A system to allow easy & flexible creation, configuration, and inclusion of visualizations made by you

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Lowest layer first

You know best what questions need to be answered or explored

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Does not do your visualization programming for you

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Does not do your visualization programming for you

Does your Galaxy programming for you

You know best what questions need to be answered or explored

Does not do your visualization programming for you

Does your Galaxy programming for you

(we want to do your Galaxy programming then get out of your way)

### Visualization

Your code + user's data

### Visualization

### Your code:

### Visualization

Your code:

a VisualizationRegistry entry

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Your code:

a VisualizationRegistry entry

a .mako template

### Visualization

Your code:

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a .mako template

(that's it – no commits, controller methods, etc.)

### Visualization

Your code:

a VisualizationRegistry entry

a .mako template

#### A familiar pattern

```
visualizations_conf.xml.sample
    <!--
        Sample registry shipped with Galaxy
        -->

universe_wsgi.ini
...
#visualizations_conf_path = visualizations_conf.xml
...

visualizations_conf.xml
        <!--
        Include your visualizations and customize your server here
        -->
```

#### A familiar pattern

# Controls how Galaxy interacts with your visualization

```
<visualization name="Your Visualization">
    <!-- ... -->
</visualization>
<visualization name="Your Visualization 2">
    <!-- ... -->
</visualization>
<visualization name="Your Visualization 3">
    <!-- ... -->
</visualization>
```

How will the user start my visualization?
/visualization/show/myvis?data=lots

How will the user start my visualization?
/visualization/show/myvis?data=lots
When will the link be rendered?
when a data source passes your tests

#### When will the link be rendered?

#### Some source of data

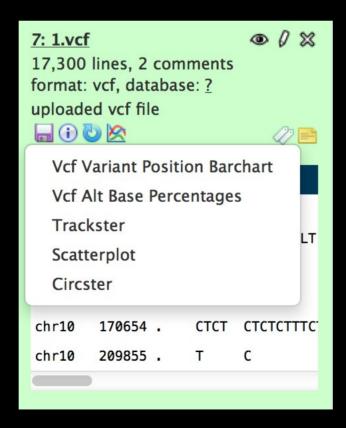
```
for dataset in history:
    visualization_links = registry.get_visualizations( dataset )
```

When will the link be rendered?

#### Some source of data

```
for dataset in history:
    visualization_links = registry.get_visualizations( dataset )
```

#### When will the link be rendered?



#### When will the link be rendered?

#### Some source of data

```
for dataset in history:
    visualization_links = registry.get_visualizations( dataset )

for dataset in library_folder:
    visualization_links = registry.get_visualizations( dataset )

for saved in user_visualizations:
    visualization_links = registry.get_visualizations( saved )

if user.is_admin():
    admin_vis_links = registry.get_visualizations( user )
```

When will the link be rendered?

Some source of data

VisualizationRegistry

Links for the user

## Outgoing Links



VisualizationRegistry



#### Some source of data

### VisualizationRegistry



VisualizationRegistry

#### tests

Some source of data

VisualizationRegistry

test model\_class

<model\_class>HistoryDatasetAssociation</model\_class>

#### Some source of data

VisualizationRegistry

test datatype

HistoryDatasetAssociation( 1.vcf ).datatype == Tabular == Pass



Link for the user

#### Some source of data

VisualizationRegistry

test are OR'd

```
<model_class>HistoryDatasetAssociation</model_class>
<test type="isinstance" test_attr="datatype"
    result_type="datatype">binary.Bam</test>

HistoryDatasetAssociation( 1.vcf ).datatype == Bam == Fails

<test type="isinstance" test_attr="datatype"
result_type="datatype">tabular.Tabular</test>

HistoryDatasetAssociation( 1.vcf ).datatype == Tabular == Pass

Link for the user
```

#### Some source of data

VisualizationRegistry

test attr string



Link for the user

```
How will the user start my visualization?
/visualization/show/myvis?data=lots
When will the link be rendered?
when a data source passes your tests
Can I pass more data through the link?
yes
```

### VisualizationRegistry

### Links for the user

#### VisualizationRegistry

#### Links for the user

#### to\_param

### VisualizationRegistry

#### Links for the user

#### to\_param

### VisualizationRegistry

#### Links for the user

#### to\_param



#### Links for the user

#### based on tests

### VisualizationRegistry

#### Links for the user

#### based on tests

### VisualizationRegistry

#### Links for the user

1<sup>st</sup> passed: 1<sup>st</sup> rendered

```
How will the user start my visualization?

/visualization/show/myvis?data=lots
When will the link be rendered?

when a data source passes your tests
Can I pass more data through the link?
```

yes

How do I get at that data then?
it's parsed and sent to your template

How do I get at that data then?

User clicked the link

visualization.render

VisualizationRegistry

Template 'resources'



#### User clicked the link visualization.render

```
def render( self, trans, visualization name, embedded=None, **kwargs ):
    Render the appropriate visualization template, parsing the
    query string `kwargs` into appropriate variables and resources
    (such as ORM models) based on this visualizations `param` data
    in visualizations conf.xml.
    URL: /visualization/show/{visualization name}
   # validate name vs. registry
    registry = trans_app_visualizations registry
    if not registry:
        raise HTTPNotFound( 'No visualization registry '
                          + '(possibly disabled in universe wsgi.ini)' )
    if visualization name not in registry listings:
        raise HTTPNotFound( 'Unknown or invalid visualization: ' +
                            visualization name )
    registry listing = registry.listings[ visualization name ]
```



#### User clicked the link visualization.render

```
def render( self, trans, visualization name, embedded=None, **kwargs ):
    Render the appropriate visualization template, parsing the
    query string `kwargs` into appropriate variables and resources
    (such as ORM models) based on this visualizations `param` data
    in visualizations conf.xml.
    URL: /visualization/show/{visualization name}
   # kwargs
   # validate name vs. registry
    registry = trans.app.visualizations registry
    if not registry:
        raise HTTPNotFound( 'No visualization registry '
                          + '(possibly disabled in universe wsgi.ini)')
    if visualization name not in registry.listings:
        raise HTTPNotFound( 'Unknown or invalid visualization: ' +
                            visualization name )
    registry listing = registry.listings[ visualization_name ]
```

#### visualization.render

### VisualizationRegistry



```
<visualization name="scatterplot">
 <data sources>
    <data source>
      <model_class>HistoryDatasetAssociation</model_class>
      <test type="isinstance" test_attr="datatype"
            result_type="datatype">tabular.Tabular</test>
      <to param param attr="id">dataset id</to param>
    </data source>
  </data sources>
  <params>
    <param type="dataset" var_name_in_template="hda"</pre>
           required="true">dataset id</param>
  </params>
  <template>scatterplot.mako</template>
</visualization>
```

#### VisualizationRegistry

#### template 'resources'

#### params

#### VisualizationRegistry

#### template 'resources'

#### to\_param => param

### VisualizationRegistry

#### template 'resources'

#### serialization

```
# some target object or other param
<HistoryDatasetAssociation>

# ... goes to the registry and gets serialized ...
<to_param param_attr="id">dataset_id</to_param>

# ... into a query string in a link that, when clicked, ...
/visualization/show/scatterplot?dataset_id=01234567890ABCD

# ... goes to the registry and gets un-serialized ...
<param type="dataset">dataset_id</param>

# ... into the target object for your template
<HistoryDatasetAssociation>
```

#### VisualizationRegistry

#### template 'resources'

#### serialization

```
# some target object or other param
<HistoryDatasetAssociation>

# ... goes to the registry and gets serialized ...
<to_param param_attr="id">dataset_id</to_param>

# ... into a query string in a link that, when clicked, ...
/visualization/show/scatterplot?dataset_id=01234567890ABCD

# ... goes to the registry and gets un-serialized ...
<param type="dataset">dataset">dataset_id</param>

# ... into the target object for your template
<HistoryDatasetAssociation>

(yay! NAQSP)
```

#### VisualizationRegistry

#### template 'resources'

#### template models

```
<param type="dataset" var_name_in_template="hda"
    required="true">dataset_id</param>

## ... into the target model! for your template
## <HistoryDatasetAssociation>
Name: ${hda.name}
## access to metadata
Data: ${hda.metadata.data_lines}
## access to 'eager loaded', related tables
Job settings: ${hda.creating_job.get_api_value()}
```

VisualizationRegistry

template 'resources'

#### secure models

```
parsed_param = controller.get_dataset( trans, encoded_dataset_id,
    check_ownership=False, check_accessible=True )

returned = trans.show_error_message(
    "There was an error rendering the visualization. " +
    "Contact your Galaxy administrator if the problem persists." +
    "<br/>Details: " + str( exception ), use_panels=False )
```

#### VisualizationRegistry

#### template 'resources'

#### many types

```
<param type="dataset">
                               ?d=<id>
                                           HistoryDatasetAssociation
<param type="visualization">
                               ?v=<id>
                                           Visualization(Revision)
<param type="hda or ldda">
                               ?d=<id>
                                           HDA or LDDA
<param type="dbkey">
                               ?dbk=hg18
                                           an existing genome build
                                           JSON -> python structure
<param type="json">
                               ?x={...}
<param type="float">
                                           parsed float
                               ?x=1.3
<param type="int">
                               ?x = -5
                                           parsed int
<param type="bool">
                               ?x=true
                                           True or False
... or lists of any of the above
<param csv="true"</pre>
                               ?d=<id1>,<id2>,<id3>
       type="dataset">
                                           a list of datasets [ ... ]
<param csv="true"</pre>
                               ?v=<id1>,<id2>,<id3>
                                           a list of visualizations
       type="visualization">
<param csv="true"</pre>
                               ?vs=3,0,-9
       type="int">
                                           a list of parsed ints
```

#### VisualizationRegistry

#### template 'resources'

#### multiple params

```
/visualization/show/complicated_visualization?
   allele_data=01234567890ABCD&
    ref_annotations=01234567890ABCE&
    ref_seq=hg18&
    chromosome=chr1&
    zoom_level=0.25&
    mark_positions=30033,35539,35542&
    quality_subgraph=F00FF0FFF8080ABE&
    view={start:25000,end:45000}
```

```
How will the user start my visualization?
      /visualization/show/myvis?data=lots
When will the link be rendered?
     when a data source passes your tests
Can I pass more data through the link?
                                       yes
How do I get at that data then?
 (we want to do your Galaxy programming
```

then get out of your way)

```
How will the user start my visualization?
/visualization/show/myvis?data=lots
When will the link be rendered?
when a data source passes your tests
Can I pass more data through the link?
yes
```

How do I get at that data then?
it's parsed and sent to your template

### Visualization

Your code:

a VisualizationRegistry entry

a .mako template

What libraries do I have to use?

(almost) any you'd like

What libraries do I have to use?

#### When to render?:

Server side Client side A mix of both

```
Server side – what to render?:

SVG
Images
HTML
```

```
Server side - rendering technology:

Matplotlib
Gnuplot + gnuplot
Rpy2 + R
Web Start + Java, etc.
```

```
Server side:

Matplotlib
Gnuplot + gnuplot
Rpy2 + R
Web Start + Java
(must be installed/available)
```

```
Server side SVG:

Matplotlib -> SVG
Gnuplot -> SVG
SVGfig (installed)
PySVG
Pygal -> SVG
(Mako) -> SVG
```

```
What libraries do I have to use?

(almost) any you'd like

Do I need to know JavaScript?

no (but js is good)
```

Do I need to know JavaScript?

Client side: SVG Canvas WebGL

Do I need to know JavaScript?

Client side:

SVG

Canvas

WebGL

Interactivity

```
Client side:
  SVG
      d3 (installed)
      Raphaël
      jQuery SVG
  Canvas
   WebGL
  Interactivity
```

```
Client side:

SVG

Canvas

paper, fabric, easel, processingJS

WebGL

PhiloGL, xtk

Interactivity
```

```
Client side:
    SVG
    Canvas
    WebGL
    Interactivity (support/UI)
    jQuery
    backbone & Galaxy models/libs
```

```
A mix of both:

Matplotlib -> SVG + d3.js

Gnuplot + JSON -> image + Canvas

Mako -> HTML + jQuery
```

```
A mix of both:
```

```
Matplotlib -> SVG + d3.js
Gnuplot + JSON -> image + Canvas
Mako -> HTML + jQuery
```

```
server = heavy lifting
client = interactivity
```

Do I need to know JavaScript?

```
A mix of both:
```

```
Matplotlib -> SVG + d3.js
Gnuplot + JSON -> image + Canvas
Mako -> HTML + jQuery
```

server = small,quick,light client = expense on user side

```
What libraries do I have to use?

(almost) any you'd like

Do I need to know JavaScript?

no (but js is good)
```

(we want to do your Galaxy programming then get out of your way)

```
What libraries do I have to use?

(almost) any you'd like

Do I need to know JavaScript?

no (but js is good)

How do I get the user's data?
```

(we want to do your Galaxy programming then get out of your way)

The Visualization Framework

### Visualization

User's data + your code

### Data

Possible sources of data:

The database Dataset contents

## Data

Possible sources of data:

The database resource APIs, the registry Dataset contents

#### Data

#### Possible sources of data:

The database

Dataset contents
bootstrapping into python/js
datasets API

## Data

#### Possible sources of data:

The database
Dataset contents
DataProviders

Get only the data you need in the format you want

Data source + format desired + settings

#### Currently defined in datatypes

```
@dataproviders.decorators.has dataproviders
class MyDataType( Tabular ):
   @dataproviders.decorators.dataprovider factory(
          'mycolumns', ColumnarDataProvider.settings )
   # returns an iterator object
   def mycolumn_dataprovider( self, dataset, **settings ):
       return ColumnarDataProvider( dataset.
          indeces=[0, 6, 3, 3, 94],
          column_types=[ 'str', 'int', 'float', 'str' ],
          **settings )
---> [ [ 'KMFDM_0002', 8675309, 1.68, '1.68', None ],
```

#### Currently defined in datatypes

Simple Filtering (line) no blanks, no comments

```
Simple Filtering
Limiting
start at line 1000, return only 500
(pagination)
```

```
Simple Filtering
Limiting
More Filtering
don't return lines matching regex
(or inverse)
```

Simple Filtering
Limiting
More Filtering
Formatting
raw lines, arrays, dictionaries
specific columns
format with metadata

```
Limiting
More Filtering
Formatting
Parsing
parse columns, key/values
parse using metadata.column_types
```

#### Multiple formats per source

```
@has_dataproviders
    @dataprovider_factory( 'column' )
    @dataprovider_factory( 'mycolumns' )
    @dataprovider_factory( 'genome-region' )
    @dataprovider_factory( 'interval-map' )
    @dataprovider_factory( 'seq-qal' )
```

# Inherited formats (no need to redefine)

#### (bootstrapping) into python:

```
bootstrapped_interval_list = list( dataset.datatype.dataprovider(
    dataset,
    'interval-map',
    limit=100,
    offset=200,
    regex_list=[ '^chr10' ], invert=True, ) )
# do something with the data
```

#### (bootstrapping) into js:

#### (via the datasets API) into js:

## Visualization

Your code + user's data

(we want to do your Galaxy programming then get out of your way)

# Persistence & Sharing

#### The Visualizations models and API

save load versioning links by slugline

# Persistence & Sharing

#### The Visualizations model and API

#### save

"Here's a link to the mumps epidemiology..."

#### load

"Got it. Good – so no one will be affected?"

#### versioning/'bookmarking'

"Oh. I set the points to transparent. Try this one."

#### links by slugline

"Well... that's probably more publishable."

## In the Future

A work in progress

Multiple target data sources

Higher layers: UI elements

Higher layers: base templates

Embedding

Run tools & create new data

Combine & separate visualizations

# Tusen takk! Thanks!

```
You, the Galaxy community!
UiO
Dave Clements
The Galaxy Team
```