

Integration of R to VTK: Adding Statistical Computation to a Visualization Toolkit

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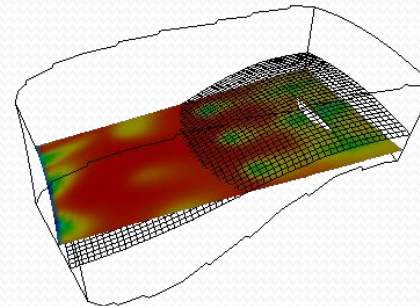
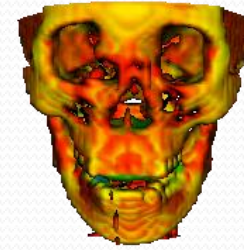
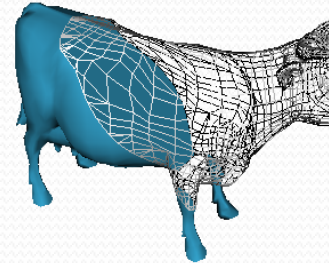
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Basics – VTK

The **Visualization Toolkit (VTK)** is an open-source, freely available software system for 3D computer graphics, image processing and visualization.

- Scalar, vector, tensor, texture, and volumetric methods
- Advanced modeling techniques such as:
 - Implicit modeling,
 - Polygon reduction,
 - Mesh smoothing,
 - Cutting,
 - Contouring,
 - Delaunay triangulation.
- VTK:
 - Has an extensive information visualization framework,
 - Has a suite of 3D interaction widgets,
 - Supports parallel processing, and
 - Integrates with various databases on GUI toolkits such as Qt and Tk.
 - Cross-platform and runs on Linux, Windows, Mac and Unix platforms.



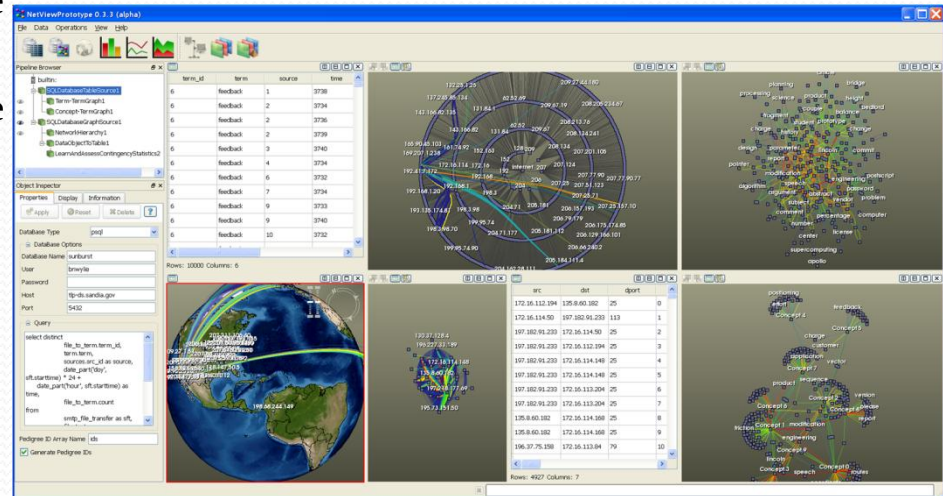


Basics – Titan

The **Titan Informatics Toolkit** is a collaborative effort between Sandia National Laboratories and Kitware Inc. It represents a significant expansion of the Visualization ToolKit (VTK) to support the ingestion, processing, and display of informatics data. By leveraging the VTK engine, Titan provides a flexible, component based, pipeline architecture for the integration and deployment of algorithms in the fields of intelligence, semantic graph and information analysis. Scalar, vector, tensor, texture, and volumetric methods

Titan Provides a set of data structures and algorithms for:

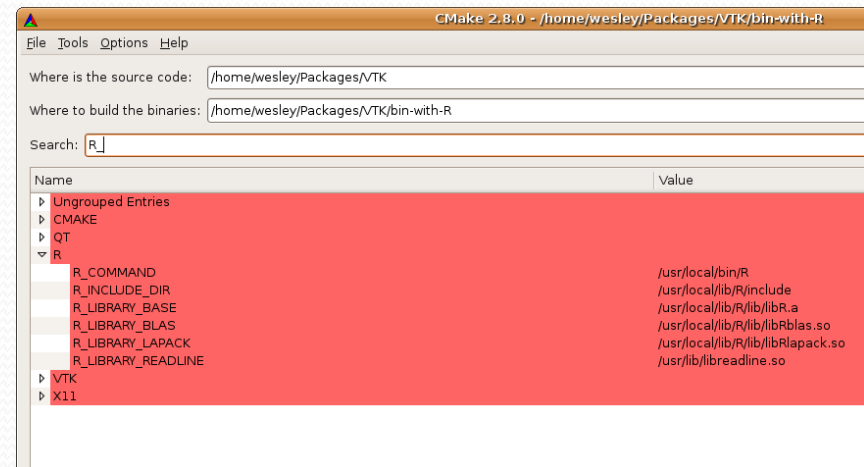
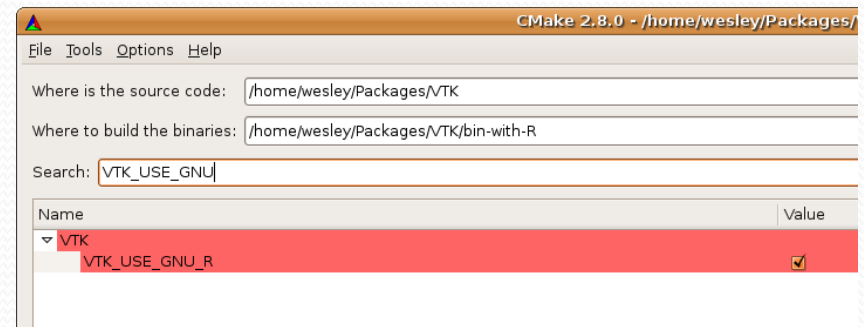
- Translation between VTK data structures and Graph/Tree data structures
- Access to databases
- Graph and Tree Layouts
- Graph and Tree Analysis



An application based on the Titan Informatics Toolkit is shown here displaying multiple views including table, graph and geospatial all semantically linked.

Enabling R in a VTK Build

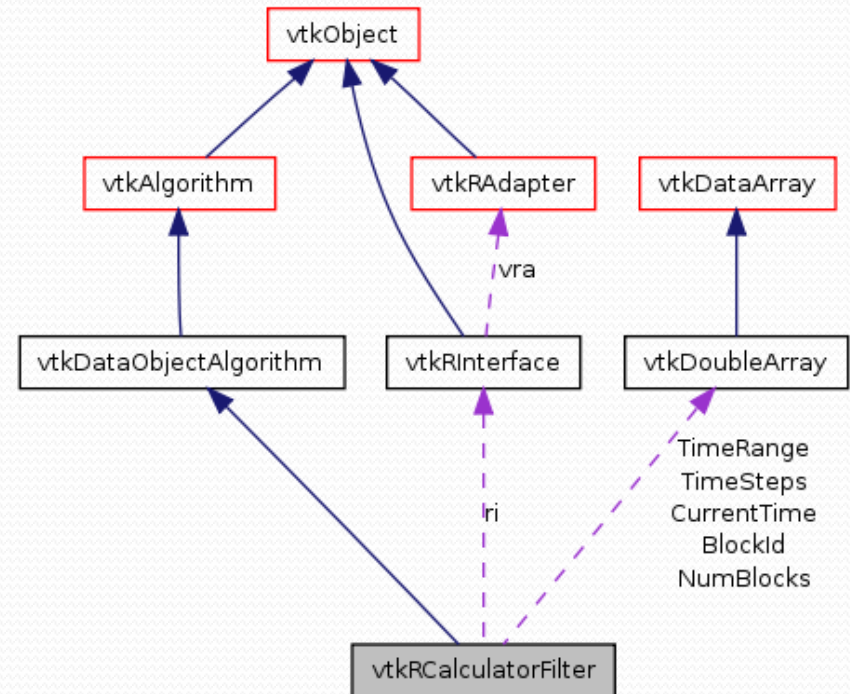
- Obtain R Source
 - Build
 - Install
- VTK Uses CMake
 - Download VTK
 - During VTK configuration
 - `cmake-gui <VTK-source>`
 - Set `VTK_USE_GNU_R` On
 - Hit `<Configure>`
 - VTK uses the “R RHOME” command to determine the R configuration
 - Verify the R environment
 - Complete the VTK configuration
 - Build VTK as normal



Interface to R

- vtkRCalculatorFilter
 - VTK filter API
- vtkRInterface
 - VTK Interface to embedded R Interpreter
 - Creates/Manages R Interpreter Instance
- vtkRAdapter
 - Lowest level of interface
 - Converts VTK Data to R SEXP
 - Converts R SEXP to VTK Data

Classes originally authored by Thomas Otahal at Sandia National Labs



```
vtkSmartPointer<vtkRCalculatorFilter> calc =
    vtkSmartPointer<vtkRCalculatorFilter>::New();
calc->SetRoutput(0);
calc->SetInputConnection(tab->GetOutputPort());
calc->PutArray("0", "metabData");
calc->GetArray("correl", "correl");
calc->SetRscript("correl<-cor(metabData)");
calc->Update();
calc->GetOutput()->Print(std::cout);
```

vtkCalculatorFilter Flow

- **Public Member Functions**

- virtual const char * **GetClassName** ()
- virtual int **IsA** (const char *type)
- void **PrintSelf** (ostream &os, **vtkIndent** indent)
- void **PutArray** (const char *NameOfVTKArray, const char *NameOfRvar)
- void **GetArray** (const char *NameOfVTKArray, const char *NameOfRvar)
- void **RemoveAllPutVariables** ()
- void **RemoveAllGetVariables** ()
- void **PutTable** (const char *NameOfRvar)
- void **GetTable** (const char *NameOfRvar)
- virtual void **SetRscript** (const char *)
- virtual char * **GetRscript** ()
- virtual void **SetScriptFname** (const char *)
- virtual char * **GetScriptFname** ()
- virtual void **SetRoutput** (int)
- virtual int **GetRoutput** ()
- virtual void **SetTimeOutput** (int)
- virtual int **GetTimeOutput** ()
- virtual void **SetBlockInfoOutput** (int)
- virtual int **GetBlockInfoOutput** ()
- virtual int **ProcessRequest** (**vtkInformation** *request, **vtkInformationVector** **inputVector, **vtkInformationVector** *outputVector)

- **Static Public Member Functions**

- static **vtkRCalculatorFilter** * **New** ()
- static int **IsTypeOf** (const char *type)
- static **vtkRCalculatorFilter** * **SafeDownCast** (**vtkObject** *o)

Initialize

- Create vtkRInterface
- Initialize locals

Execute

- Convert input->SEXP
- Load script if requested
- Evaluate script
- Convert SEXP->output

Destroy

- Remove vtkRInterface
- Clean up locals

vtkCalculatorFilter Flow

- Public Member Functions

- virtual const char * **GetClassName** ()
- virtual int **IsA** (const char *type)
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Initialize

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Standard VTK
Boilerplate

Convert input->SEXP
Load script if requested
Evaluate script
Convert SEXP->output

Destroy

- Remove vtkRInterface
- Clean up locals

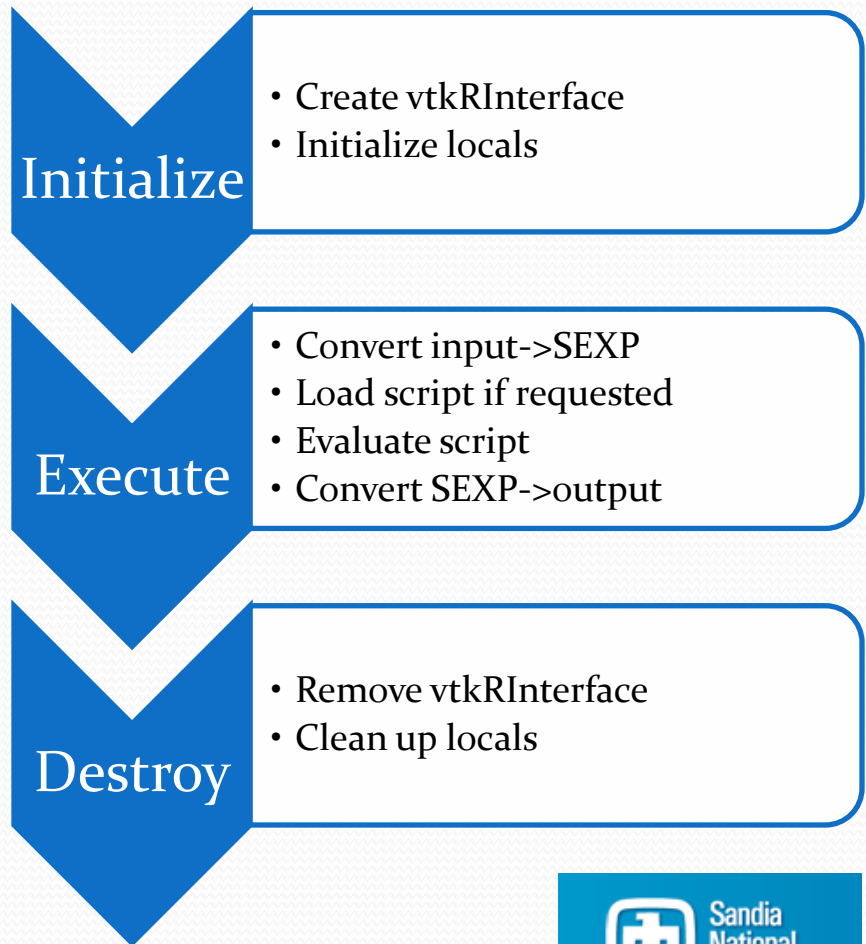
vtkCalculatorFilter Flow

Inputs:

- void **PutArray** (const char *NameOfVTKArray, const char *NameOfRvar)
- void **PutTable** (const char *NameOfRvar)
- void **RemoveAllPutVariables** ()

Outputs:

- void **GetArray** (const char *NameOfVTKArray, const char *NameOfRvar)
- void **RemoveAllGetVariables** ()
- void **GetTable** (const char *NameOfRvar)



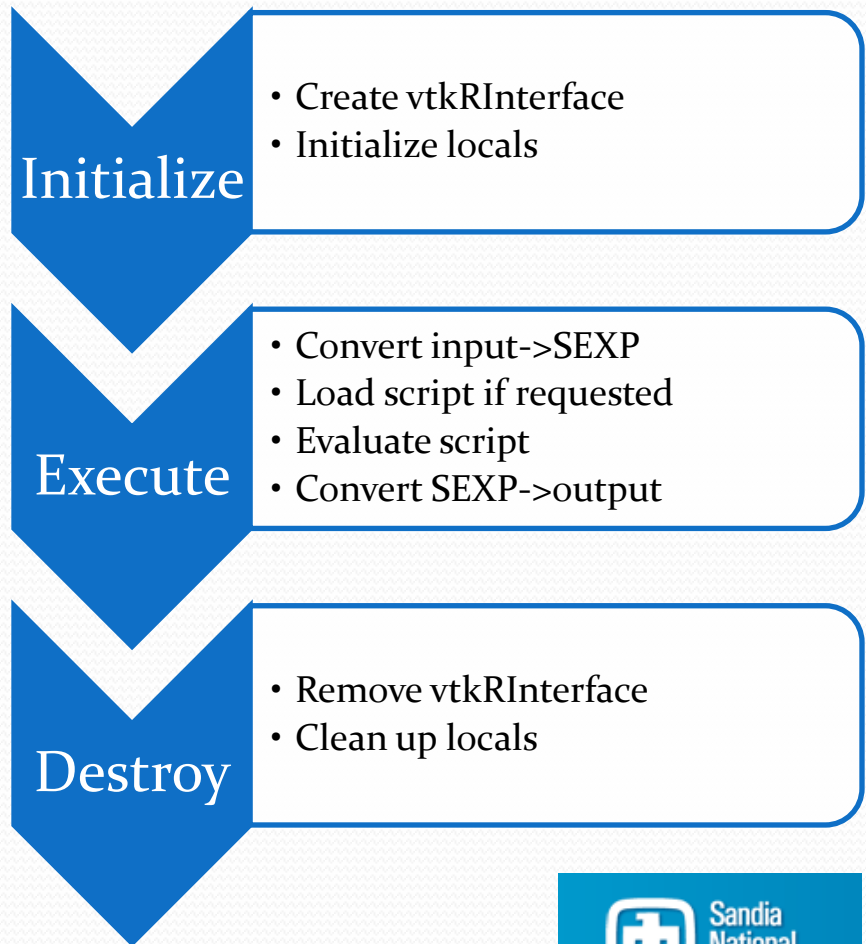
vtkCalculatorFilter Flow

R Scripts Control:

- virtual void [SetRscript](#) (const char *)
- virtual char * [GetRscript](#) ()
- virtual void [SetScriptFname](#) (const char *)
- virtual char * [GetScriptFname](#) ()
- virtual void [SetRoutput](#) (int)
- virtual int [GetRoutput](#) ()

Execution Control:

- virtual void [SetTimeOutput](#) (int)
- virtual int [GetTimeOutput](#) ()
- virtual void [SetBlockInfoOutput](#) (int)
- virtual int [GetBlockInfoOutput](#) ()



Auto-Generation

- Now consider a specific set of R functionality
- Filter needs to:
 - Identify inputs
 - Identify the R operations
 - Identify R outputs
- Why not automate this?
- Automation is an interaction between:
 - A R filter template
 - `vtkRGenericInterface.h.in`
 - `vtkRGenericInterface.cxx.in`
 - A GUI/Interaction module
 - E.g. `vtkWebModuleHandler.h`
 - E.g. `vtkWebModuleHandler.cxx`
 - A CMake macro to connect the two and define the inputs

Auto-Generation

- **Macro:**

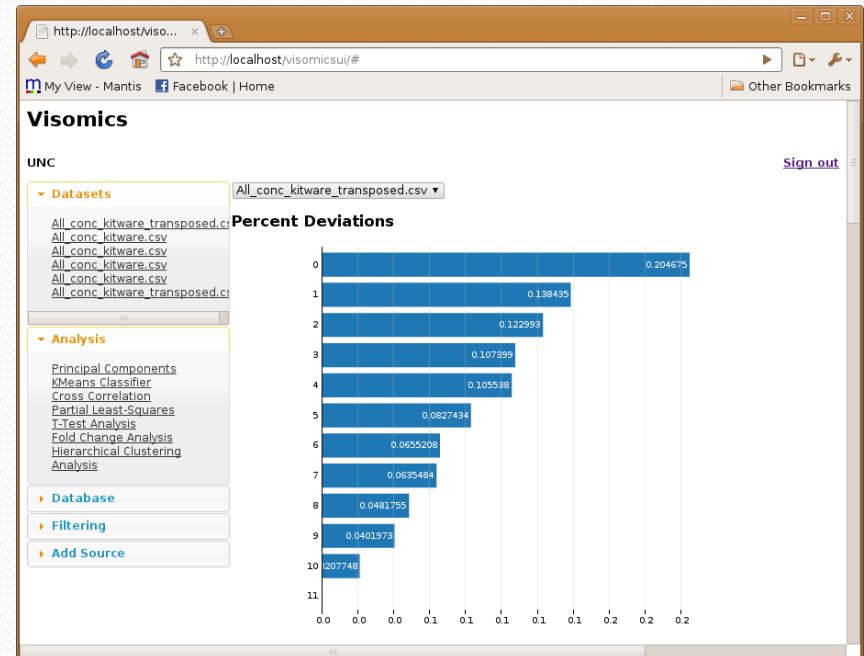
```
Visomics_CREATE_MODULE(  
  MODULE_NAME      "voXCorrelHandler"  
  ANALYSIS         "xcorrel"  
  VISUALIZATION_BASE "voWebModuleHandler"  
  MODULE_INPUTS    "metabData"  
  MODULE_OUTPUTS   "correl"  
  R_COMMAND        "correl<-cor(metabData)"  
  VISOMICS_SHOW_CORRELATION  
                  "1"  
  VISOMICS_SHOW_ROTATED_COORDINATES  
                  "0"  
  VISOMICS_SHOW_ROTATION_MATRIX  
                  "0"  
  VISOMICS_SHOW_STANDARD_DEVIATION  
                  "0"  
  VISOMICS_SHOW_GENERIC_ARRAY  
                  "0"  
)
```

- **Currently Available:**

- "voHierarchicalClusterHandler"
- "voFoldChangeHandler"
- "voTTestHandler"
- "voPLSHandler"
- "voXCorrelHandler"
- "voPCAHandler"

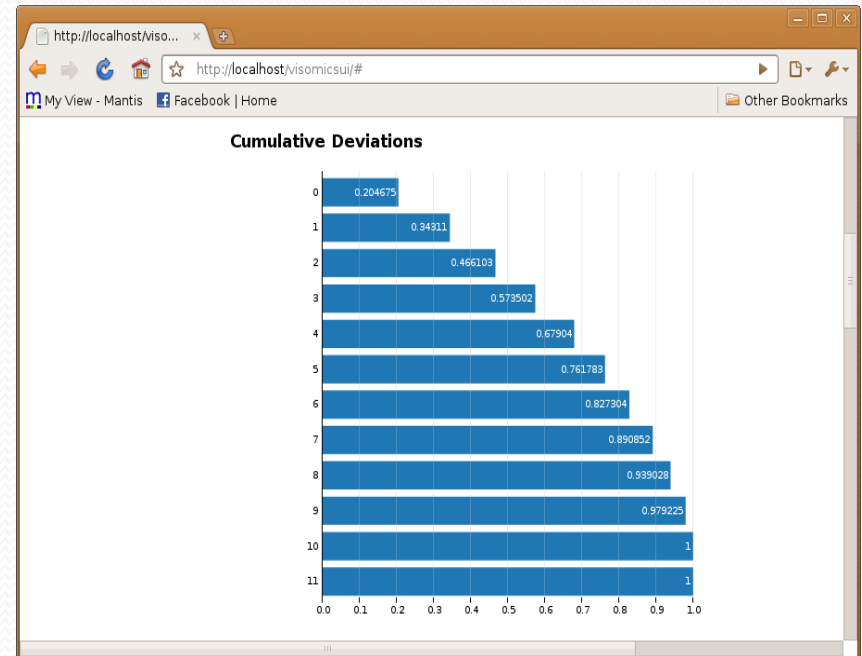
'Omics Application

- One application is a research vehicle for 'omics efforts
- Input is measured gene expression/metabolite concentrations
 - Multiple experiments
 - Cases and Controls
- Output is relationships
 - Correlations
 - Significance



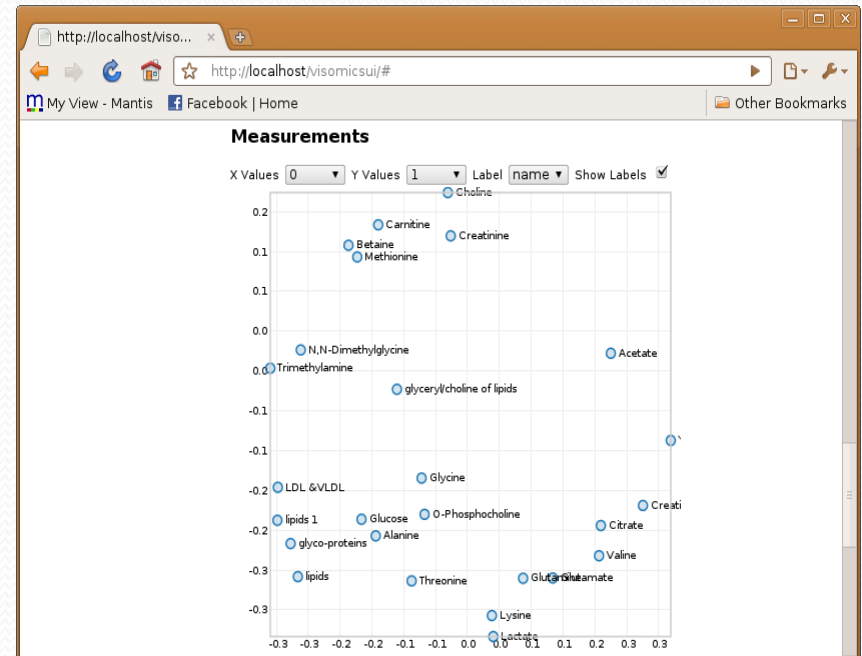
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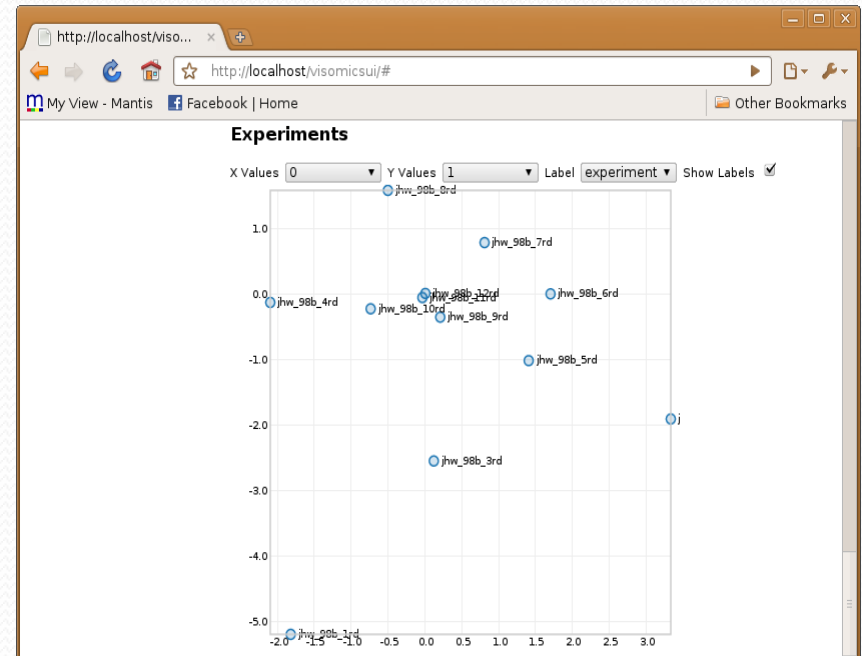
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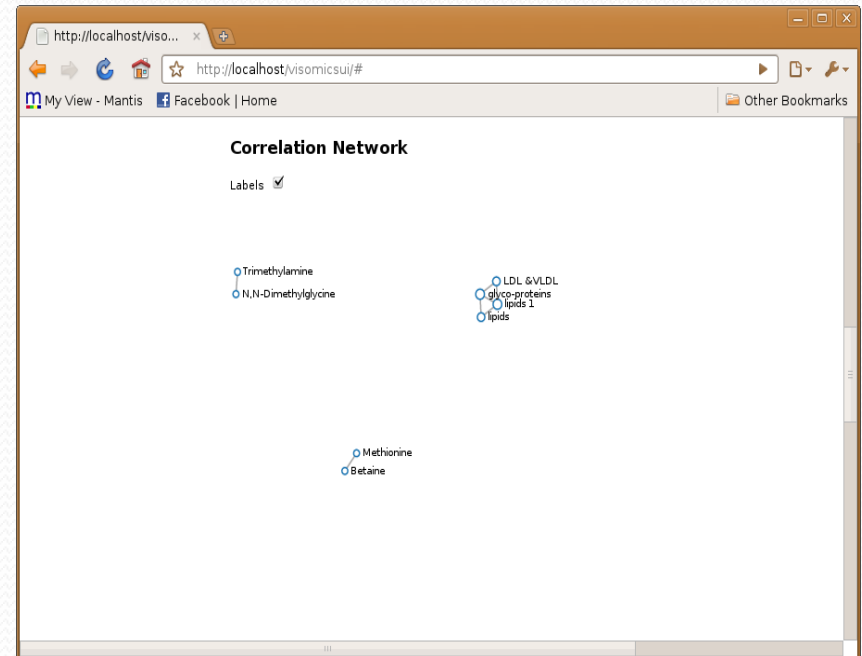
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End