

AGINFRA+ data science workshop

PLANT PHENOTYPING EXPERIMENTS ANALYSIS

1. Accessing the Food Security VRE
2. Exploring phenotyping studies
3. Retrieve data
4. Visualize data
5. Analyze data

1. ACCESSING THE FOOD SECURITY VRE

To access the VRE, open a web browser and go to the website : <https://aginfra.d4science.org/> On this AgInfra+ Gateway site, please log in with your account details.

You should see the VREs that are accessible to you. Select the Food Security VRE. This will open the website in your browser.

The screenshot displays the Food Security VRE dashboard. The top navigation bar includes links for Food Security, Administration, Members, Analytics, Semantics, Discovery, Visualization, and Plant Phenotyping installations. The main content area is divided into three columns. The left column shows 'Statistics' and 'Your Stats in FoodSecurity' with a user profile and activity metrics. The middle column shows a 'Share updates' section with a text input and a 'Share' button, followed by a 'News feed' with a post from Panagiota Koltida. The right column shows an 'About' section with the Food Security logo and a description of the VRE.

2. EXPLORING PHENOTYPING STUDIES AND RETRIEVING DATA

Open the Studies Exploration. Select the server PHIS_SANDBOX.

BRAPI Compliant Servers Exploration Studies Exploration Study Variables Preview Observations Data

Choose the BRAPI server to look in:

PHIS_SANDBOX

Season (Enter a year to filter)

Study Name (search with the name of the study)

Choose a specie:

all

Find

Studies Exploration

This application is meant to explore open data stored in BRAPI compliant databases.

PHIS_SANDBOX studies

Show 25 entries Search:

studyDbId	studyName	startDate	endDate	active
http://www.opensilex.org/demo/DIA2017-1	MAU17-PG	2017-05-19	2017-09-22	false
http://www.opensilex.org/demo/DMO2012-1	ZA12	2012-01-01	2012-12-31	false

studyDbId studyName startDate endDate active

Showing 1 to 2 of 2 entries Previous 1 Next

You should see at least 2 observations. In this exercise we will work with the study ZA12 data. We are interesting in plant height data. Let's check if there are plant height data on this study.

Copy the ExperimentURI and go to the tab "Preview study observations". You can see the first observations measurements of the study.

BRAPI Compliant Servers Exploration Studies Exploration Study Variables Preview Observations Data

Enter the studyDbId

http://www.opensilex.org/demo/DMO2012-1

See Data Download the table

Study Observations

Here you can retrieve 20 first observations data from one study

This study contains 506 observations

Show 25 entries Search:

observationDbId	observationLevel
http://www.opensilex.org/demo/id/data/tdohn3xtthfwbgkav2hg35ct6upj5rxu7drj2t7oalg64pnqf2e9c12181584b0c8ebe904e1694a311	http://www.opensilex.org/vocabulary/oeso#Pl
http://www.opensilex.org/demo/id/data/svmz52tcypbh2ziukpw46ouxmwzgjytcabdlts4vweedsrkunhqa2ac74a7f1034a859da09b14dde7b27e	http://www.opensilex.org/vocabulary/oeso#Pl
http://www.opensilex.org/demo/id/data/cclsgjtw4fyagqbpfd27fsjmk7jq3n6hawwdt6felt2zjqgecq6970e1d22cf8460a8bf0f996f7ca14ea	http://www.opensilex.org/vocabulary/oeso#Pl
http://www.opensilex.org/demo/id/data/zdg7nkwejkfhkdgxz34ae5q6cxz3d6vk22u5bxhqn2jalix2q706009e5a7cd444db4f59855129d3476	http://www.opensilex.org/vocabulary/oeso#Pl
http://www.opensilex.org/demo/id/data/77tzcpglwq5vbqeomlfr2wqfsftawb36lbqk4s7vxf64smzczyad4617f025f8147b38ee1a6aca11973b4	http://www.opensilex.org/vocabulary/oeso#Pl
http://www.opensilex.org/demo/id/data/ae34ggmzh3mbofvt4pmjpr234t62jxhad5qrd6dr264q3vdfef3773a64cc0d4e0cb5f280f326d01906	http://www.opensilex.org/vocabulary/oeso#Pl

3. RETRIEVE DATA

You will use a dataminer algorithm to get all data of the ZA12 study. Go to *Analytics/Dataminer* to execute the algorithm **Brapi Get Studies Observations**. (this algorithm is inside the category "Data extraction")

Fill the different parameters as presented below (the login and password parameters can be used to access to private studies):

The screenshot shows the Dataminer web interface. On the left, the 'Operators' panel lists several operators: CHARTS (1), CURVE FITTING (2), and DATA EXTRACTION (5). Under DATA EXTRACTION, there are three operators: 'Brapi Get Studies', 'Brapi Get Studies Observations', and 'Brapi Get Variables'. The 'Brapi Get Studies Observations' operator is selected. On the right, the 'Operator' configuration panel is shown. It has a 'Tools' section with 'Remove All Operators' selected. Below this, a text box describes the operator: 'Returns all observations where there are measurements for the given study and if so the given observation variables (Published by Alice Boizet (aliceboizet) on 2019/09/16 14:18 GMT)'. The 'Parameters' section includes fields for 'DBServerURL' (PHIS_SANDBOX), 'studies' (.org/demo/DMO2012-1), 'variables' (all), 'login' (none), and 'password' (none).

Then *Start Computation*

You should see this execution loading bar :

The screenshot shows the Dataminer web interface with the 'Computations Execution' panel active. It displays the status of a computation for 'Brapi Get Studies Observations'. The status is 'Computation Complete', indicated by a green bar. Below this, it says 'The computation Brapi Get Studies Observations finished.' and 'The algorithm produced Multiple Results.' There is a 'Files' section with a 'Log of the computation' and a 'Download File' button.

This algorithm creates a csv file and a json file with the study data. They are stored in your workspace under the dataminer folder.

4. VISUALIZE DATA

There are 2 ways to visualize the data. You can do it by using the visualization tools or you can use Rstudio to generate graphs.

Visualize data in the visualization tool

The workspace is not integrated with this tool yet. So you will first need to download the csv file created by the dataminer. If you didn't do it at the end of the dataminer computation, you can go to your workspace:



button at the upper left corner. The files are stored in DataMiner/Output Data Sets.

When you have downloaded the csv file, you can go to the *visualization/Create Graphs* to create a new graph.

In Data, upload the csv file you got from the studies exploration tool.

Fill the General tab this way :

General

Data

Filters

Transformations

Documents

Label*

ZA12_plantHeight

Description

Description

Type

Scatter

Available Types

Select other chart types that will be available as options

Group By

observationUnitName

X Axis*

observationTimeStamp

X Axis Label*

date

Y Axis*

value

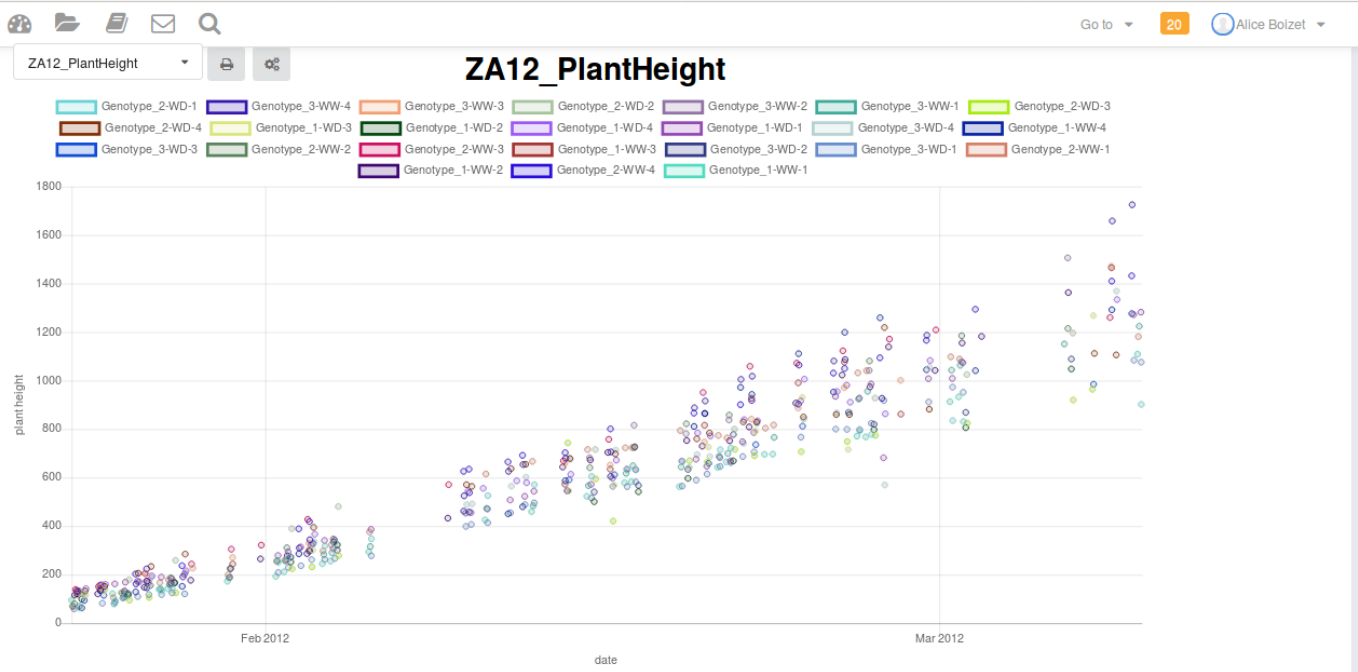
Y Axis Label*

plant height

Color

observationUnitName

You can go to *visualization/Create Graphs* and see your chart.

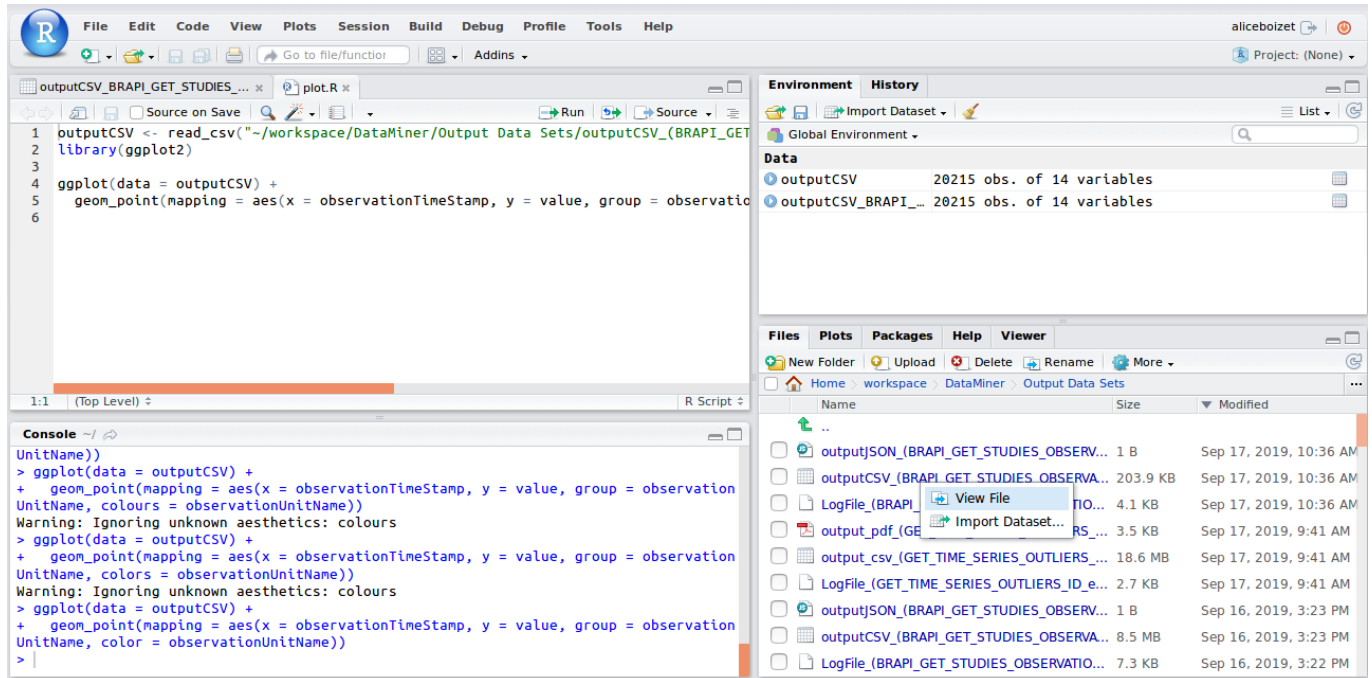


Visualize data in Rstudio

An other good way to visualize is to use R. Indeed there are great libraries such as ggplot2 to build charts. There are also very helpfull galleries with a lot of code examples (<https://www.r-graph-gallery.com/>)

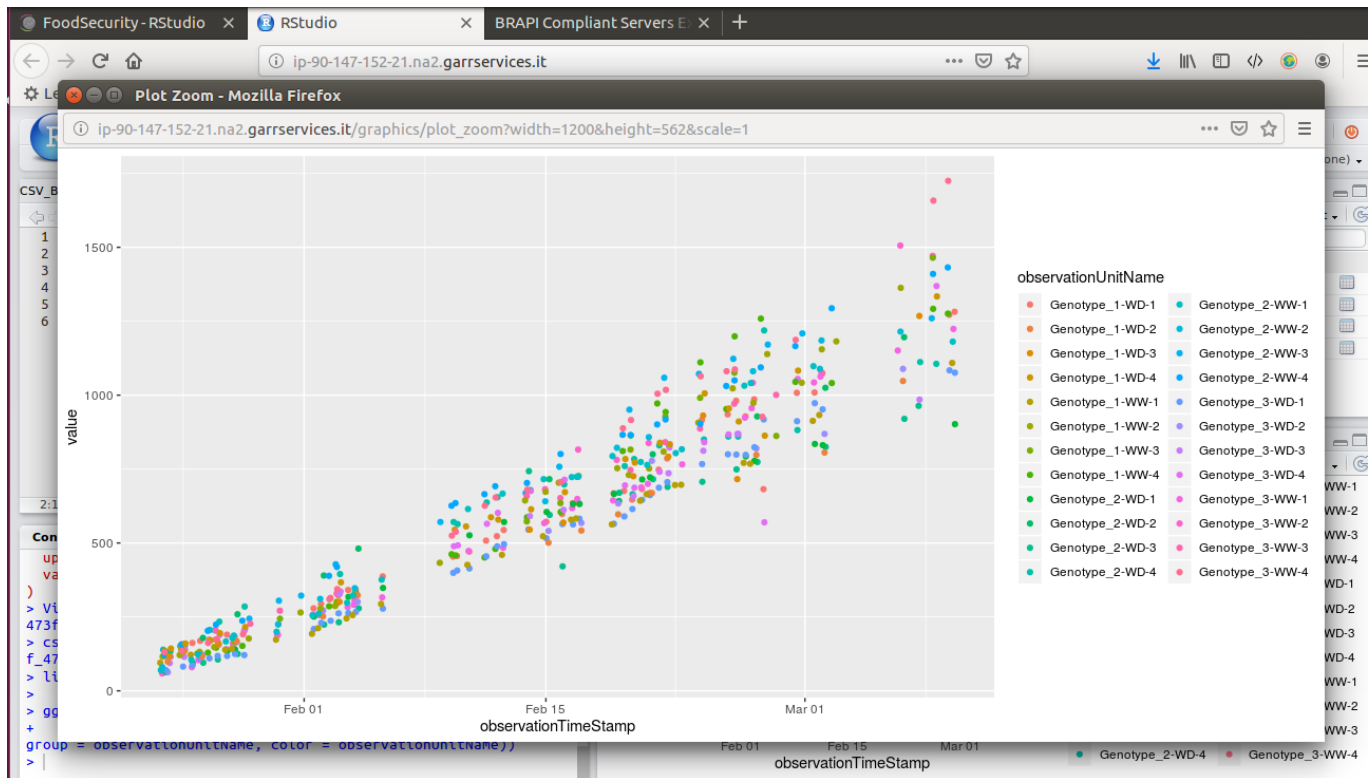
Go to *Analytics/Rstudio* to open **Rstudio**. Find your data csv file (created by the dataminer) in the *Files* tab at the bottom right corner. It is stored in your workspace under *DataMiner/Output Data Sets* folder.

Click on the file to "import Dataset".



You can now use the R console to visualize your data. To create a simple chart, you can use the code below which creates the following chart:

```
library(ggplot2)
ggplot(data = outputCSV) +
  geom_point(mapping = aes(x = observationTimeStamp, y = value, group = observationUnitName, color = observationUnitName))
```



ANALYZE DATA

Reuse an algorithm to detect outliers

You would like to detect the outliers on your data. To do that, you could use Rstudio (or even Jupyter) and make a script. But you are very lucky, another VRE member has already done a similar script and he has imported it in the dataminer tool as a black box (the algorithm name is **Get Time Series Outliers**). So all you need is to go to your workspace and right click on your data csv file and select "execute DM task".

Name	Owner	Type	Last Update	Size
outputJSON_(BRAP1_GET_STUDIES_OBSERVA...)	Alice Boizet	application/json	17 Sep 10:36 AM 2019	1 KB
outputCSV_(BRAP1_GET_STUDIES_OBSERVA...)	Alice Boizet	text/csv	17 Sep 10:36 AM 2019	203 KB
LogFile_(BRAP1_GET_STUDIES_OBSERVA...)	Alice Boizet	text/plain	17 Sep 10:36 AM 2019	4 KB
output_pdf_(GET_STUDIES_OBSERVA...)	Alice Boizet	application/pdf	17 Sep 09:41 AM 2019	3 KB
output_csv_(GET_STUDIES_OBSERVA...)	Alice Boizet	text/csv	17 Sep 09:41 AM 2019	19,020 KB
LogFile_(GET_STUDIES_OBSERVA...)	Alice Boizet	text/plain	17 Sep 09:41 AM 2019	2 KB
outputJSON_(BRAP1_GET_STUDIES_OBSERVA...)	Alice Boizet	application/json	16 Sep 03:23 PM 2019	1 KB
outputCSV_(BRAP1_GET_STUDIES_OBSERVA...)	Alice Boizet	text/csv	16 Sep 03:23 PM 2019	8,675 KB
LogFile_(BRAP1_GET_STUDIES_OBSERVA...)	Alice Boizet	text/plain	16 Sep 03:22 PM 2019	7 KB
outputFile_(PHI1_GET_STUDIES_OBSERVA...)	Alice Boizet	text/csv	16 Sep 11:03 AM 2019	14,483 KB
LogFile_(PHI1_GET_STUDIES_OBSERVA...)	Alice Boizet	text/plain	16 Sep 11:03 AM 2019	3 KB

Fill the parameters as bellow:

Create Task Configuration for: outputCSV_(BRAPI_...

Execute in the VRE

The Algorithm

Description

Parameter

Key:

Value:

Parameter

Key:

Value:

Parameter

Key:

Value:

Parameter

Key:

Create Task Configuration for: outputCSV_(BRAPI_...

Parameter

Key:

Value:

Parameter

Key:

Value:

Parameter

Key:

Value:

Parameter

Key:

Value:

Add Parameter



Parameters description : If removeOutliers=0 the outliers will be kept and identifiable. If createPDFwithPlots=1 a pdf file with plots of each time serie will be created

Click on *Create configuration*, then run the algorithm.

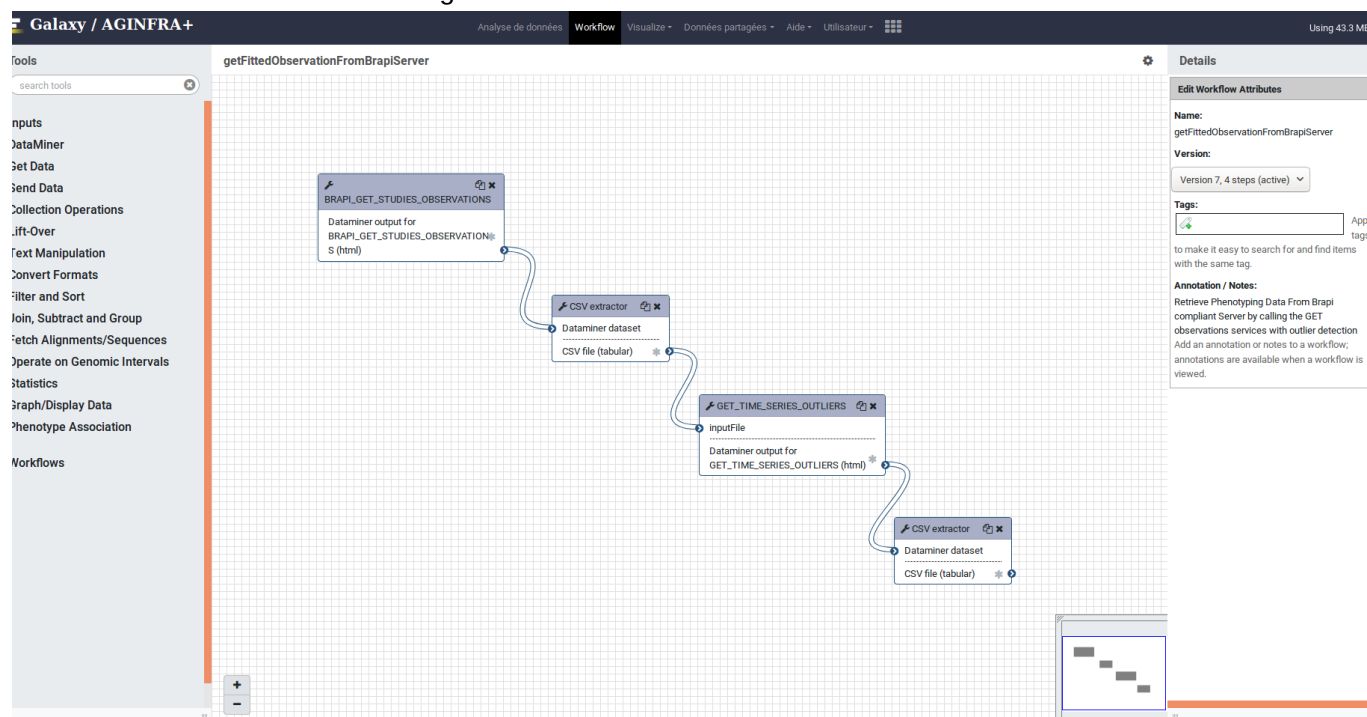
After the execution, you should find the output files in your workspace (*dataminer/output data sets folder*)

Make a Galaxy workflow

You would like to be able to access data from PHIS studies and detect outliers in a single process as a routine, so that you could easily do the same process on next studies. To do that, you can build a workflow in Galaxy. Galaxy can be found in the analytics tab. The dataminer algorithms are automatically transferred as galaxy tools. Build a workflow which first Retrieve observation data of the study ZA12 and then detects outliers.

Tip : you need to use node csv_extractor in order to convert the dataminer tools output into csv

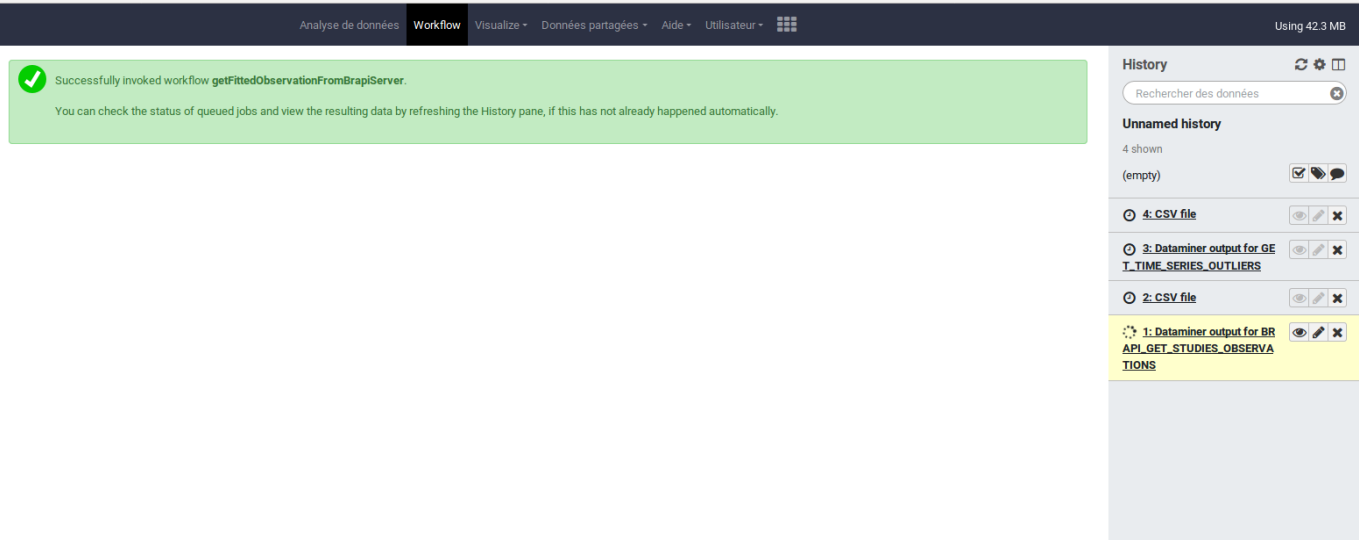
You should have a workflow looking like this :



Don't forget to save your workflow by clicking on



When your workflow is ready, you can run it. You can see each step execution on the right of the screen.



After the workflow execution, you can view the outputs. You can click on the view button on the 3rd step (Dataminer output for get_time_series) when you can click on the 2 output files to download them.

The csv validator tool enable to convert the dataminer csv output into a galaxy datatable file which means that it can be reused as an input on another galaxy tool and you can visualize the data directly in Galaxy. To do that, you can click on the view button of the last step of the workflow.