



SuGR

Welcome to SuGR (SAFIR02 Upload Gustave Roussy), the Shiny application that allows you to check, edit your files and submit them on Synapse.

1- Enter your Synapse User Name and Password

A login form with a light gray background. It contains two input fields: "User Name:" and "Password:". Below the password field is a "Log in" button with a dark gray background and white text.

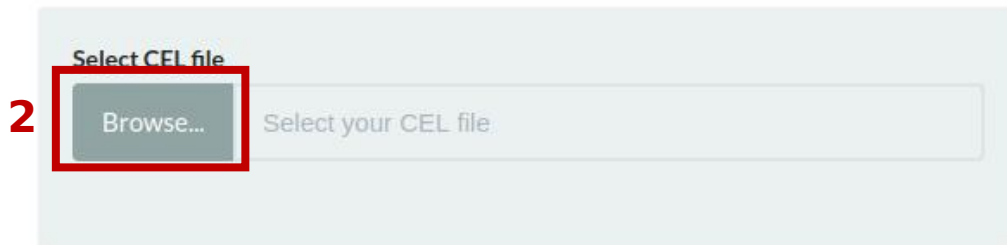
Please contact Alicia TRAN DIEN (alicia.tran-dien@gustaveroussy.fr) from  Bioinformatics platform for any requests.

2- Upload your files

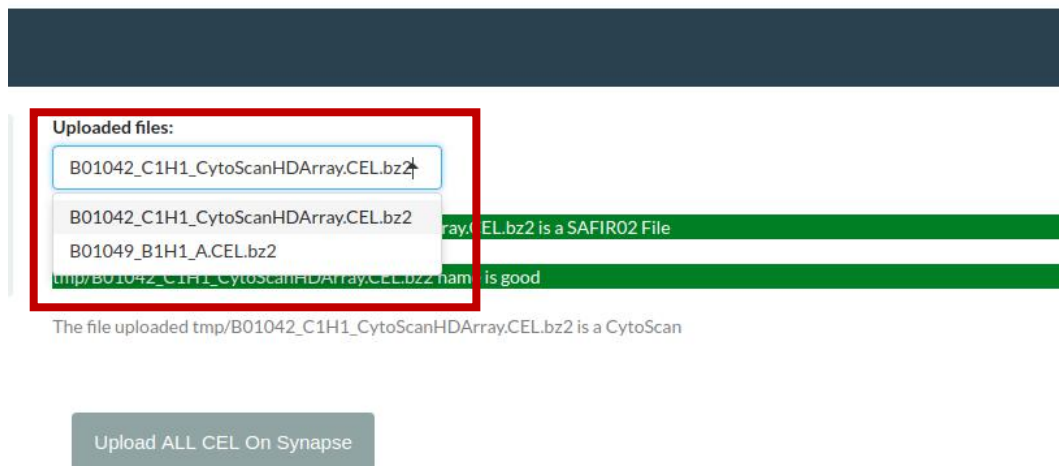
- You can select multiple files at the same time.
- If you don't see your file, it means that the file you are trying to upload was not supported by SuGR. Please ensure that your file has either a .CEL/.CEL.bz2 or a .tsv extension. Otherwise, the warning below will show up.

A red rectangular box with white text that reads: "Oops!! The file format is incorrect please try again".

a- Upload CEL file



- Select your file in the drop down menu



ct Alicia TRAN DIEN (alicia.tran-dien@gustaveroussy.fr) from  Bioinformatics platform for any requests.

- Upload all the files on Synapse

Uploaded files:

B01042_C1H1_CytoScanHDArray.CEL.bz2

B01042_C1H1_CytoScanHDArray.CEL.bz2

B01049_B1H1_A.CEL.bz2

ay.CEL.bz2 is a SAFIR02 File

tmp/B01042_C1H1_CytoScanHDArray.CEL.bz2 name is good

The file uploaded tmp/B01042_C1H1_CytoScanHDArray.CEL.bz2 is a CytoScan

Upload ALL CEL On Synapse

b- Upload NGS file

- You can select multiple files at the same time.

CEL

NGS

1

Choose NGS File

2

Browse...

Select your NGS file

You can download a SampleSheet Example if you want

SampleSheetExample

You can download the list of genes name below

GeneName

- Wrong gene name typo is mark with a red background cell. You can edit the table by double clicking on the cell and type Enter.

Don't forget to click on Save changes after editing.

Warning : Clicking again on the file in the drop down menu will erase your modifications.

Upload ALL NGS On Synapse

Save changes

| | PatId | Global_Conclusion | Manual_Var_Comment | Manual_Var_Classif | Gene_Symbol | Protein_Char |
|----|--------|------------------------------|--------------------|--------------------|-------------|--------------|
| 1 | T04004 | Presence d'une mutation TP53 | Variant deletere | VP | TP53 | p.C141Y |
| 2 | T04004 | | | VNP | Brca2 | p.R324T |
| 3 | T04004 | | | 0 | Brca2 | p.V2466A |
| 4 | T04004 | | | 0 | NOTCH4 | p.G534S |
| 5 | T04004 | | | 0 | TSC1 | p.M322T |
| 6 | T04004 | | | 0 | FGFR4 | p.V10I |
| 7 | T04004 | | | 0 | MAP3K1 | p.V906I |
| 8 | T04004 | | | 0 | FGFR4 | p.P136L |
| 9 | T04004 | | | 0 | NOTCH4 | p.D272G |
| 10 | T04004 | | | 0 | MAP3K1 | p.D806N |
| 11 | T04004 | | | 0 | TP53 | p.P72R |
| 12 | T04004 | | | 0 | NOTCH4 | p.K117Q |
| 13 | T04004 | | | 0 | NOTCH4 | p.T320A |

- On the side panel, you can see the list of files that will be uploaded on Synapse

List of files that will be uploaded on Synapse

| | inputfile | sy |
|---|--|-----|
| 1 | data/tmpData/AliciaTD/B01041_C1H1_CytoScanHD_Array.CEL.bz2 | syr |
| 2 | data/tmpData/AliciaTD/B01042_C1H1_CytoScanHD_Array_1_CEL.bz2 | syr |

Purge list to upload

Upload ALL NGS On Synapse

Save changes

| | PatId | Global_Conclusion | Manual_Var_Comment | Manual_Var_Classif | Gene_Symbol | Protein_Char |
|----|--------|------------------------------|--------------------|--------------------|-------------|--------------|
| 1 | T04004 | Presence d'une mutation TP53 | Variant deletere | VP | TP53 | p.C141Y |
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| 5 | T04004 | | | 0 | TSC1 | p.M322T |
| 6 | T04004 | | | 0 | FGFR4 | p.V10I |
| 7 | T04004 | | | 0 | MAP3K1 | p.V906I |
| 8 | T04004 | | | 0 | FGFR4 | p.P136L |
| 9 | T04004 | | | 0 | NOTCH4 | p.D272G |
| 10 | T04004 | | | 0 | MAP3K1 | p.D806N |
| 11 | T04004 | | | 0 | TP53 | p.P72R |
| 12 | T04004 | | | 0 | NOTCH4 | p.K117Q |
| 13 | T04004 | | | 0 | NOTCH4 | p.T320A |

- You can purge the list of files to be upload on Synapse by clicking on the button below.

List of files that will be uploaded on Synapse

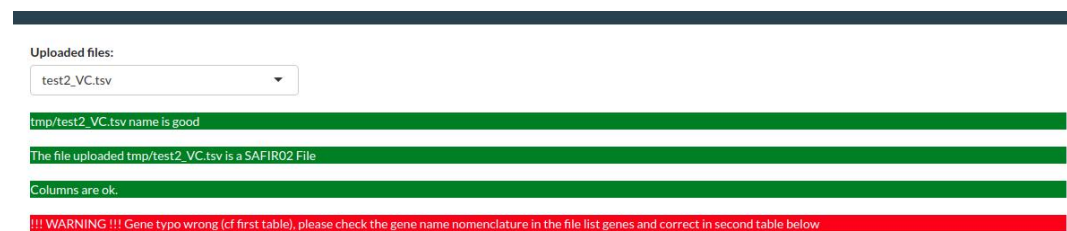
| | inputfile | sy |
|---|---|-----|
| 1 | data/tmpData/AliciaTD/B01041_C1H1_CytoScanHD_Array.CEL.bz2 | syn |
| 2 | data/tmpData/AliciaTD/B01042_C1H1_CytoScanHD_Array_1_.CEL.bz2 | syn |

Purge list to upload

3- Warning Color

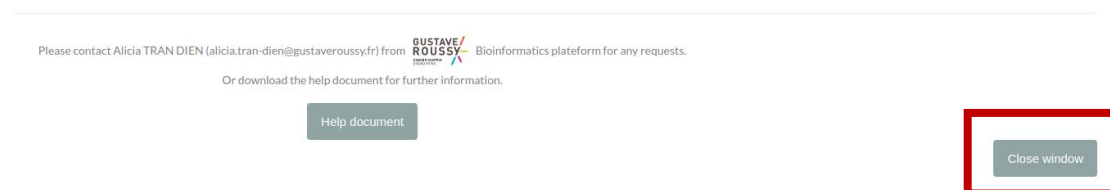
- A green warning indicates that the file is correct and can be uploaded
- An orange warning indicates that the file is incorrect but the problem has been corrected (ex: space found in file name) and the file ready to be uploaded. **Note that once the file has been rename, you can not click again on the drop down menu. You will have to upload it again**
- A red warning indicates that the file is incorrect and that a new version has to be edited and/or download.

Make sure that there are no red warnings before uploading on synapse.



4- Disconnecting

You can disconnect from Synapse and close the app/window by clicking on the button below.



5- Contact

If you need any help or requests, please contact Alicia TRAN DIEN (alicia.tran-dien@gustaveroussy.fr) from Gustave Roussy Bioinformatics platform.