



# Data Challenge Presentation

Teams: Fork Gladius + Issue Saber

## Fork Gladius

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## PHYSICS CASE



Picture of AGATA detector taken in GANIL

**$^{98}\text{Zr}$**   
**30.7 s**

**$\beta^- = 100\%$**

**$^{90}\text{Br}$**   
**1.91 s**

**$\beta^- = 100\%$**   
 **$\beta^- n = 25.3\%$**

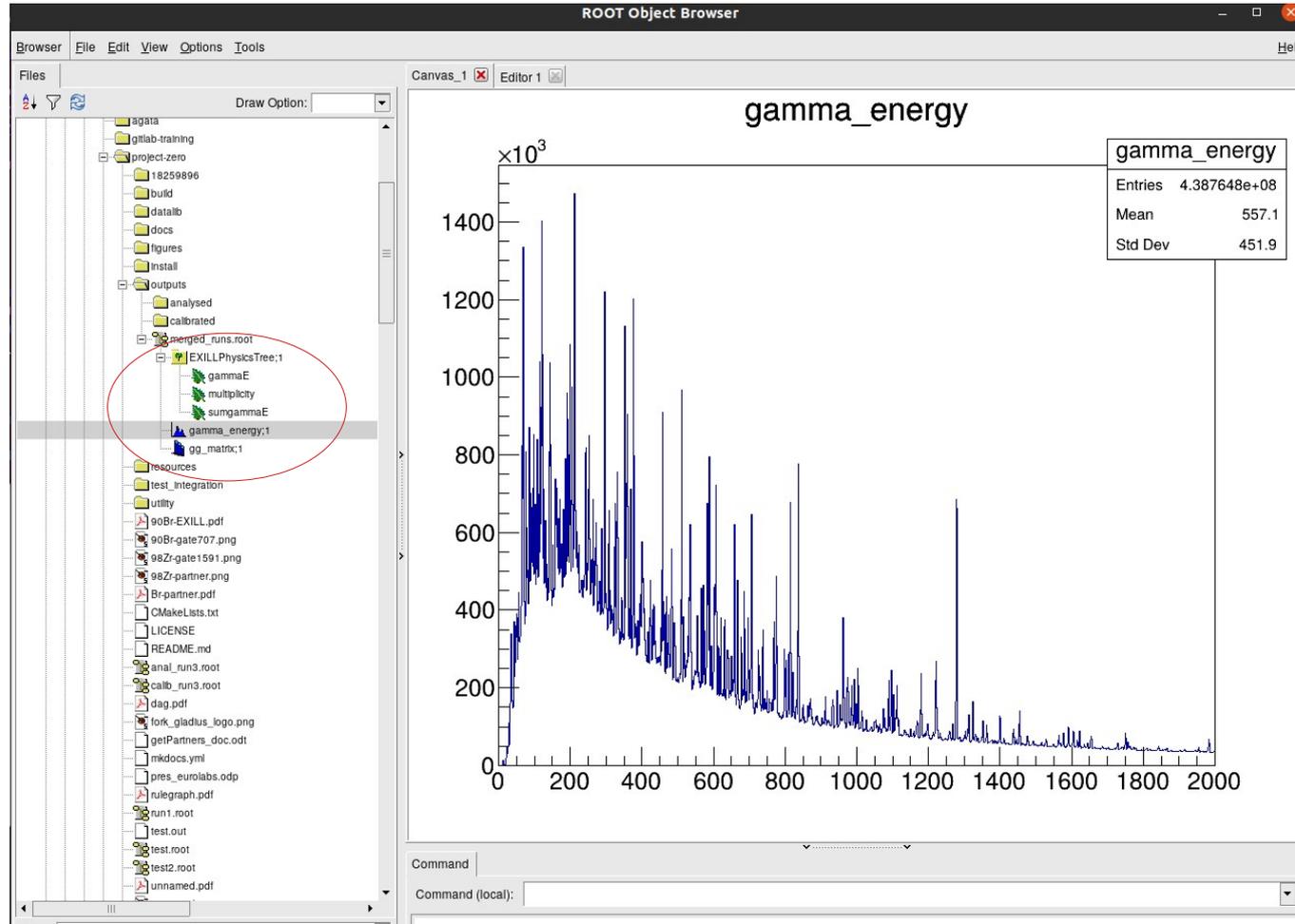
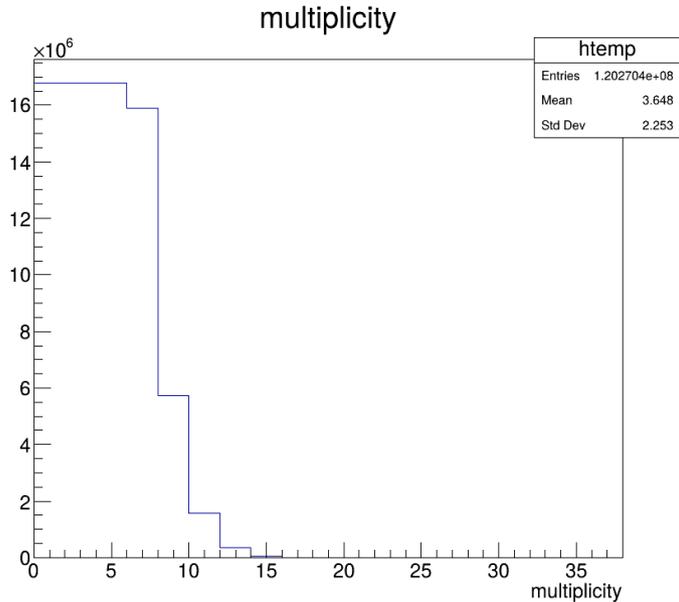
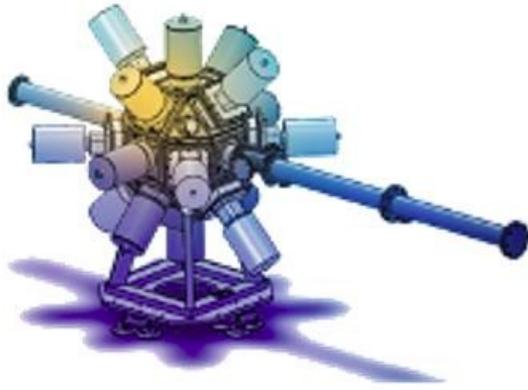
Setup : VAMOS + AGATA

Gamma rays spectroscopy

Gammas emitted in flight (Doppler correction needed)

=> Only single gamma spectrum

# EXILL: Data



# GitLab Page

<https://project-zero-f040ed.pages.in2p3.fr/>



Activities Firefox Web Browser 08:25 en

98zr\_it\_decay\_1.9\_un X NNDC | National Nuc X NuDat 3 X List of levels for 98zr X Datasets for 98zr X EnsdfDispatcherServ X Inbox (34) - konstpall X Project Zero X

Import bookmarks... Getting Started PyPI - The Python Pac... Marshmallow - YarnArt Report\_tracker\_test... Clebsch-Gordan calcul... Barad-dur - Free Pat... Everything Free Cooki... Πρόσδιο πιότο | Ακηç... LRS 821.pdf

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## Fork-Gladius (project-zero fork)

This project contains the fork of the project-zero made by the amazing fork-gladius team.

### Setting up the work environment

#### Micromamba environment

You first need to install micromamba:

```
`${SHELL}` <(curl -L micro.mamba.pm/install.sh)
```

You can then create the root\_py12 environment:

```
micromamba env create -f resources/root_py12.yaml
```

And activate it:

```
micromamba activate root_py12
```

#### Compiling the code

It is recommended to create a build and install folder:

```
mkdir build install && cd build
```

As the micromamba environment includes ninja, it is recommended to use it to compile the code. ninja automatically allows the RAM in order to optimize your compilation time. In the build folder:

```
cmake -GNinja -DCMAKE_INSTALL_PREFIX=../install ..
```

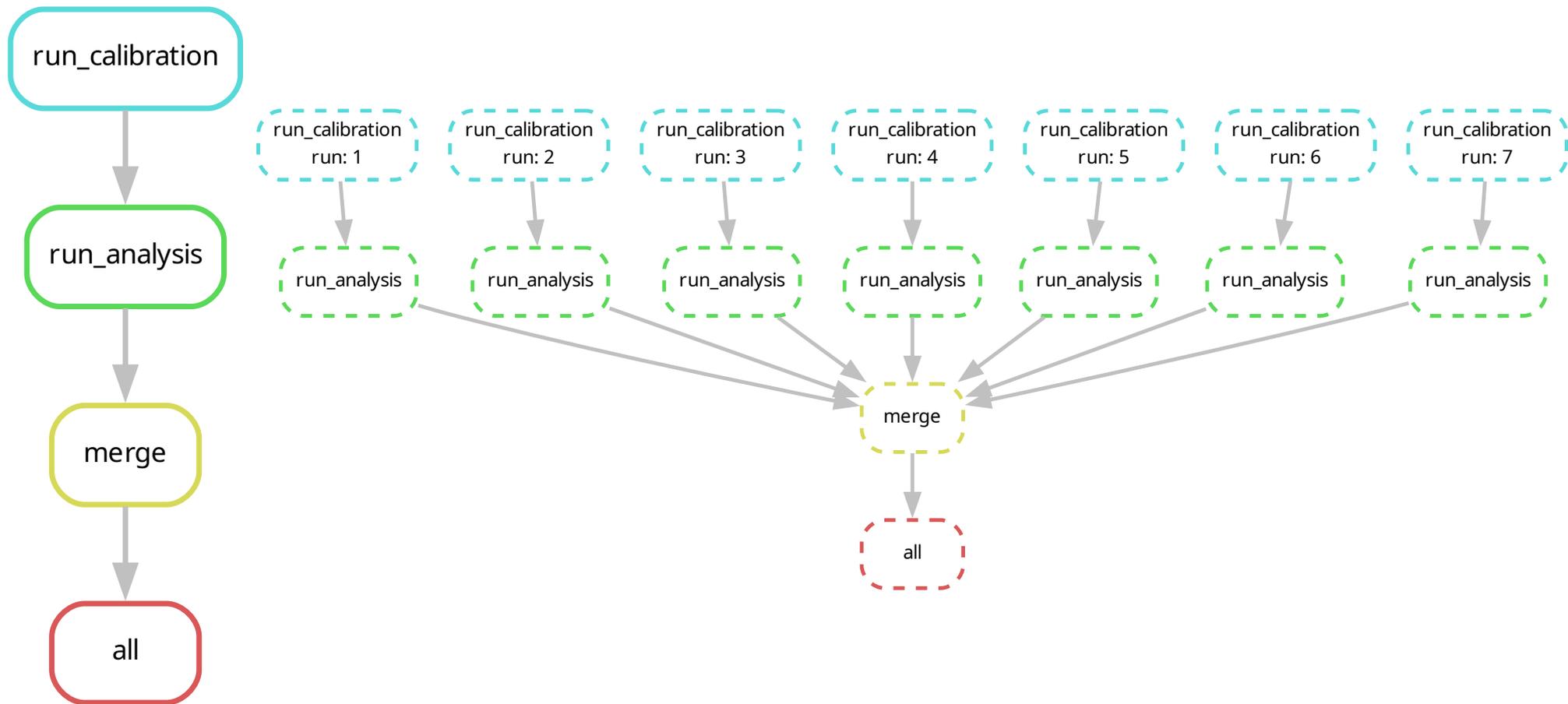
You can then install it:

```
ninja install
```

You can now use the executable in install/bin

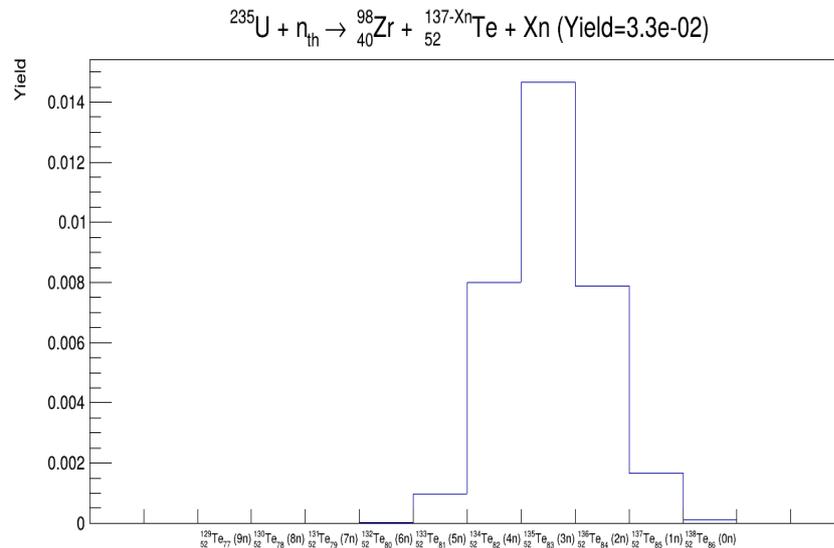
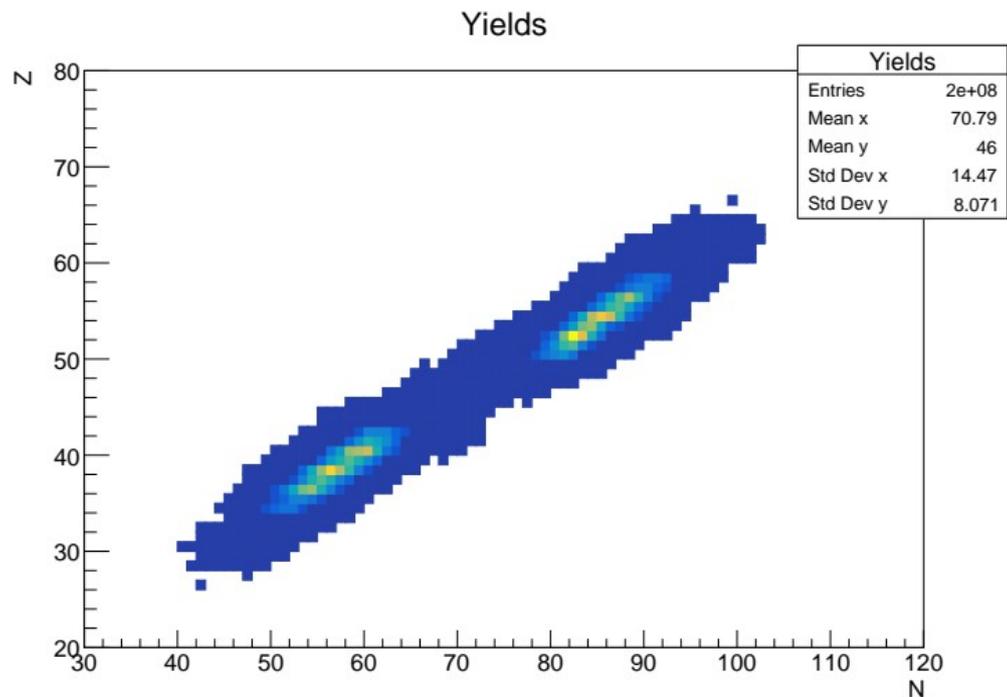
### Utility codes

# Snakemake structure

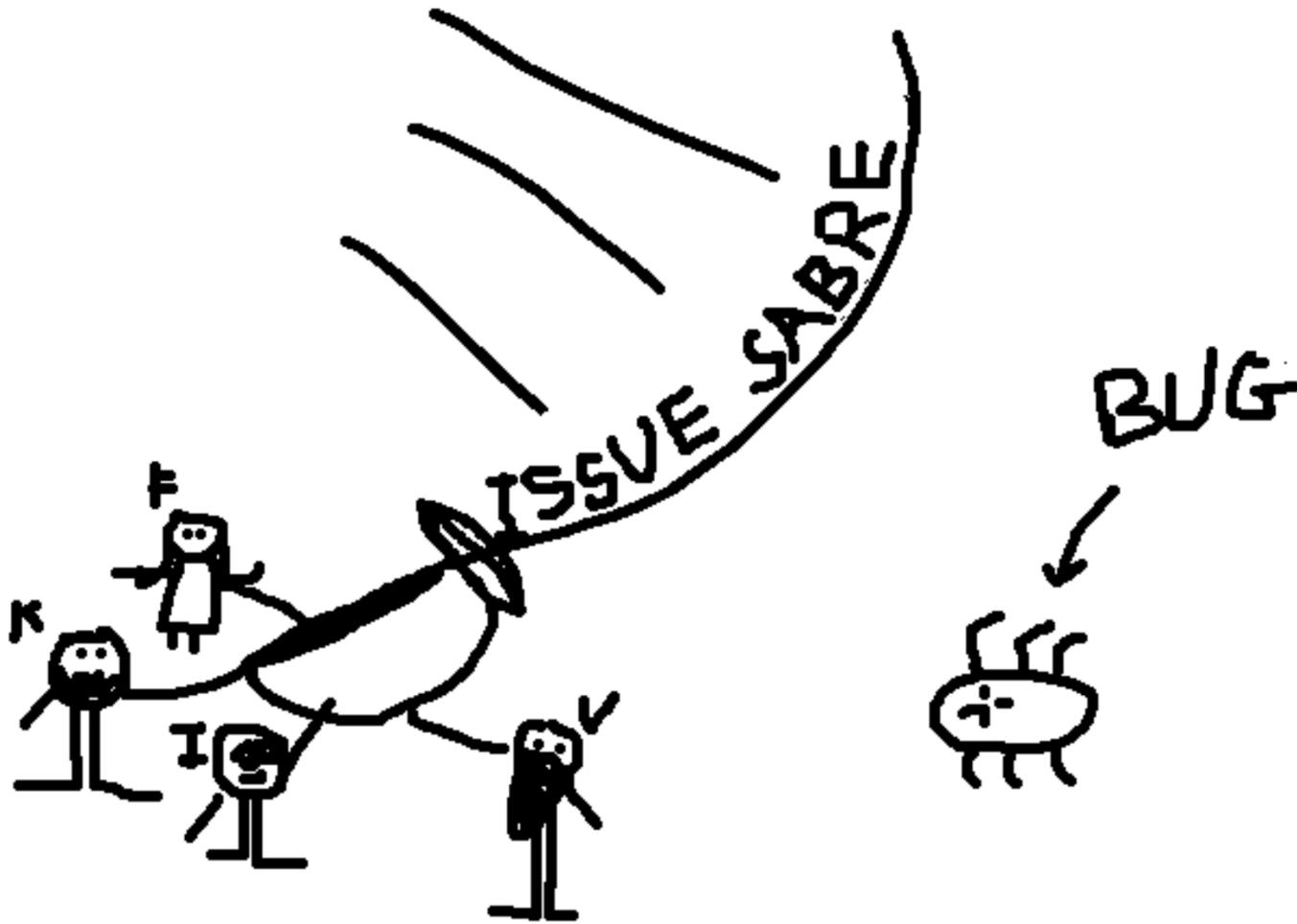


# get\_partner Code

EXILI\_Partners.root



# Issue-saber

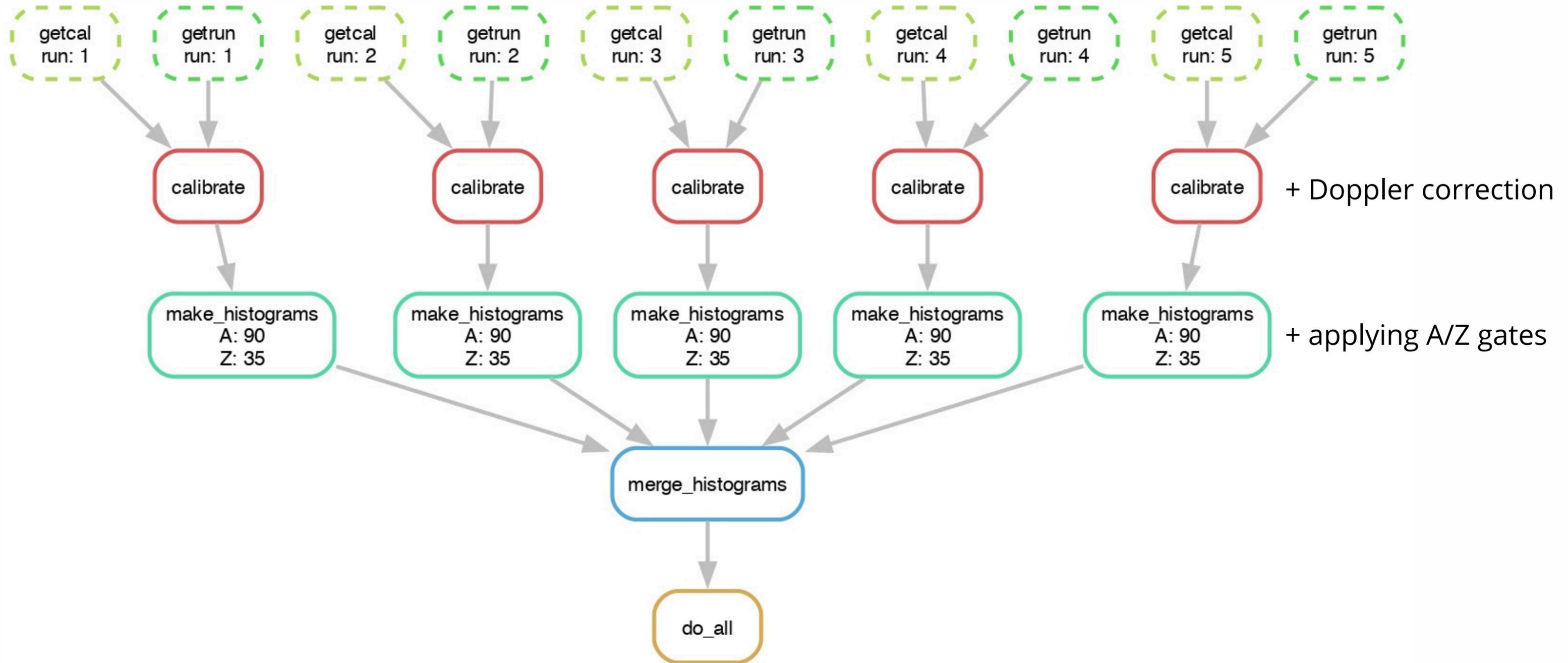


main agata Find file Code

Merge branch 'main' of gitlab.in2p3.fr:eurolabs-os-school/hands-on-2026/issue-saber/agata ✓ c1260a55 History  
VICTOR MARTINEZ NOUVILAS authored 6 minutes ago

Name	Last commit	Last update
data/lib	Resolve "Check the documentation is up ..."	2 days ago
env	Add root_py12	3 days ago
images	Added spectra images for 98Zr and 1221 ...	7 minutes ago
resources	adding images	19 minutes ago
test	adding reproducible file to resources	2 days ago
utility	changes in Snakefile (example path), RE...	16 hours ago
.gitlab-ci.yml	Update .gitlab-ci.yml file	2 days ago
CMakeLists.txt	Resolve "Create a test programm that co..."	2 days ago
LICENSE	Resolve "Update README so the EXILL g..."	17 hours ago
README.md	changes to README	15 hours ago
Snakefile	Resolve "Update README so the EXILL g..."	17 hours ago
Snakefile_config.yaml	changes in Snakefile (example path), RE...	16 hours ago
Spectrum90Br.png	Upload New File	23 minutes ago
dag.pdf	Upload New File	8 minutes ago
outputfiles_snakemake.png	Output_files	25 minutes ago
README.md		

# ANALYSIS WORKFLOW



# CODES AND ISSUES

```
agata > ! Snakefile
1 wildcard_constraints:
2   run = r"d{1,2}"
3
4 configfile: # Default configuration file
5   f"{workflow.basedir}/Snakefile_config.yaml"
6
7 def get_data_location():
8   return config['dataLocation']
9
10 def get_runs():
11   return [i for i in range(config['run_min'], config['run_max'] + 1)]
12
13 def get_gates():
14   gate_str_list = []
15   for gate in config["gates"]:
16     name = str(gate["gateA"]) + "_" + str(gate["gateZ"])
17     gate_str_list.append(name)
18   return gate_str_list
19
20 rule do_all:
21   input:
22     expand("histograms/merged_histogram_{gate}.root", gate=get_gates() )
23
24 rule getcal:
25   output:
26     f"{config['dataLocation']}/cal_run{{run}}.yaml"
27   shell:
28     "curl https://zenodo.org/records/18253801/files/cal_run{wildcards.run}.yaml > {output} "
29
30 rule getrun:
31   output:
32     f"{config['dataLocation']}/run{{run}}.root"
33   shell:
34     "curl https://zenodo.org/records/18253801/files/run{wildcards.run}.root > {output} "
35
36 rule calibrate:
37   input:
38     cal = str(expand("{dataLocation}", dataLocation = get_data_location())[0]) + "/cal_run{run}.yaml",
39     data = str(expand("{dataLocation}", dataLocation = get_data_location())[0]) + "/run{run}.root"
40   output:
41     "outfiles/run{run}-cal.root"
42   shell:
43     f"{workflow.basedir}/" + "install/bin/analysis_agata --in {input.data} --out {output} --cal {input.cal}"
44
45 rule make_histograms:
46   input:
47     "outfiles/run{run}-cal.root"
48   output:
49     "histograms/hist_run{run}_{A}_{Z}.root"
50   shell:
51     f"{workflow.basedir}/" + "install/bin/histo_code.py --in {input} --tree-name AVPhysicsTree --out {output} --gateA {wildcards.A} --gateZ {wildcards.Z}"
52
53 rule merge_histograms:
54   input:
55     expand("histograms/hist_run{run}_{A}_{Z}.root", run = get_runs())
56   output:
57     "histograms/merged_histogram_{A}_{Z}.root"
58   shell:
59     "hadd -f {output} {input}"
60
```

Snakemake structure

```
agata > ! Snakefile_config.yaml
1 # Default configuration
2 # - note that yaml allow comments, so put some if needed
3 run_min: 1
4 run_max: 21
5 gates:
6   - gate:
7     gateA: 98
8     gateZ: 40
9   - gate:
10    gateA: 90
11    gateZ: 35
12
13 dataLocation: "/home/zeghni-l/Documents/18253801"
14
```

Configuration for snakemake

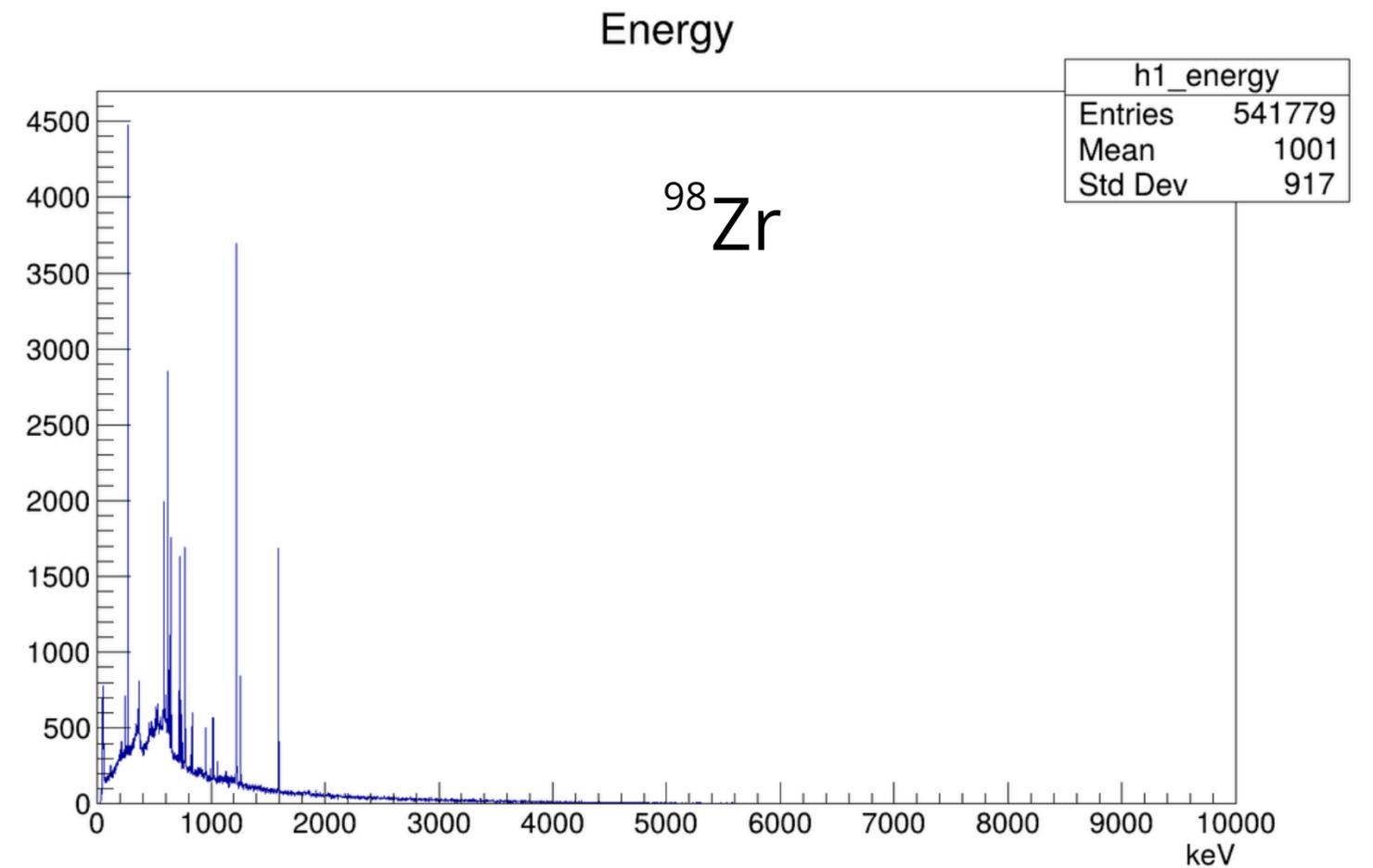
Some issues encountered:

- Merging conflicts
- Path related issues
- README file
- MetaData missing

# RESULTS

```
agata > utility > histo_code.py > main
1  #!/usr/bin/env python3
2  """
3  Generate histograms from ROOT TTree, using pyROOT
4
5  Usage:
6  generate_histo_pyroot.py --in INPUT_FILE --out OUTPUT_FILE [--tree-name NAME] --gateA GATEA --gateZ GATEZ
7  generate_histo_pyroot.py -h | --help
8
9  Options:
10 -h --help          Show this screen.
11 --in INPUT_FILE    ROOT input file
12 --tree-name NAME   TTree name within the ROOT file [default: events]
13 --out OUTPUT_FILE  ROOT output file
14 --gateA GATEA      Gate on A
15 --gateZ GATEZ      Gate on Z
16 """
17
18 from docopt import docopt
19 import ROOT
20
21 def main(input_file_name, input_tree_name, output_file_name, gateA, gateZ):
22
23     fin = ROOT.TFile(input_file_name, "READ")
24     tree = fin.Get(input_tree_name)
25
26
27     # Declare the histogram
28     h1 = ROOT.TH1D("h1_energy", "Energy;keV", 10000, 0, 10000)
29
30     # Fill the histograms
31     # - "E >> h1_energy" -> Use branch E to fill h1 energy
32     # - "ct > 0" -> fill for rows that meet ct > 0
33     # - "goff" -> Don't show anything
34     condition = "A > " + str(int(gateA) - 0.2) + " && A < " + str(int(gateA) + 0.2) + " && Z > " + str(int(gateZ) - 0.2) + " && Z < " + str(int(gateZ) + 0.2)
35     tree.Draw("gammaEDC >> h1_energy", condition, "goff")
36
37     fout = ROOT.TFile(output_file_name, "RECREATE")
38     h1.Write()
39     fout.Close()
40
41
42 if __name__ == "__main__":
43     args = docopt(__doc__)
44
45     main(
46         input_file_name = args['--in'],
47         input_tree_name = args['--tree-name'],
48         output_file_name = args['--out'],
49         gateA = args['--gateA'],
50         gateZ = args['--gateZ']
51     )
```

Python code to obtain histograms  
Sets up the gates for A and Z



Gamma spectrum

# AGATA Data

$^{98}\text{Zr}$  Peak Energies:

268 keV

620 keV

1223 keV

1591 keV

$^{90}\text{Br}$  Peak

Energies:

123 keV

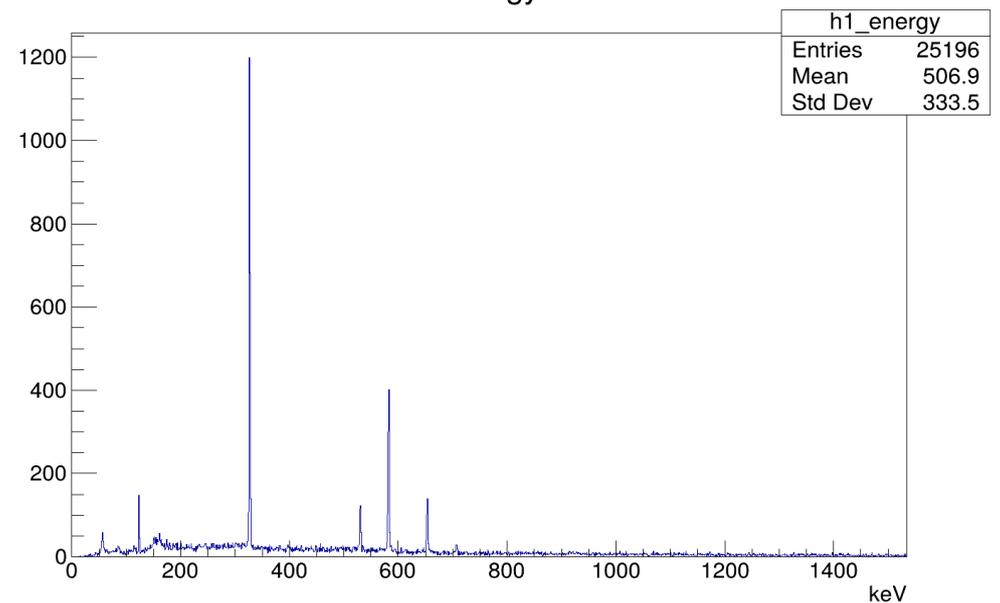
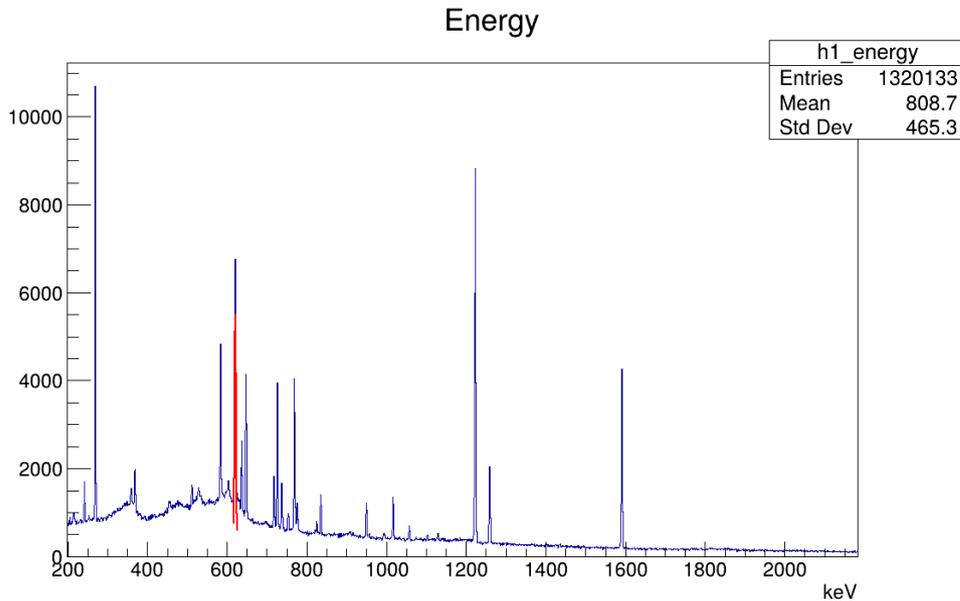
327 keV

530 keV

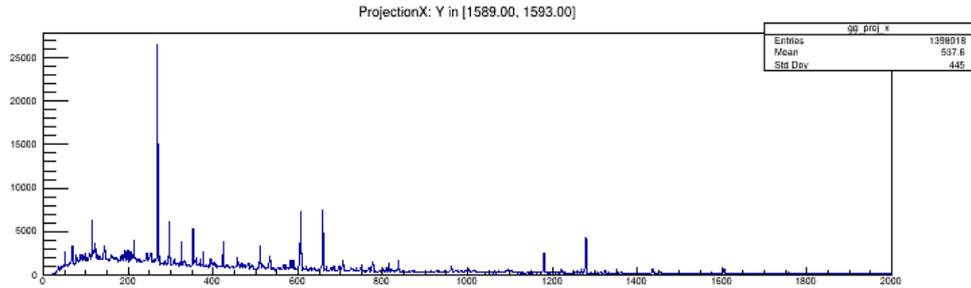
582 keV

653 keV

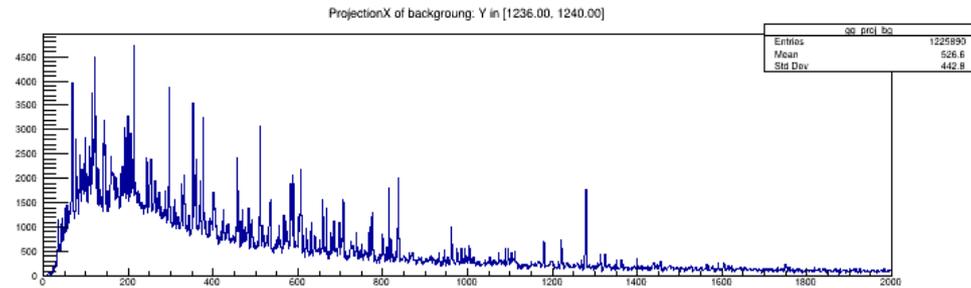
Energy



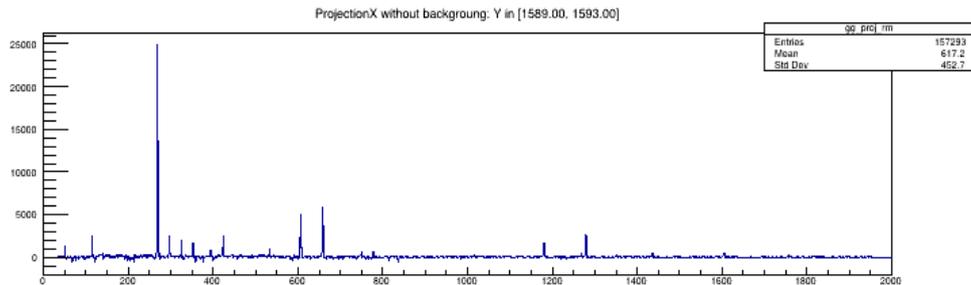
# EXILL: $^{98}\text{Zr}$ $\gamma$ spectrum in coincidence with the $\gamma$ -ray at 1591 keV



- Full spectrum

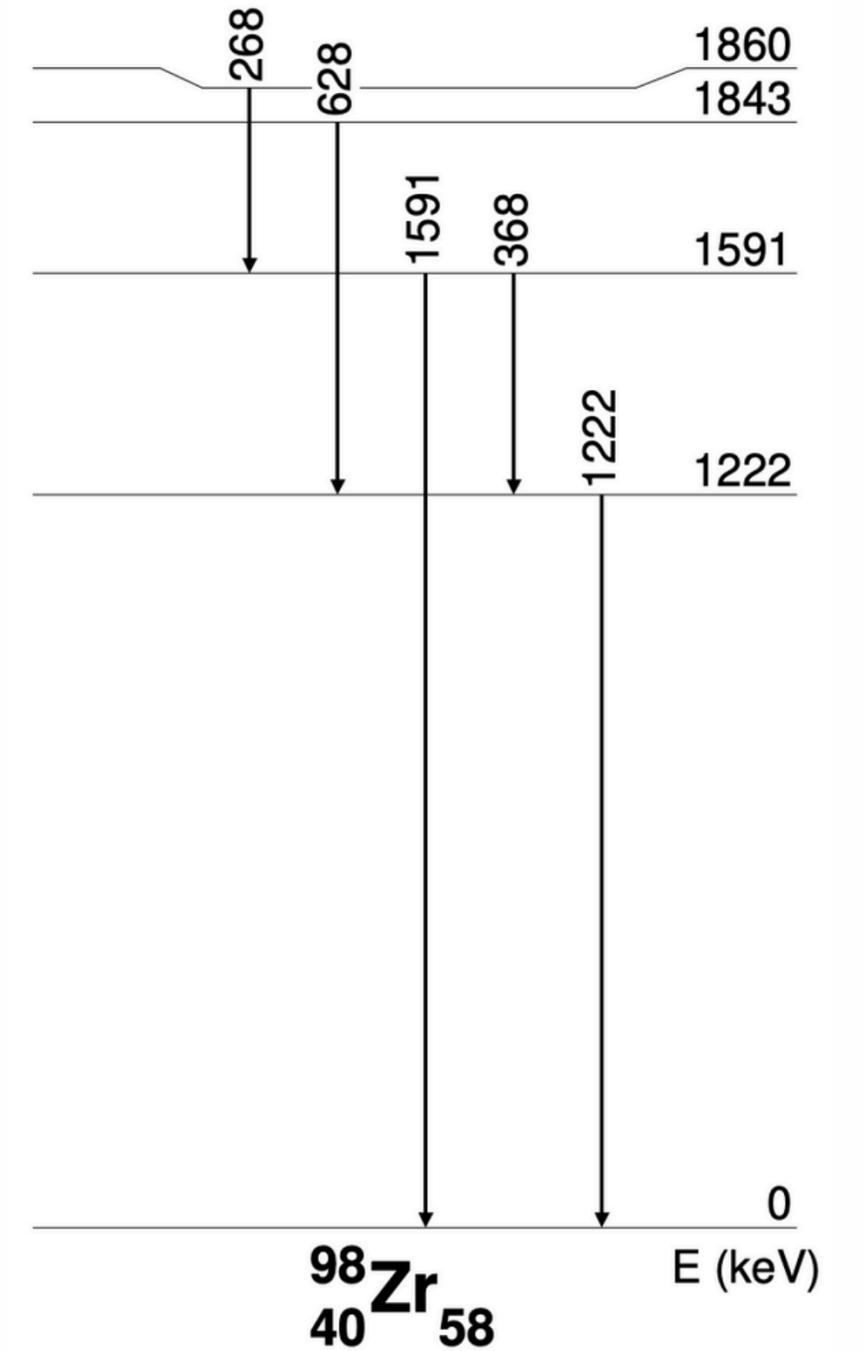
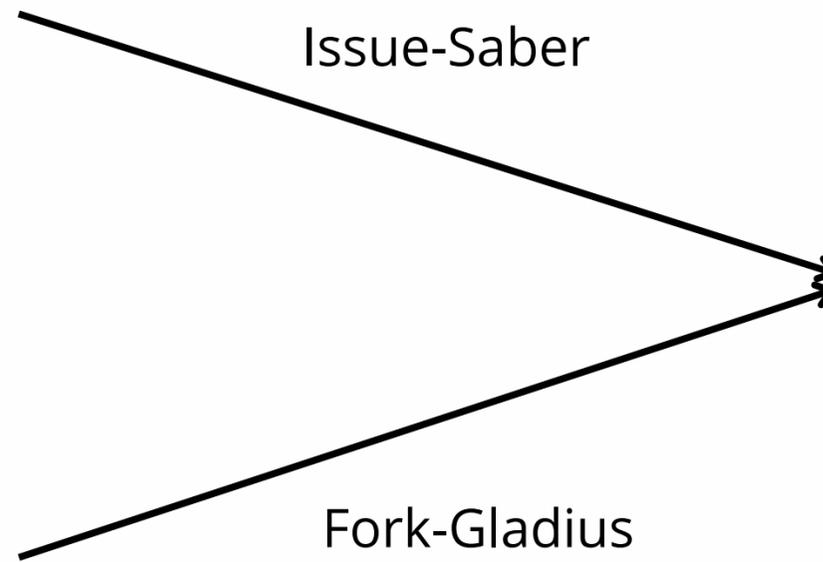
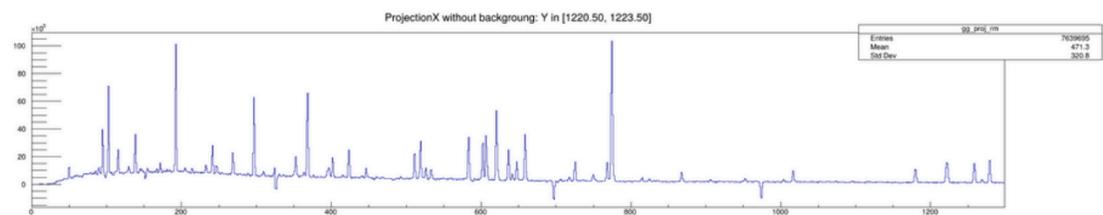
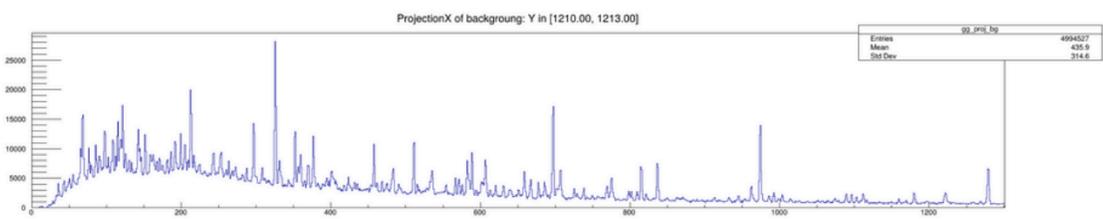
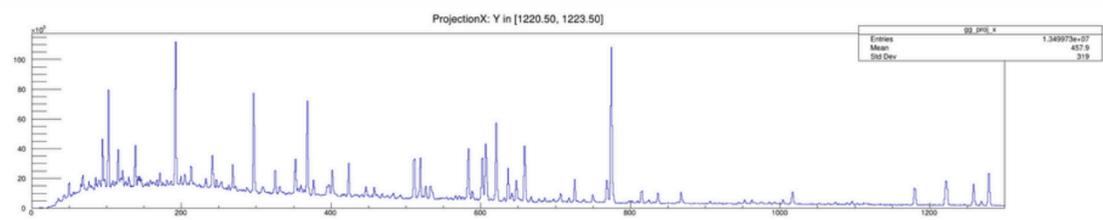
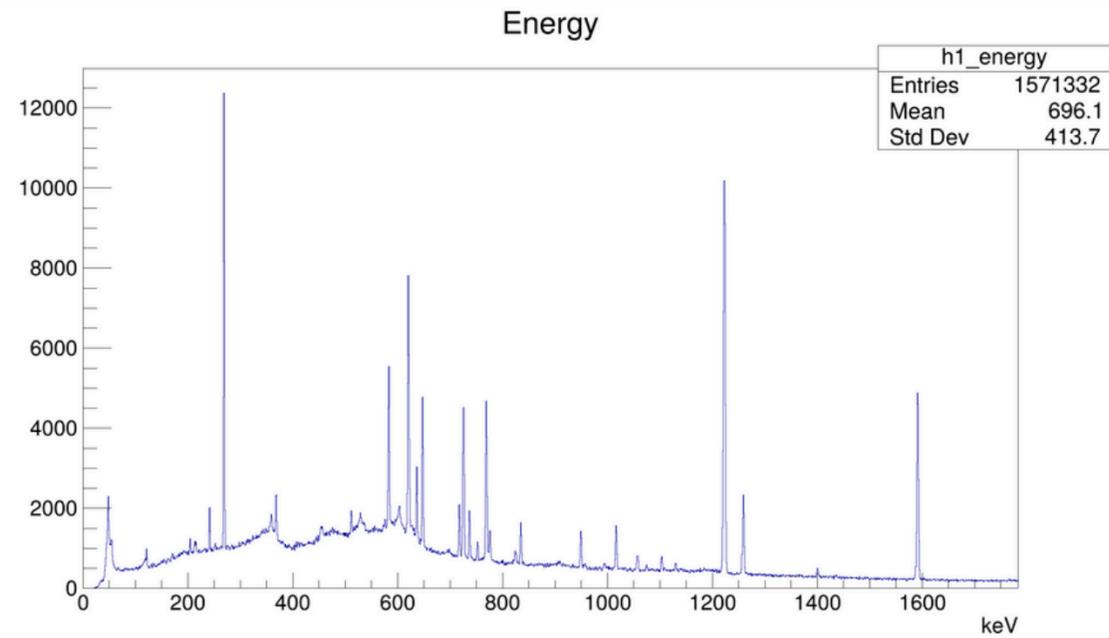


- Background



- Background-subtracted spectrum

# FINAL RESULTS



Level scheme