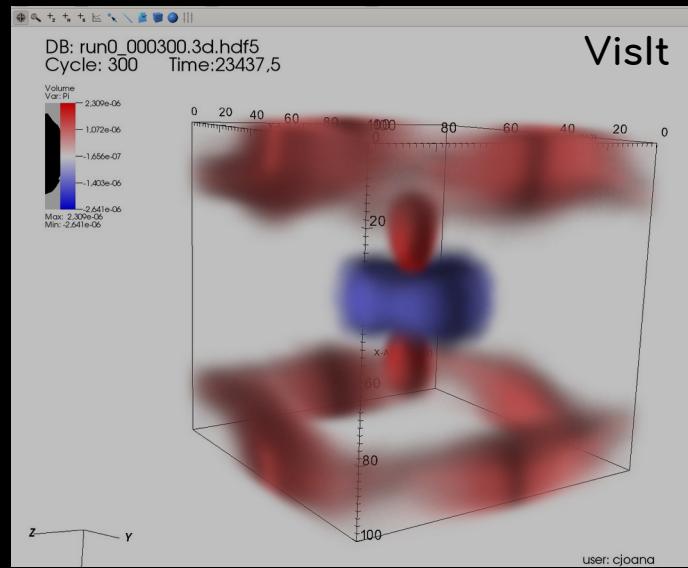


# Visualisation tools for GRChombo

```
import yt

dfn = './data/run0p_000300.3d.hdf5'
ds = yt.load(dfn)
L, _, _ = ds.domain_width

normal = 'z'
var = "K"
center = [L/2, L/2, L/2]
plot = yt.SlicePlot(ds, normal=normal,
                     fields=var, center=center)
plot.set_cmap(var, 'RdBu_r')
plot.save('./plots/{v}_slice.png'.format(v=var))
```



Cristian Joana - UCLouvain (CURL)

GRChombo Meeting 30/03/2022

# Visualisation tools for GRChombo

---

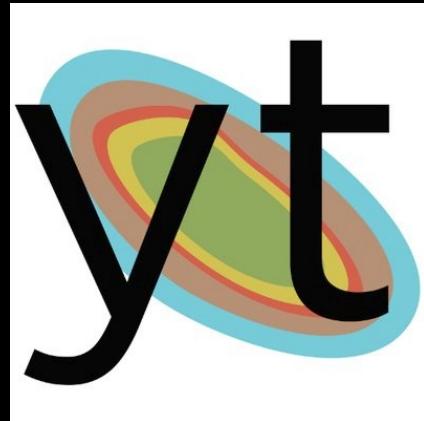
- Using YT (python)
  - Installation
  - Utilities
  - Examples
- Using Visit (GUI)

For movies → Josu slides 2019

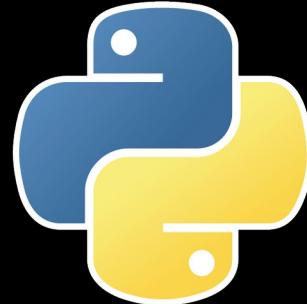


# YT - project

---



+



<https://yt-project.org/docs/dev/>

(documentation)

<https://yt-project.org/community.html> (mailing list, Slack, etc.)



# YT - Installation

---

- Via conda:

```
$ conda install -c conda-forge yt
```

- Via PiP:

```
$ pip install yt
```

- Via github repository:

```
$ git clone https://github.com/yt-project/yt  
$ cd yt && python setup.py install
```



# YT - Loading files

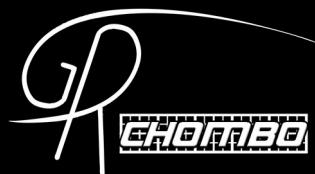
---

## Loading hdf5 file

```
# define dataset's path
dfn = './data/run0p_000300.3d.hdf5'

#load dataset
ds = yt.load(dfn)

# Equivalent to:
# ds = yt.frontends.chombo.ChomboDataset(dfn)
```



# YT - Loading files

---

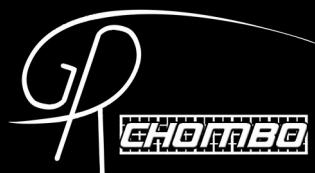
## Loading hdf5 file

```
# define dataset's path
dfn = './data/run0p_000300.3d.hdf5'

units_override = {"length_unit": (1.0, "l_pl"),
                  "time_unit": (1.0, "t_pl"),
                  "mass_unit": (1.0, "m_pl")}
unit_system = 'planck'

# load dataset
ds = yt.load(dfn,
              unit_system=unit_system, units_override=units_override)

# ds = yt.frontends.chombo.ChomboDataset(dfn,
#                                         unit_system=unit_system, units_override=units_override)
```



# YT - Handeling data

## Loading data variables

```
#load dataset  
ds = yt.load(dfn)
```

```
# Examples using the variable "K"  
  
reg = ds.r[:, :, :] # flat array  
print('shape reg:', reg['K'].shape )  
  
reg3d = ds.r[:, :, :, 120j, 120j, 120j] # or 3D when specified the resolution  
print('shape reg3D:', reg3d['K'].shape )  
  
L, _, _ = ds.domain_width  
slc = ds.r[:, :, 120j, L/2]  
print('shape slc:', slc['K'].shape )  
  
shape reg: (2097152,)  
shape reg3D: (120, 120, 120)  
shape slc: (120, 120)
```



# YT - Handeling data

Example: extraction of data & AMR coordinates

```
import matplotlib.pyplot as plt

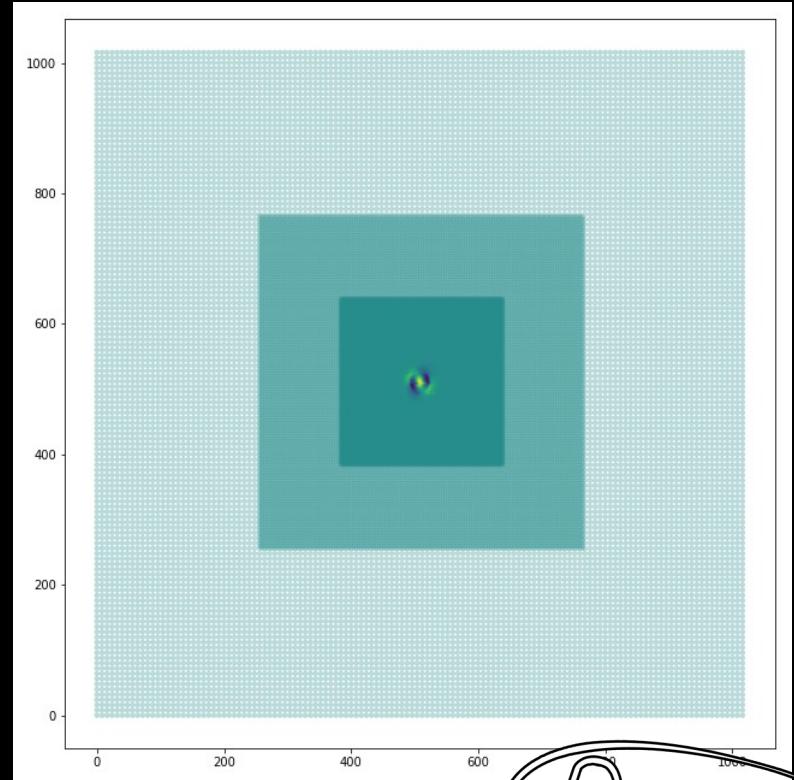
#Loading values of 'Avec0' and coordinates (taking into account AMR)
reg = ds2.all_data() # indexing data as flat array (contain all variables)
values = reg['Avec0'] # flat array that contains variable "Avec0"

# Loading grid-cell centered coordinates
X = reg['x']
Y = reg['y']
Z = reg['z']

# define position as bin-border of the grid
xpos = reg['x'] - reg['dx']/2
ypos = reg['y'] - reg['dy']/2
zpos = reg['z'] - reg['dz']/2

_, _, L = ds.domain_width
c_z = zpos[zpos >= L*0.05][0] #chosing grid-coord of interest
mask_cslice = np.array(zpos == c_z, dtype=bool) # mask for data selection

# plot
fig, ax = plt.subplots(figsize=(9,9))
ax.scatter(xpos[mask_cslice], ypos[mask_cslice], c=values[mask_cslice],
           s=20, edgecolor='r', alpha=0.3)
plt.tight_layout()
plt.savefig("./plots/plot_amr.png")
```



# YT - Handeling data



## Setting up derived variables

```
def _cell_volume(field, data): # 'field', 'data' arguments needed
    vol = data["chi"]**(-1.5) * data["dx"]**(3)
    return vol

ds.add_field(('chombo', 'cell_vol'), sampling_type="cell",
             units = "l_pl**3", function=_cell_volume)

reg = ds.r[:, :, :, :] # flat array
con_L = np.sum(reg['dx']**3)**(1/3)
eff_L = np.sum(reg['cell_vol'])**(1/3)

print("conformal / effective grid-size: {c:.2e} {e:.2e}".format(c=con_L, e=eff_L))
print("domain L", ds.domain_width[0])

conformal / effective grid-size: 1.00e+05 l_pl 2.79e+05 l_pl
domain L 100000.0 code_length
```

NB: `dx`, `dy`, ... & `x`, `y`... are automatically yt-generated grid variables.

# YT - Handeling data

---

Setting new fields : gradients

```
ds.add_gradient_fields([('chombo', 'K')) # uses second-order centered differences
```

```
[('chombo', 'K_gradient_x'),  
 ('chombo', 'K_gradient_y'),  
 ('chombo', 'K_gradient_z'),  
 ('chombo', 'K_gradient_magnitude')]
```

```
print(reg['K_gradient_x'])
```

```
[ 7.49475932e-12  7.48383738e-12  7.47916149e-12 ... -7.53537997e-12  
 -7.50883771e-12 -7.48724252e-12] 1/l_pl
```



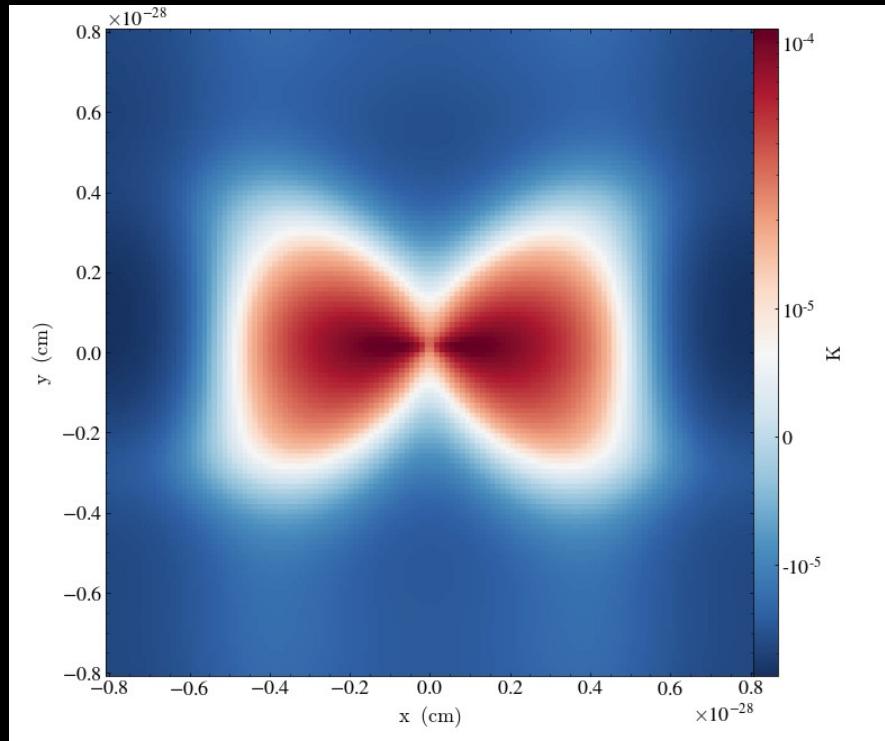
# YT – Plotting utilites

yt.SlicePlot()

```
dfn = './data/run0p_000300.3d.hdf5'
ds = yt.load(dfn)
L, _, _ = ds.domain_width

normal = 'z'
var = "K"
center = [L/2, L/2, L/2]
plot = yt.SlicePlot(ds, normal=normal,
                     fields=var, center=center)
plot.set_cmap(var, 'RdBu_r')
plot.save('./plots/{v}_slice.png'.format(v=var))

['./plots/K_slice.png']
```



→ <https://yt-project.org/doc/visualizing/plots.html>



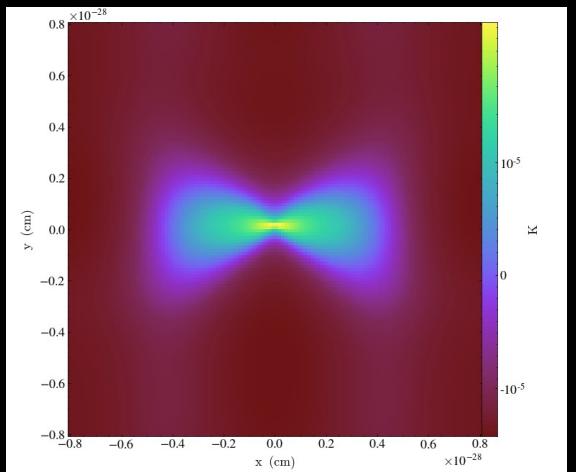
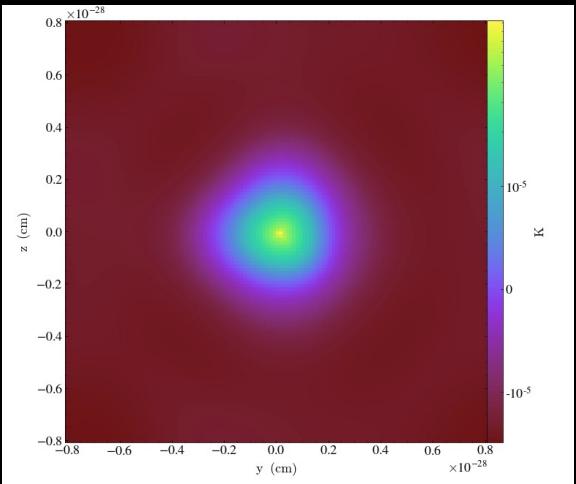
# YT – Plotting utilites



yt.ProjectionPlot()

```
dfn = './data/run0p_000300.3d.hdf5'
ds = yt.load(dfn)
ds.add_field([('chombo', 'cell_vol'),
              sampling_type="cell", units = "l_pl**3", function=_cell_volume)

var = "K"
plot = yt.ProjectionPlot(ds, fields=var, axis='x',
                         method='integrate', weight_field='cell_vol')
plot.save('./plots/{v}_projection_x.png'.format(v=var))
plot = yt.ProjectionPlot(ds, fields=var, axis='z',
                         method='integrate', weight_field='cell_vol')
plot.save('./plots/{v}_projection_z.png'.format(v=var))
```



→ <https://yt-project.org/doc/visualizing/plots.html>

# (non) YT – Plotting

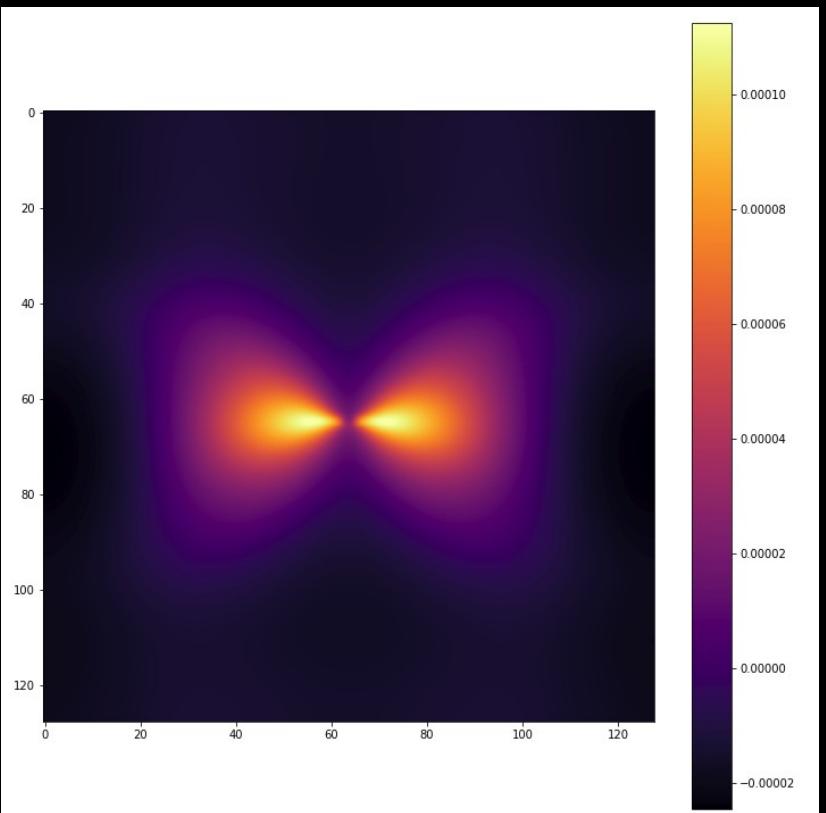
Via matplotlib

```
import matplotlib.cm as cm
import matplotlib.pyplot as plt

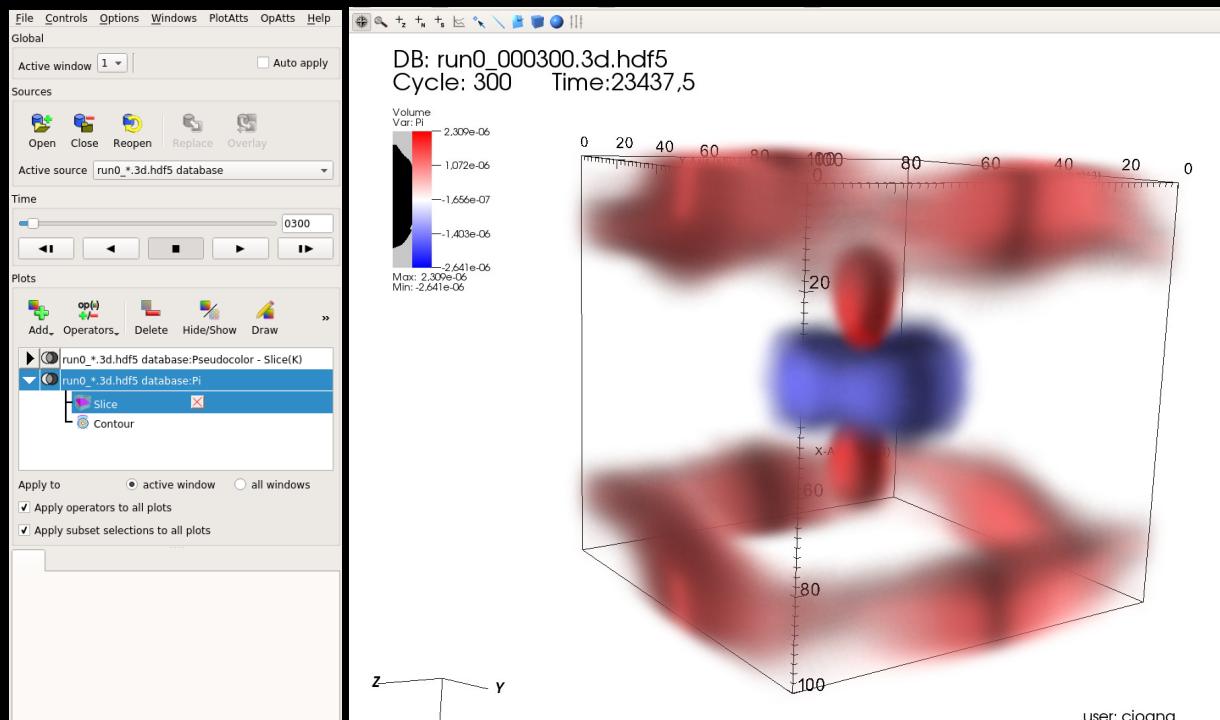
%matplotlib inline

dfn = './data/run0p_000300.3d.hdf5'
ds = yt.load(dfn)
L, _, _ = ds.domain_width
slc = ds.r[:,::128j,::128j, L/2]

fig, ax = plt.subplots(figsize=(10,10))
plot1 = ax.imshow( slc['K'], interpolation='spline16',
                   cmap=cm.inferno)
fig.colorbar(plot1, ax=ax)
plt.savefig('./plots/K_slice_matplotlib.png')
```



# Visit



Download & documentation:

<https://wci.llnl.gov/simulation/computer-codes/visit/downloads>

[https://visit-sphinx-github-user-manual.readthedocs.io/en/develop/gui\\_manual/](https://visit-sphinx-github-user-manual.readthedocs.io/en/develop/gui_manual/)



# Visit - Installation



You probably already have it installed. If not, **don't worry**, you will manage :)

+ info:

<https://github.com/GRChombo/GRChombo/wiki/Visualising-outputs>

[https://github.com/GRChombo/GRChombo/wiki/files/grchombo\\_ubuntu.pdf](https://github.com/GRChombo/GRChombo/wiki/files/grchombo_ubuntu.pdf)

shared by Leonard Wernecke



## 1.9 Installing VisIt

shared by Leonard Wernecke

VisIt is the software used by the developers of GRChombo to make beautiful plots and visual simulations. It is compatible with .hdf5 files, so it is a nice idea to install it.

Let us start by going to the following webpage

<https://wci.llnl.gov/simulation/computer-codes/visit/executables>

and downloading both the install script (copy the page to a file, in my case I have created the file `visitinstall.sh`) and the Ubuntu 14.04 executable. At the time of writing, version 2.13.0 was downloaded.

Then give permission so that the file can be executed

```
chmod 755 visitinstall.sh
```

and type

```
./visitinstall.sh 2.13.0 linux-x86_64-ubuntu14 /usr/local/visit
```

When prompted, choose the “No System Configuration” option. Then open your `~/.bashrc` file again and include at the bottom of the file the line

```
export PATH="/usr/local/visit/bin:$PATH"
```

Save the file and close it. Close all terminal windows and open a new one.

# Visit - Installation



In Ubuntu: (not tested)

Last version avail. : Vislt 3.1.1 (Feb 2020)

The screenshot shows a Stack Exchange post titled "Vislt 2.1.3 Ubuntu 18.10 setup without root". The post has 0 votes and no answers. It contains instructions for installing Visit without root privileges:

- go to the download site: <https://wci.llnl.gov/simulation/computer-codes/visit/executables>
- download the "Linux - x86\_64 64 bit" version
- extract:

```
tar xvf visit2_13_3.linux-x86_64-ubuntu18.tar.gz`
```
- run:

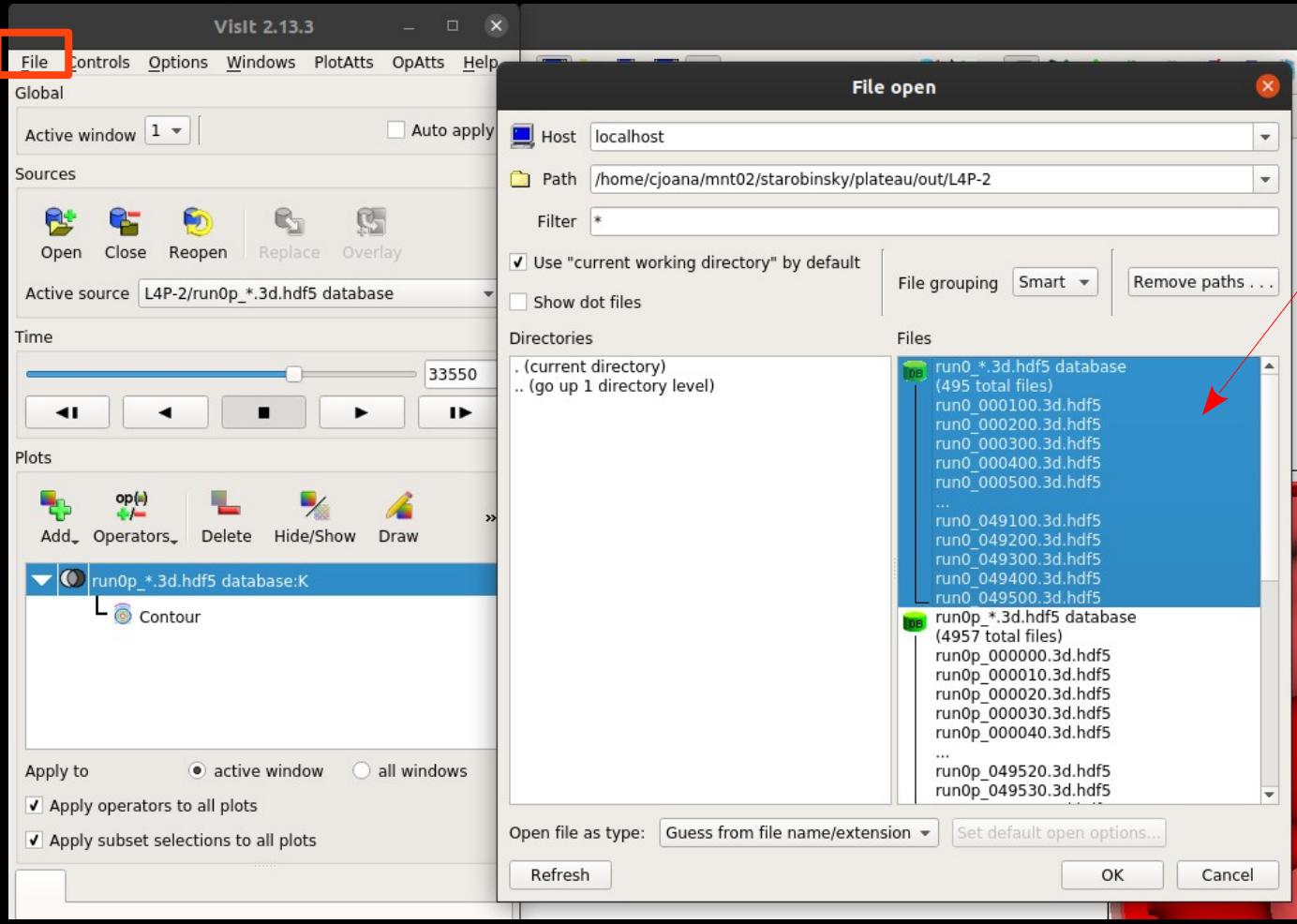
```
./visit2_13_3.linux-x86_64/bin/visit
```

The post also mentions it was used at <https://stats.stackexchange.com/questions/376361/how-to-find-the-sample-points-that-have-statistically-meaningful-large-outlier-r>.

share improve this answer

answered May 1 '19 at 9:31 by Ciro Santilli 新疆改造中

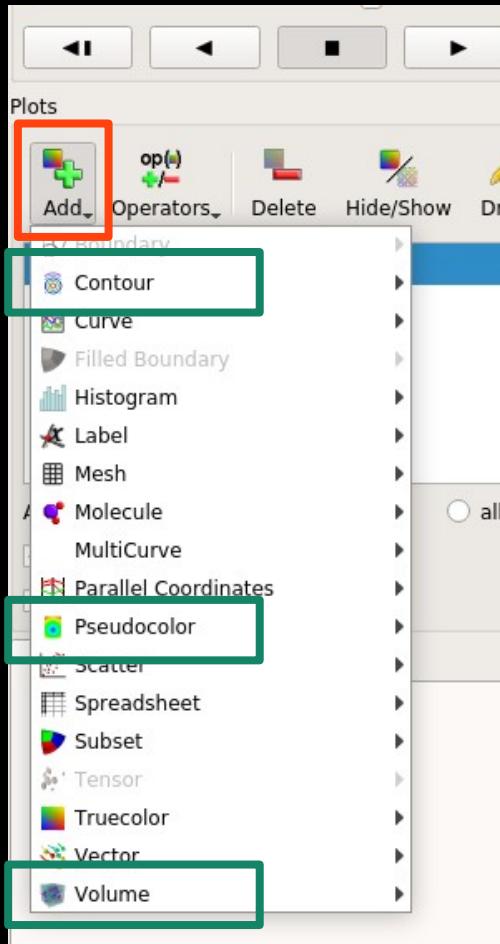
# Visit - GUI commands



Select multiple files with the same prefix

./<prefix>\_000000.hdf5

# Visit - GUI commands

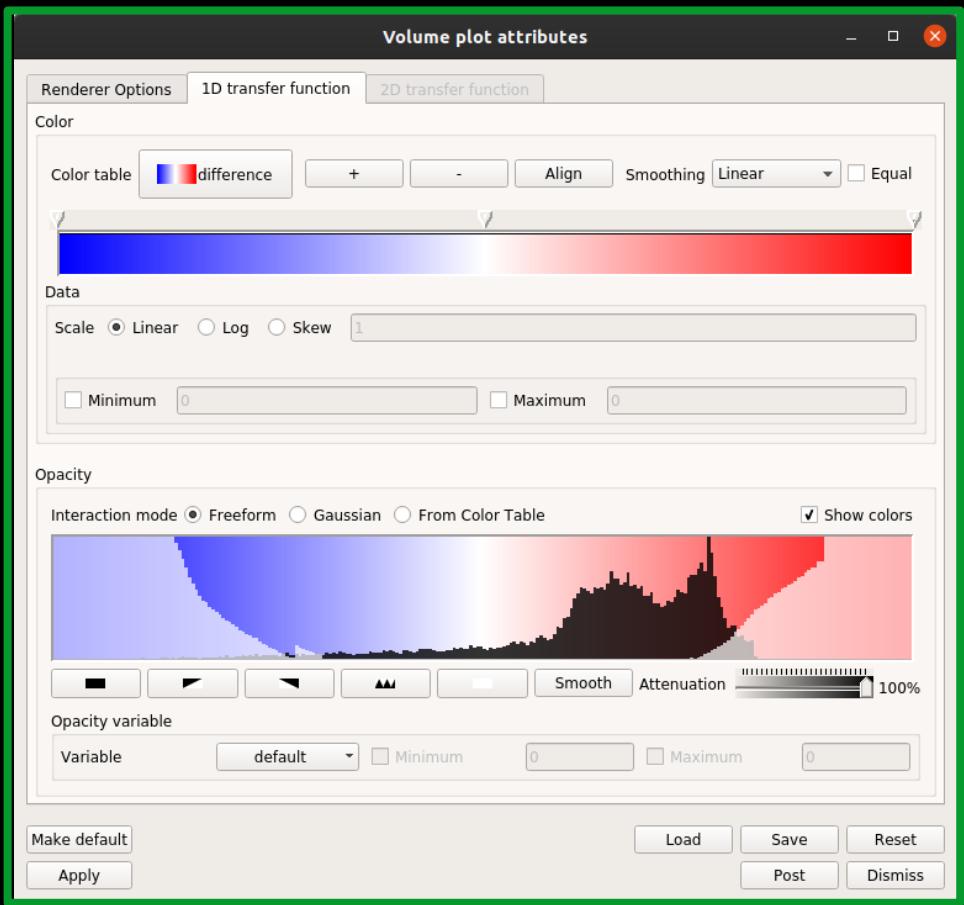
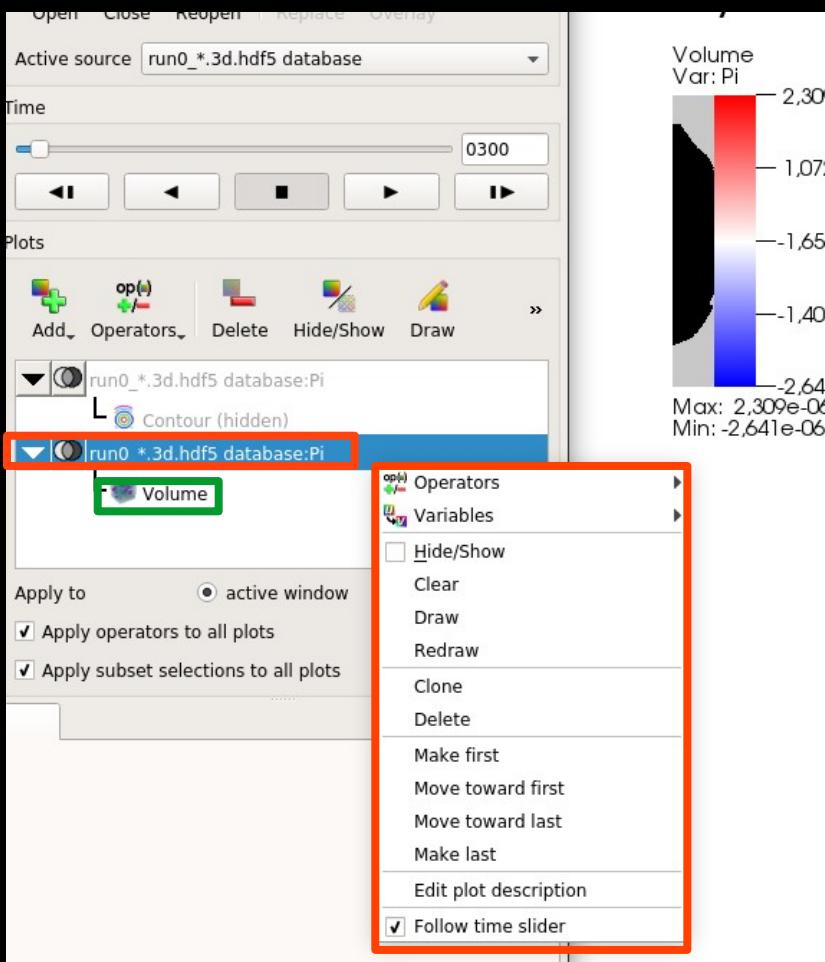


Different plotting functions:

- Contour, for 2D & 3D plots
- Pseudocolor, mainly for 2D plots.
- Volume, mainly for 3D plots

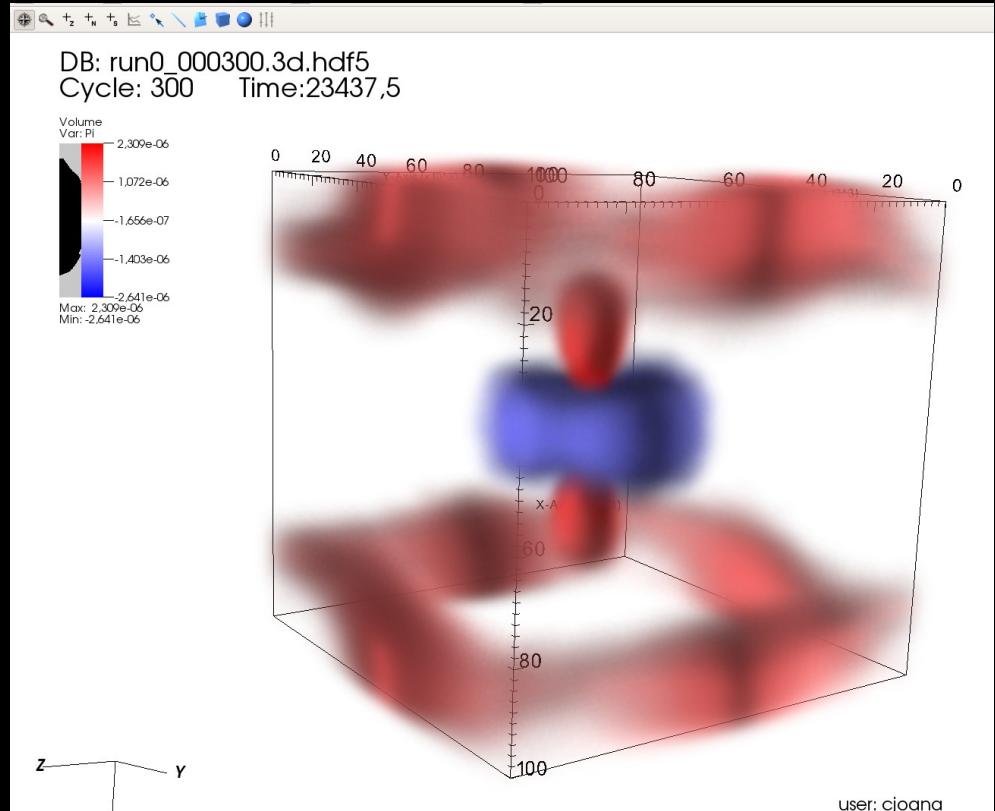
Feel free to play with the other options!

# Visit - GUI commands

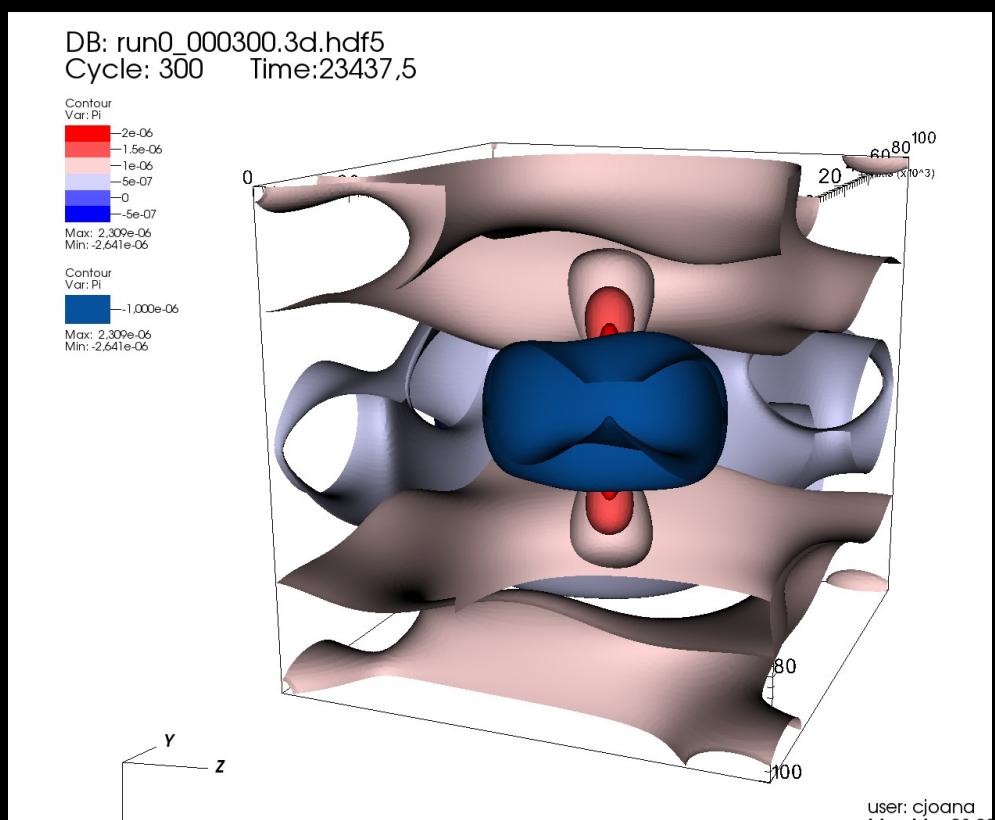


# Visit - GUI commands

Volume (3D)

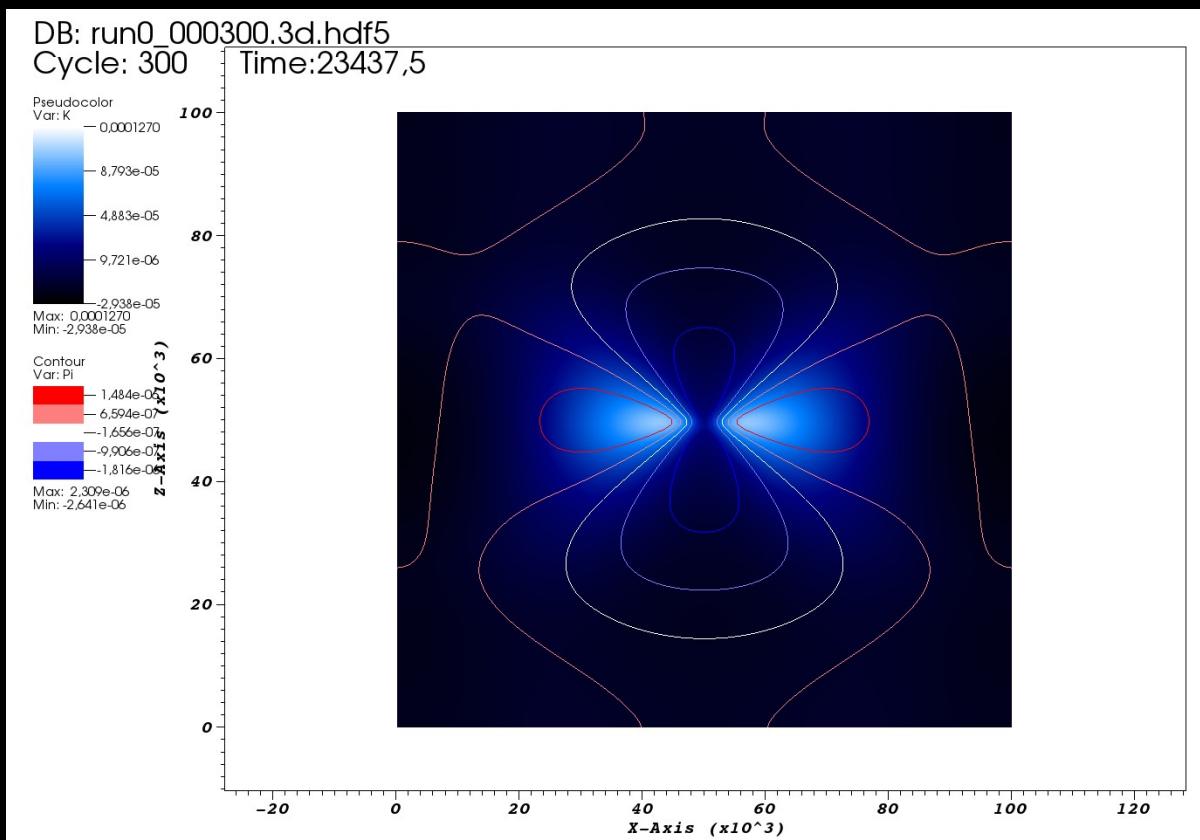
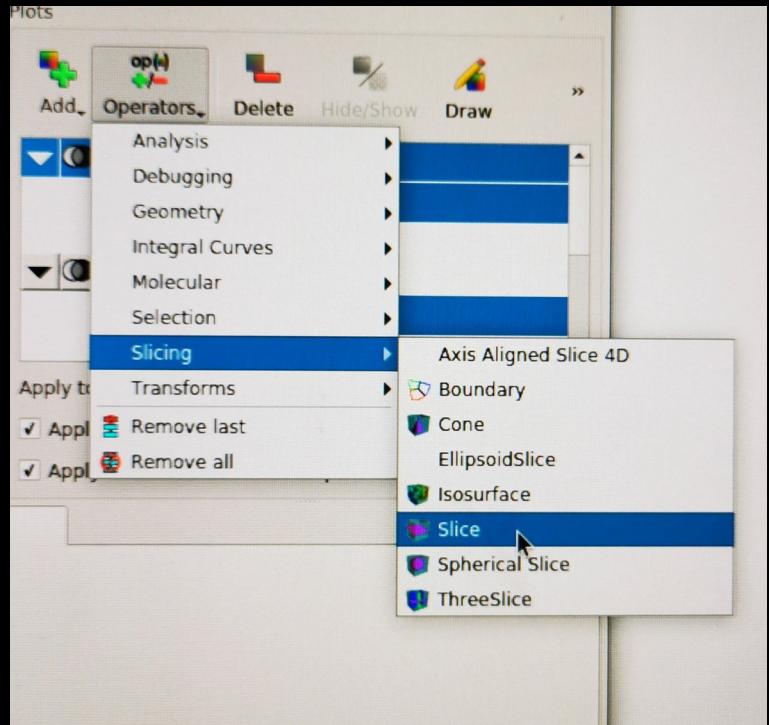


Contour (3D)

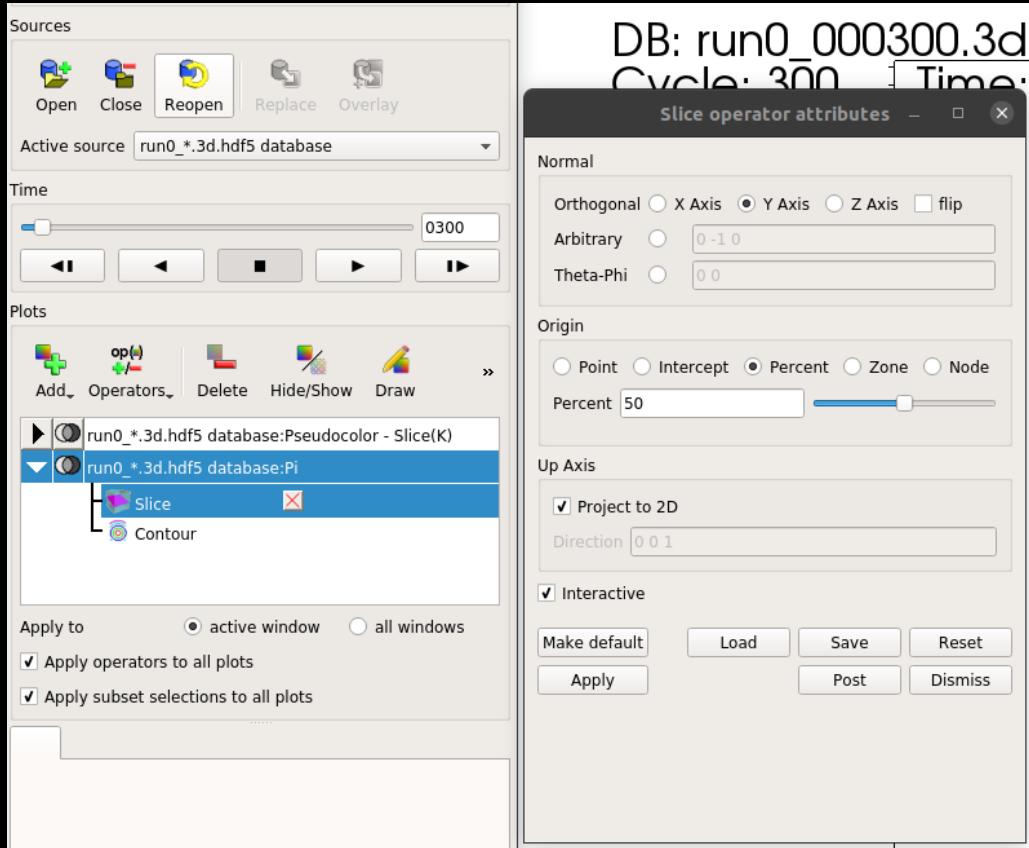


# Visit - GUI commands

## Pseudocolor + Contour (2D)



# Visit - GUI commands

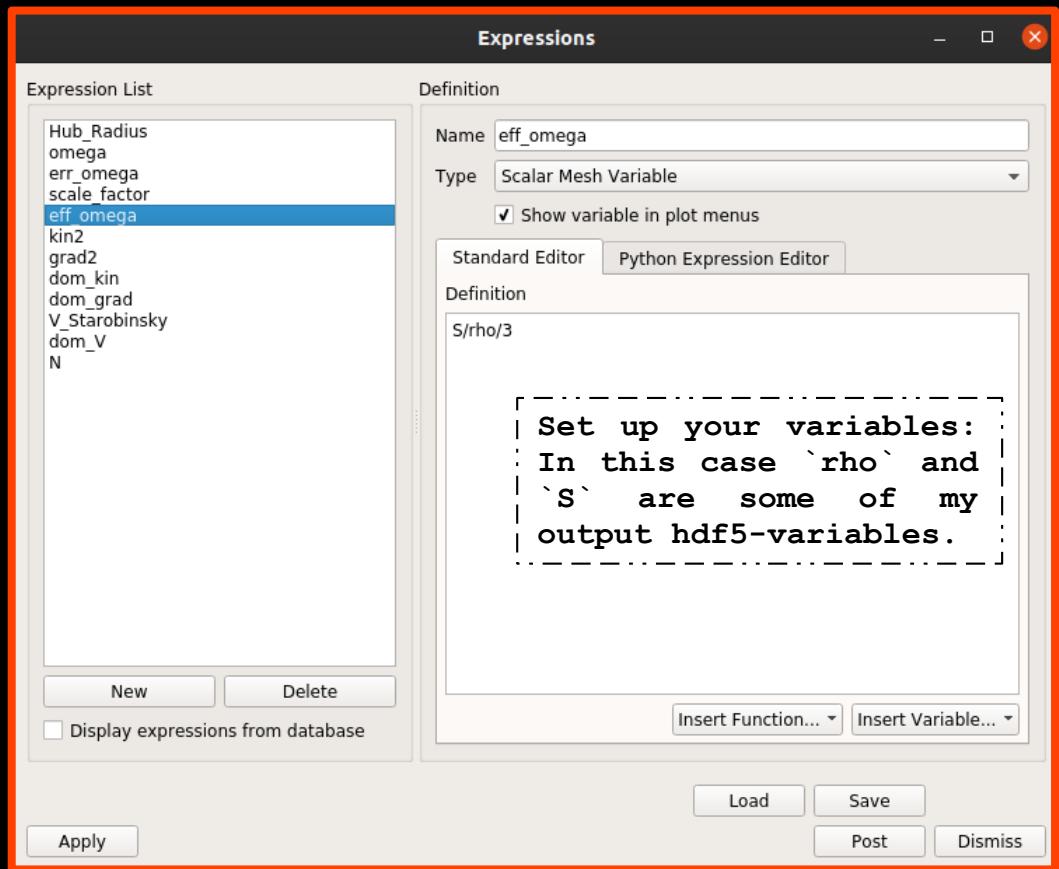
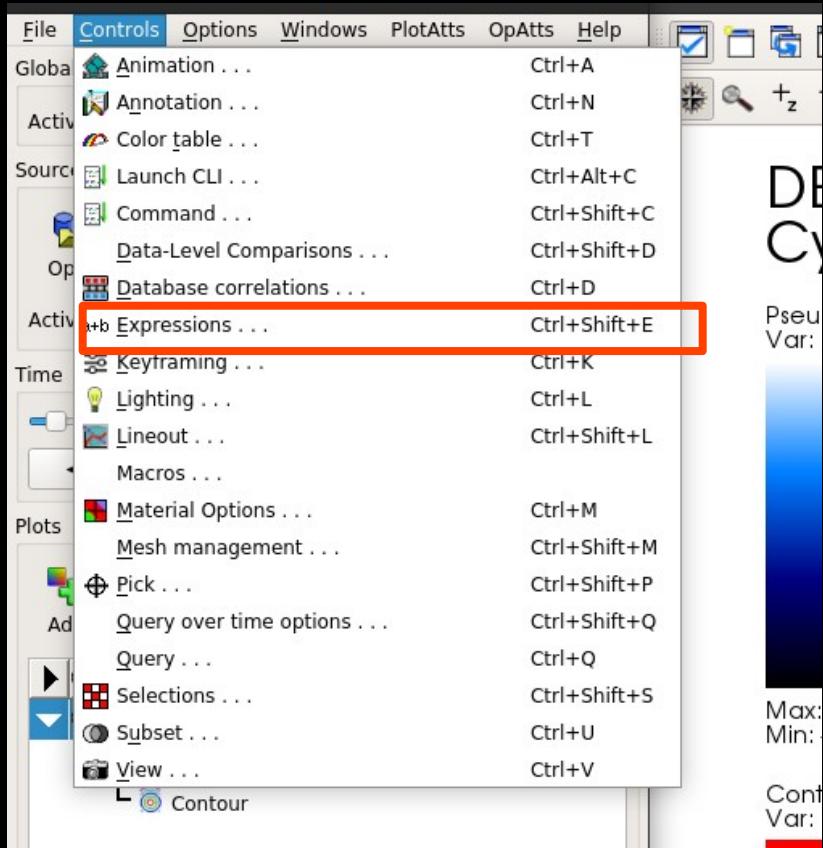


Suggestion:

- 1) Go to Slice settings
- 2) Select Orthogonal axis (e.g. Y)
- 3) Choose 'Percent' to select the cord.

# Visit - GUI commands

Add derived variable:

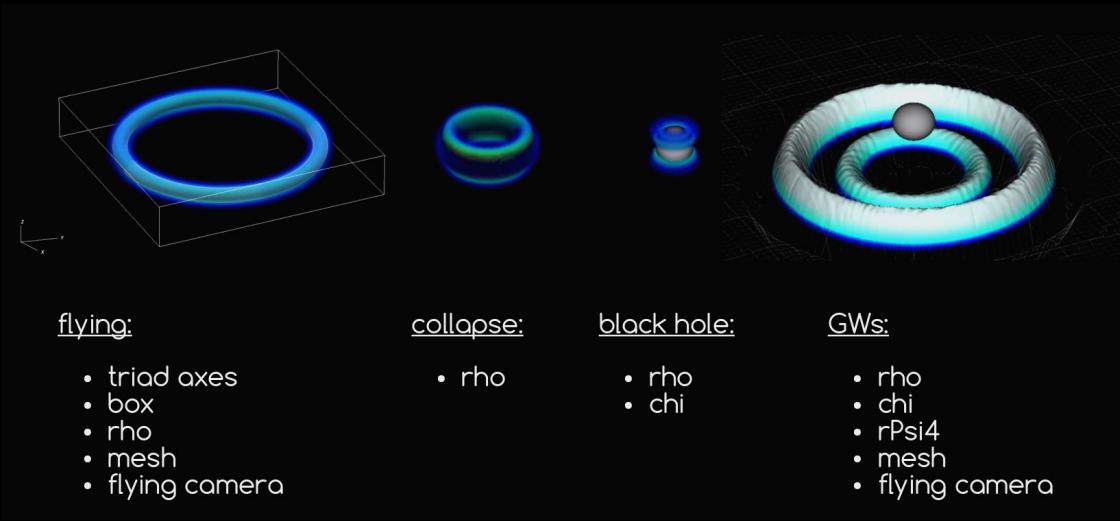


# VisIt – Script mode

---

- VisIt can also be used in `script` mode. But this is not covered on these slides.
- In `script` mode, beautiful animation can be made:

→ See Josu's slides from 2019, or ask him ;)





# Questions ?

