()[p] = coords[p] * s/100.
        am.setChanged()
        am.notifyObservers()
        QtGui.qApp.processEvents()
        time.sleep( 0.01 )
```



Anatomist and numpy

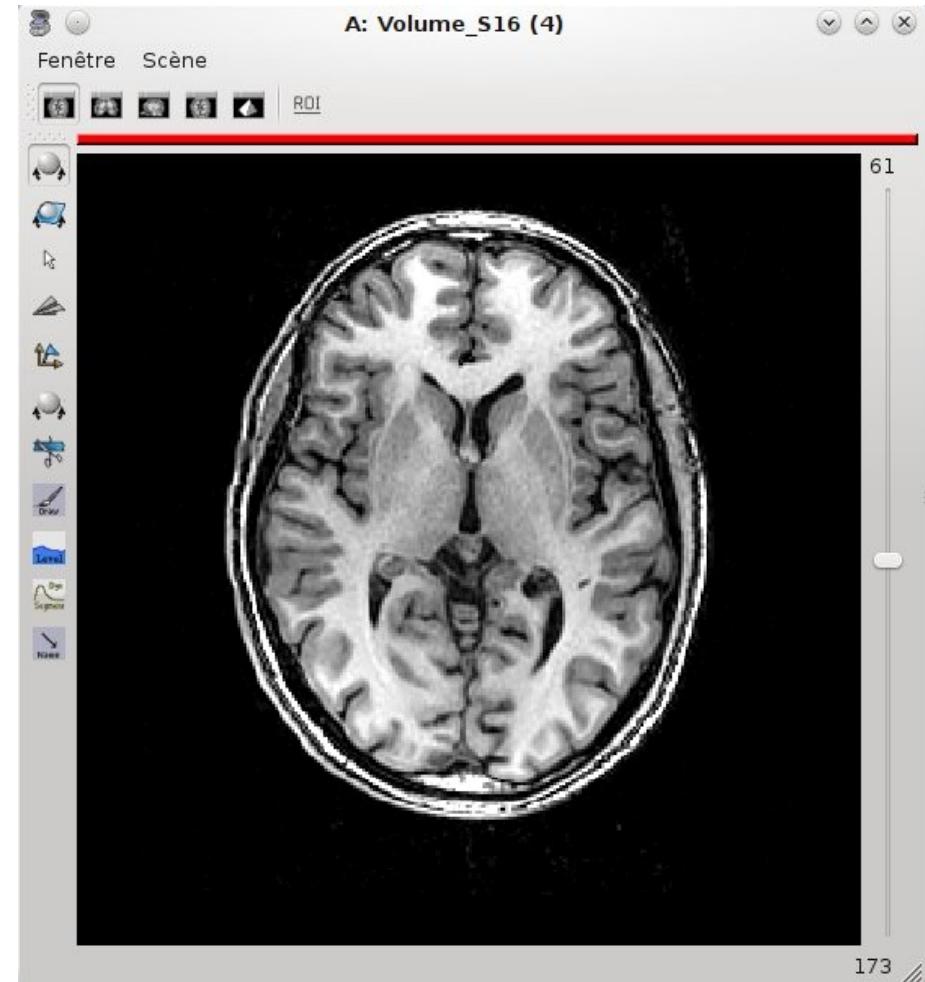


```
import anatomist.direct.api as anatomist_api
from soma import aims
import numpy
from PyQt4 import QtGui

a = anatomist_api.Anatomist()
qApp = QtGui.QApplication([])
vol = aims.read('subject01.nii')
bmask = aims.read('brain_subject01.nii')
masked = aims.Volume(vol)
avol = a.toA0Object(vol)
abmask = a.toA0Object(bmask)
amasked = a.toA0Object(masked)
w = a.createWindow('Axial')
w.addObjects(amasked)
qApp.processEvents()

# get numpy arrays on volume data
abm = numpy.array(bmask, copy=False)
am = numpy.array(masked, copy=False)
av = numpy.array(vol, copy=False)
# masking brain
am[abm==0] = 0
amasked.setChanged()
amasked.notifyObservers()
qApp.processEvents()

# iterative blurring
for i in range(50):
    am[1:-1, 1:-1, 1:-1, :] = (am[1:-1, 1:-1, 1:-1, :] * 3 \\
        + am[-2, 1:-1, 1:-1, :] + am[2:, 1:-1, 1:-1, :] \\
        + am[1:-1, -2, 1:-1, :] + am[1:-1, 2:, 1:-1, :] \\
        + am[1:-1, 1:-1, -2, :] + am[1:-1, 1:-1, 2:, :]) / 9
    amasked.setChanged()
    amasked.notifyObservers()
    qApp.processEvents()
```



Direct mode and GUI

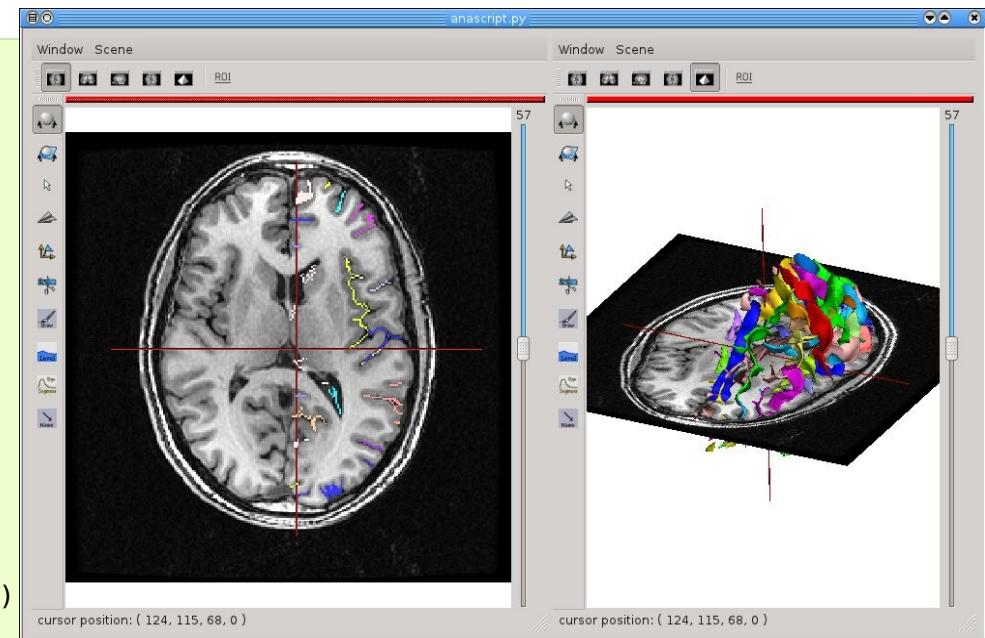


- Anatomist views internal representations are Qt widgets: using them in custom GUI is allowed

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

import anatomist.direct.api as anatomist_api
from PyQt4 import QtCore, QtGui
import sys

qapp = QtGui.QApplication( sys.argv )
# disable default anatomist main window
a = anatomist_api.Anatomist( '-b' )
# make a custom GUI with 2 views
mainw = QtGui.QMainWindow( None )
grid = QtGui.QWidget( mainw )
mainw.setCentralWidget( grid )
layout = QtGui.QGridLayout( grid )
w1 = a.createWindow( 'Axial' )
layout.addWidget( w1.getInternalRep(), 0, 0 )
w2 = a.createWindow( '3D' )
a.execute( 'LinkedCursor', window=w1, position=(124,115,68) )
layout.addWidget( w2.getInternalRep(), 0, 1 )
mainw.show()
# display something in the views
vol = a.loadObject( 'subject01.nii' )
nomenc = a.loadObject( 'sulci/sulcal_root_colors.hie' )
graph = a.loadObject( \
    'sulci/Lsubject01_default_session_auto.arg' )
a.execute( 'GraphDisplayProperties', objects=[graph],
    nomenclature_property='label' )
a.addObjects( ( vol, graph ), ( w1, w2 ), add_graph_nodes=True )
w2.camera( view_quaternion=[0.508661, 0.133626,
    0.192899, 0.828371], zoom=1.5 )
qapp.exec_()
del w1, w2, graph, vol, a
```



The `getInternalRep()` method of objects/windows grants access to a lower-level API which is the direct bindings to the C++ library API.

Custom GUI using Qt Designer



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

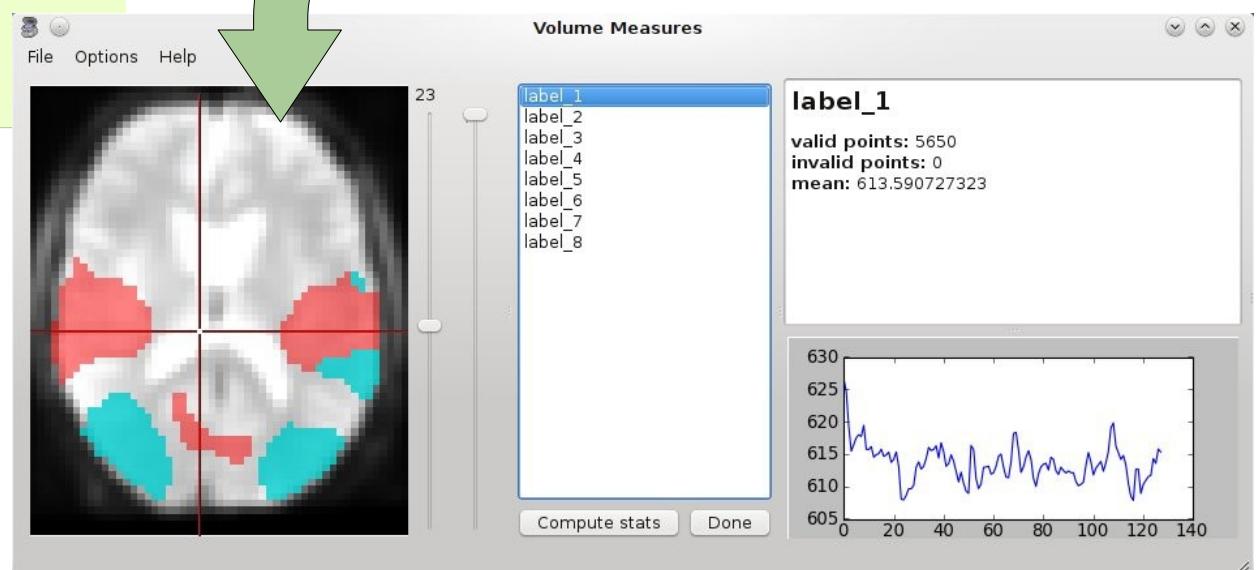
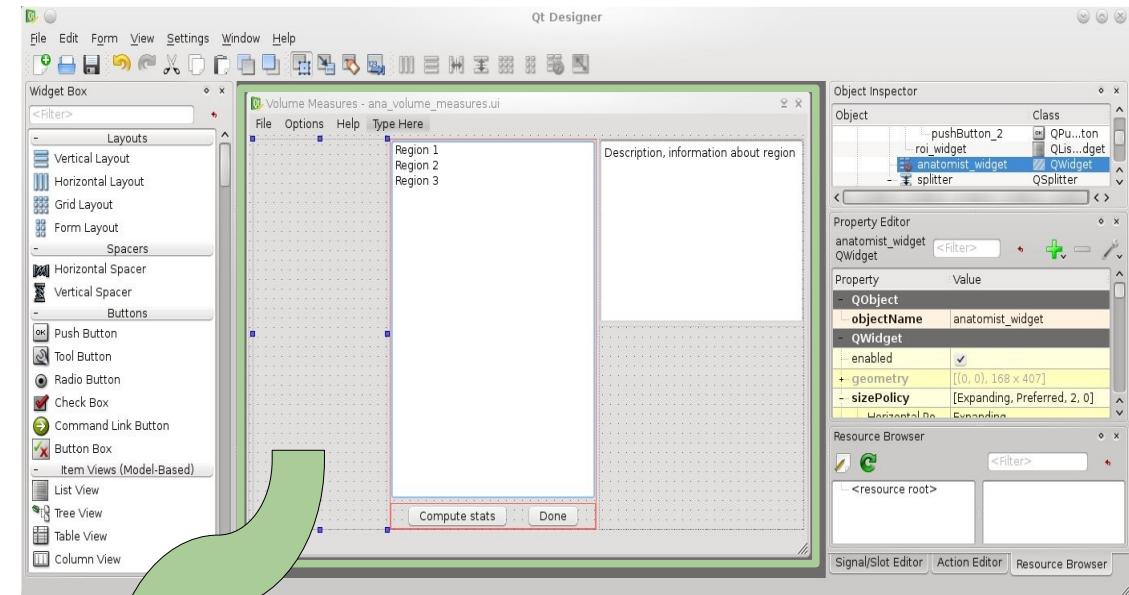
from PyQt4.uic import loadUi
from PyQt4.QtGui import QApplication, QVBoxLayout
import anatomist.direct.api as anatomist_api
Import sys

qapp = QApplication( sys.argv )
ui = 'ana_volume_measures.ui'
mainw = loadUi( ui )
anatomist = anatomist_api.Anatomist( '-b' )

# open an axial window
layout = QVBoxLayout( mainw.anatomist_widget )
ana_window = anatomist.createWindow( 'Axial',
    no_decoration=True )
ana_window.setParent( mainw.anatomist_widget )
layout.addWidget( ana_window.getInternalRep() )

# [...]

mainw.show()
```



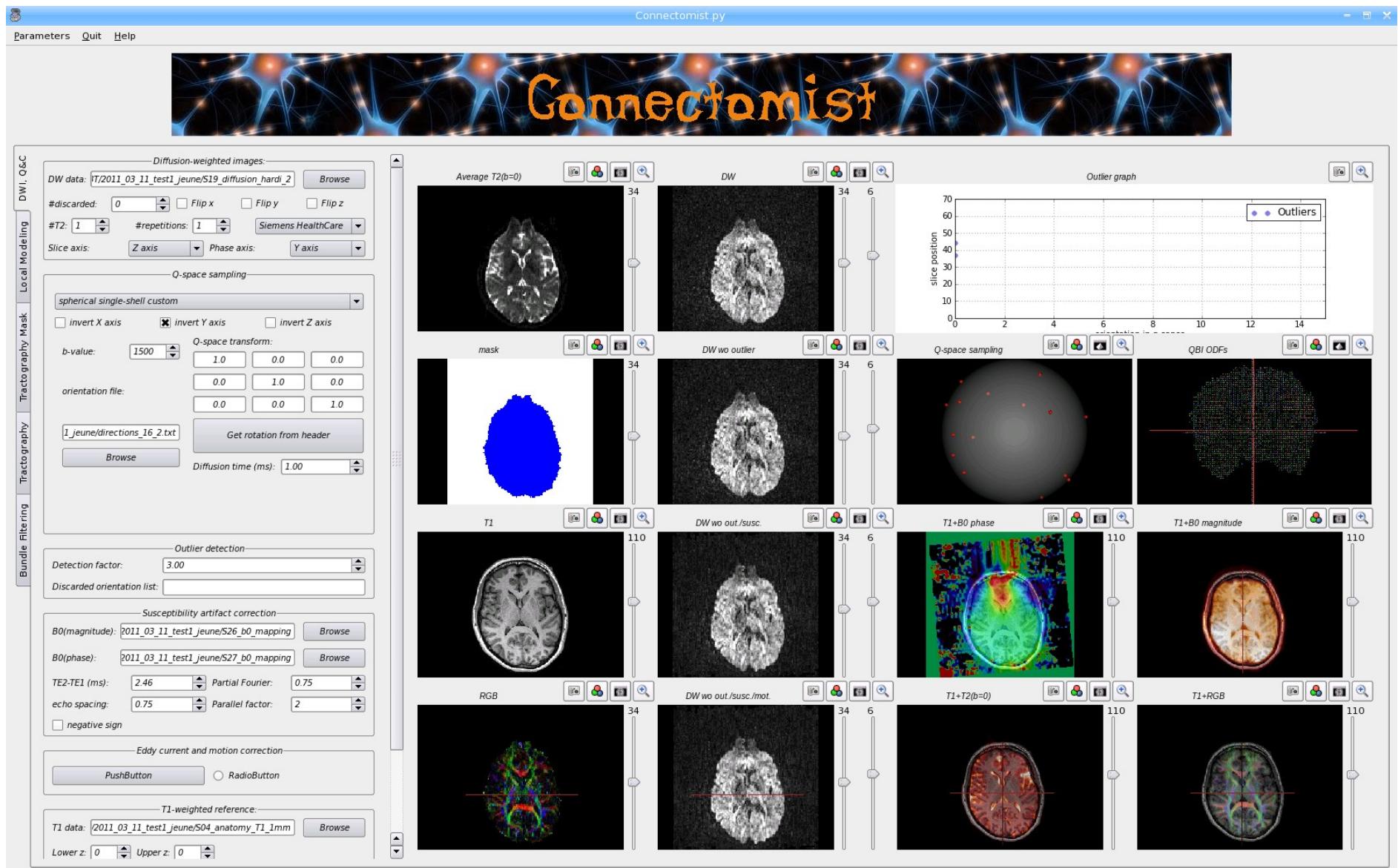
Conclusion



- Anatomist allows many pieces to be combined, python allows to do so quickly and easily.
- Simple programming interface for basic manipulations
- Everything is extensible (down to low-level, via C++ / python classes inheritance)
- Possibility to easily build custom dedicated applications



Conclusion



(image: courtesy of C. Poupon et al.)



Any questions ?

