



Ancillary analysis of AGATA@LNL



D. Brugnara

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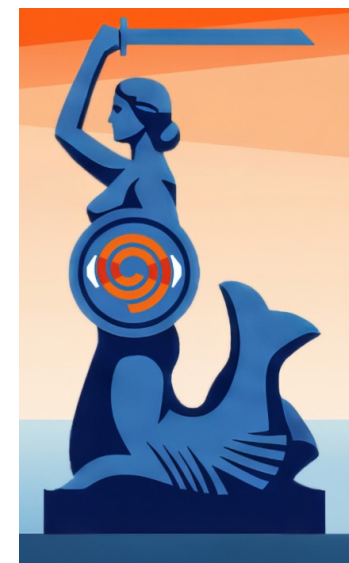
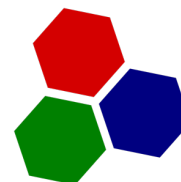
L. Zago

M. Balogh

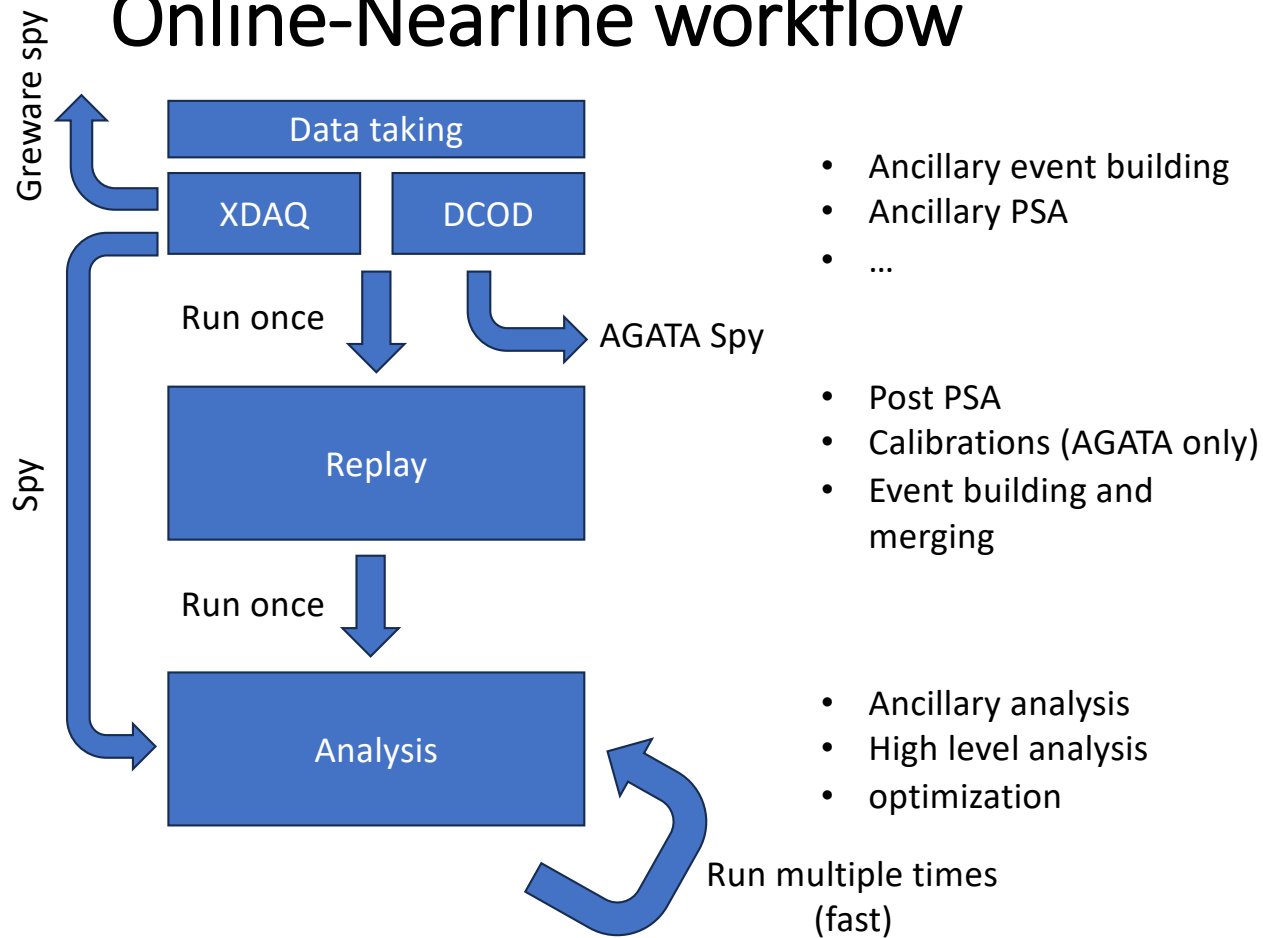
S. Pigliapoco

L. Busak

and the **local LNL group**



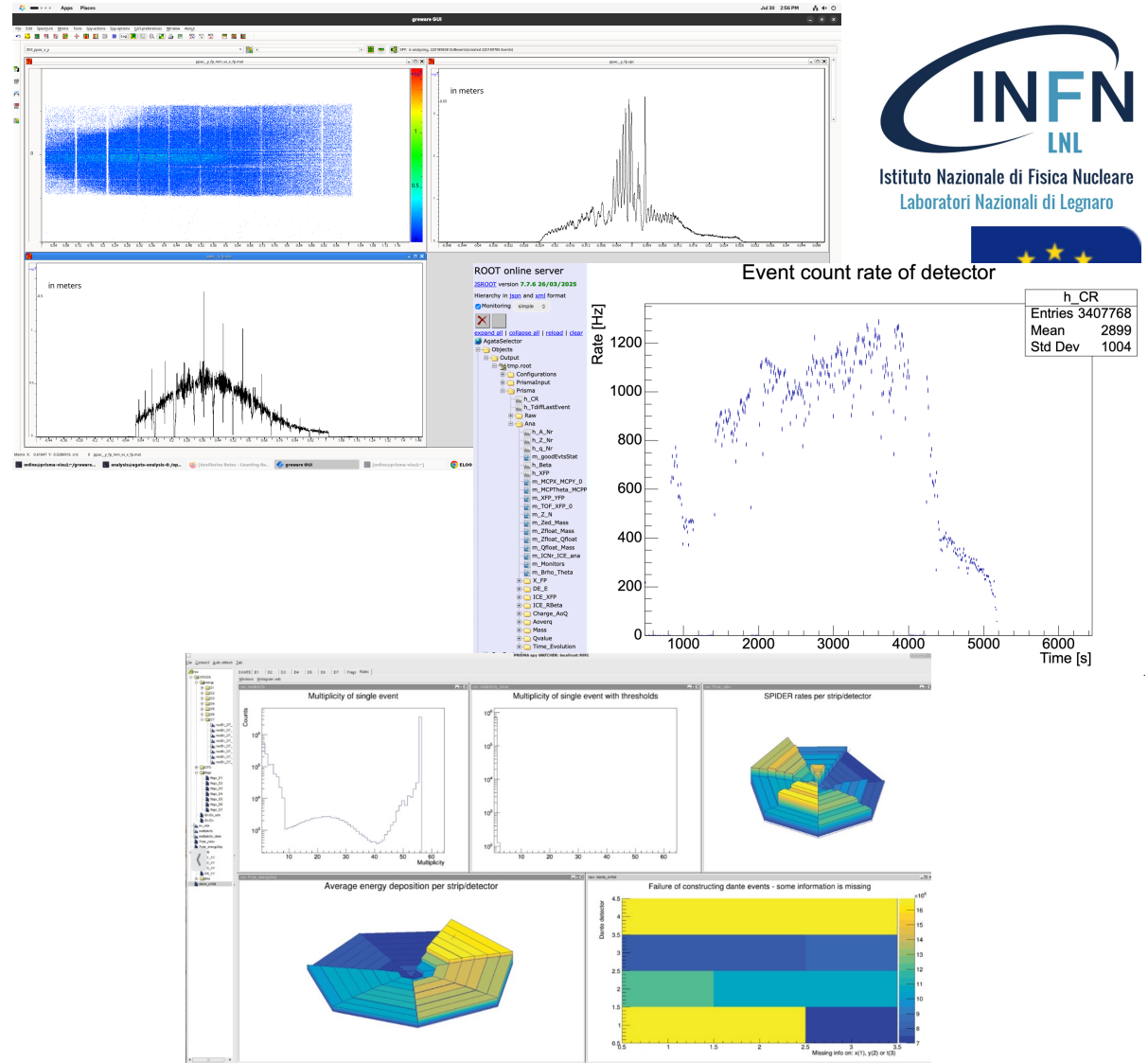
Online-Nearline workflow



- AGATA pre-processing
- AGATA PSA
- ...

Ancillary SPY

- GREWARE
 - Stable and fast
 - Well known by users and maintained for PRISMA
 - It currently does not run the same analysis
- Agataselector (**new development!**):
 - Currently spies only PRISMA data
 - Runs the same analysis as the nearline and offline
 - JSRoot is convenient but sometimes slow
- LNLspy (developed by M. Balogh):
 - ROOT-based ser friendly interface
 - Could be integrated with the selector

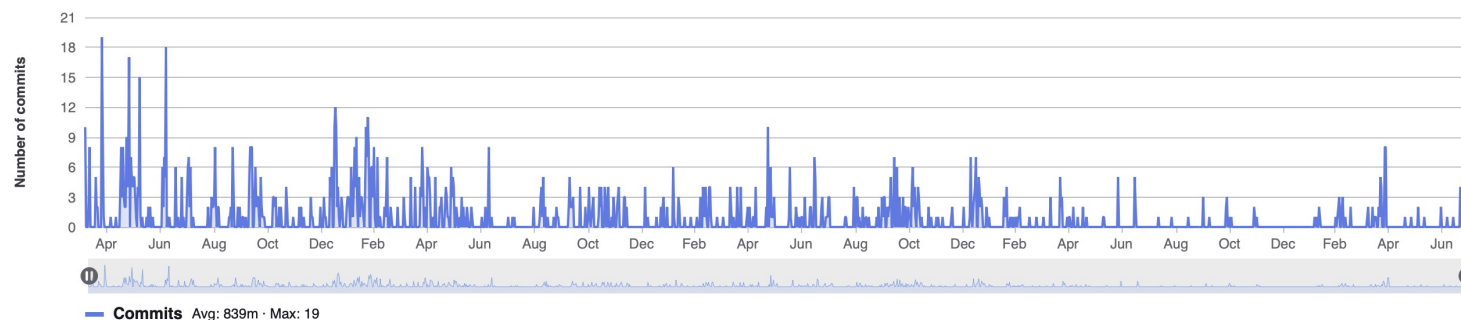


Overview

- Local team effort to provide the users with a comprehensive that handles **nearline and offline analysis**
- **Unique codebase, debugged and cross checked by users and local team**
- Modular and configurable
- **13 detectors present in the analysis**
- 59 experiments up to date and 3 commissioning analyzed
- Well maintained codebase

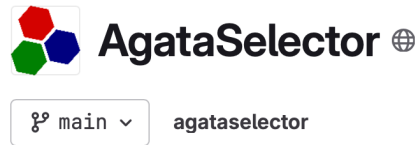
Commits to `main`

Excluding merge commits. Limited to 6,000 commits.



- Total: **1321 commits**
- Average per day: **0.8 commits**
- Authors: **45**

The repository



- Has a DOI:

DOI 10.5281/zenodo.8329198

May 07, 2025



Update file PRISMA_README.md
Filippo Angelini authored 4 months ago

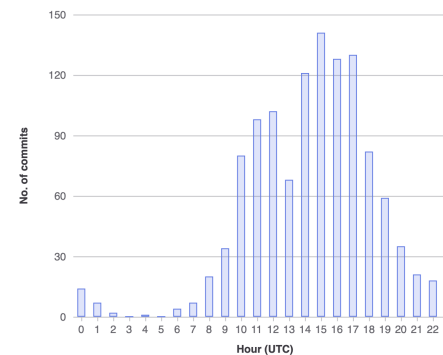


42eb99ee



- Contains tests for:
 - Check for compilation
 - Check for data integrity:
 - Doppler correction quality (mean and sigma of commissioning 1 data)
 - Data loss

Commits per day hour (UTC)



Project information

Data analysis code for the AGATA+ancillary data based on the ROOT framework.

data analysis Physics



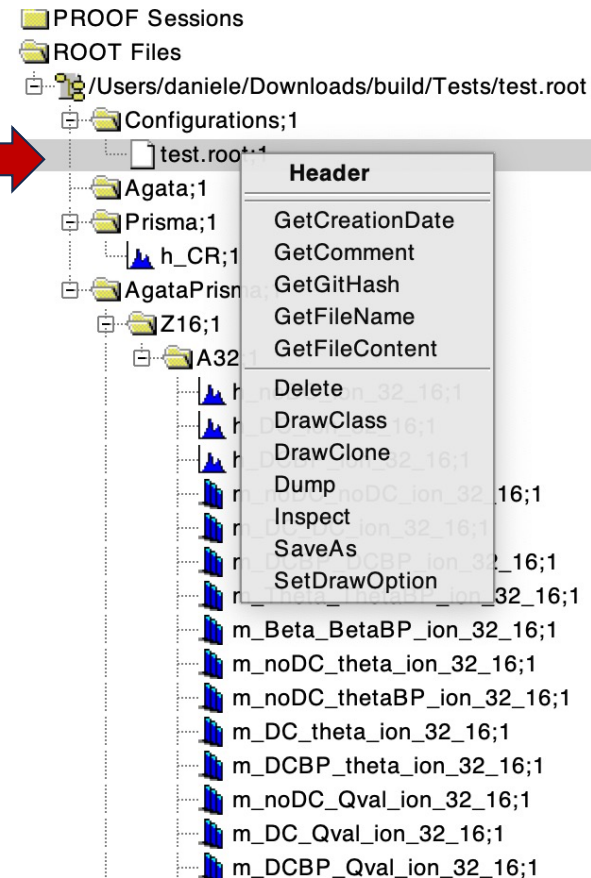
1,558 Commits

7 Branches

6 Tags

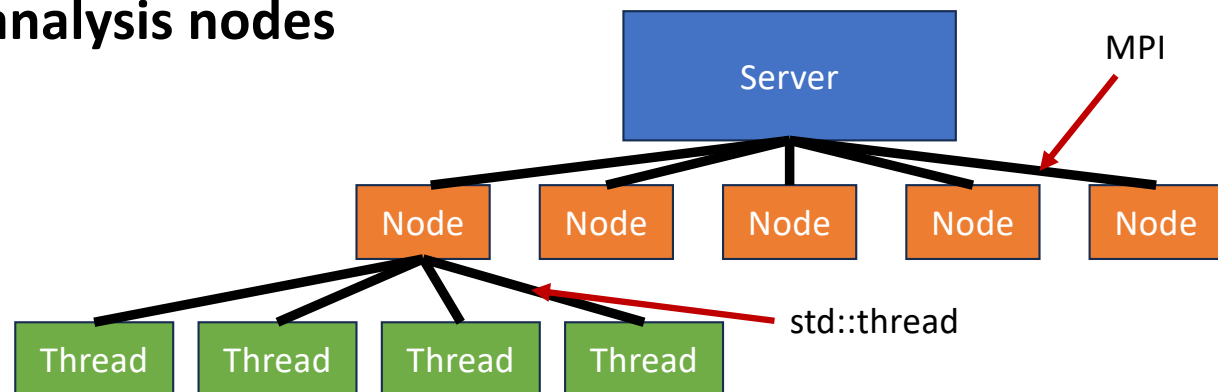
Reproducibility

- The output files contain the parameters used to generate it:
 - The entire selector.conf
 - The lookup tables
 - The git hash
 - The date of creation
- This means that the analysis can be reproduced simply by printing the selector.conf used for this specific file and checking out the correct hash
- The nearline analysis and configurations of each experiment can be recalled by compiling it with the right flags, so the user can start the analysis from day 1.



Distributed analysis

- MPI-based parallelization on multiple machines (boost-mpi)
- Needs to be compiled with “`cmake -DCMAKE_CXX_COMPILER=mpic++ -DUSE_MPI=On .`”
- Very simple server/worker structure
- Allows to run the full analysis multiple times during the nearline **thanks to the AGATA analysis nodes**



Documentation

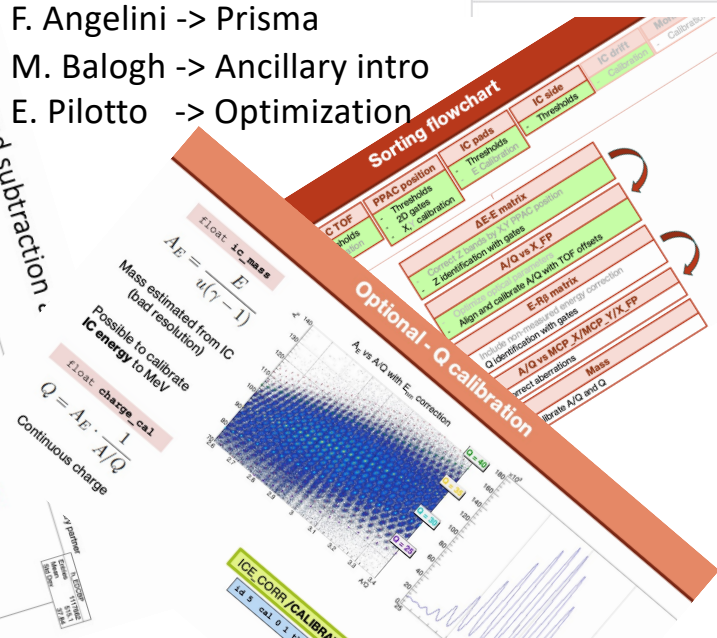
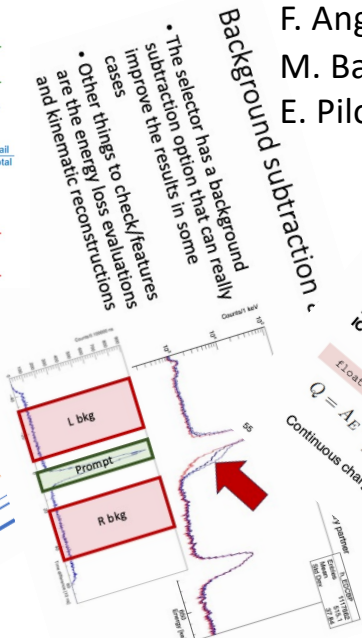
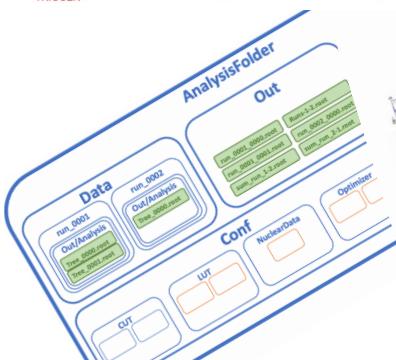
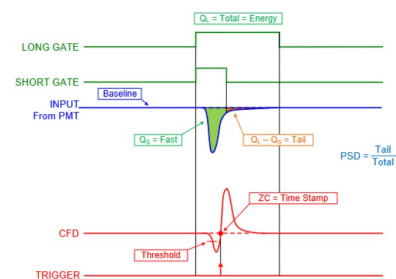
Approx. 300 slides and several **README.md** with the introduction and the details of the ancillary analysis from the latest AGATA Analysis workshop 2025 (thanks to Jérémie). This complements the detailed AGATA LLP+GLP guides.

Thanks to:

F. Angelini -> Prisma

M. Balogh -> Ancillary intro

E. Pilotto -> Optimization



ROOT file processing

A list of ALL existing selector-prodced root files present in...
ENABLED_HISTOS and MATRIX_... is extracted and u-
runs. From each file a matrix with MATRIX_NAME is extracted and u-

Important
It is required that the matrix has equal segment names on X axis and energy/spectrum u-
Something something label [keV].

Fit is performed in FIT_RANGE_START - FIT_RANGE_END range. Peak position is estimated by identifying bin with m-
available for PEAK_FIT_FUNCTION option:

$$f(x) = \frac{V}{\pi} \cdot (.x - \mu)^2 + \sigma^2 + p_0 + px$$

- Lorentzian

S2025_ancillaries_detail

Skewed Gaussian AgataAnalysisWS2025_ancillaries_general.pdf

 AgataAnalysisWS2025_prisma_analysis.pdf

 AgataAnalysisWS2025_prisma_introduction.pdf

AgataAnalysisWS2025_selector.pdf



Istituto Nazionale di Fisica Nucleare
Laboratori Nazionali di Legnaro



Configuration of the analysis

- Based on a configuration file and Lookup tables
- These files can be modified multiple times since analyzing an entire experiment often does not require more than 2 hours (depending on computing capabilities)

Board | channel | map | thr low | thr high | various par | ... | time offset | ncal | cal par 1 | ...

LUT_LABR.dat 1.10 KiB													
#	LaBr Co+Cs	#board (V1730)	channel map	name	thr_lo	thr_hi	theta	phi	TimeOffset	npar_gl p0_q1	p1_q2	npar_qs p0_qs	p1_qs
1	0	0	00	0	16000	90.422684	124.92098	0	2	-8.590549465	0.5683940043	2	-16.614035 0.584031
2	1	1	01	0	16000	84.308418	97.489398	0	2	4.994643769	0.441859949	2	10.570262 0.443247
3	1	2	02	0	16000	90.572894	73.768608	0	2	-4.882706373	0.4567364497	2	-9.782321 0.473778
4	1	3	03	0	16000	99.948116	51.748253	0	2	-2.68135951	0.4616749283	2	-9.040133 0.473527
5	1	4	04	0	16000	93.353077	26.981224	0	2	-3.368474921	0.4774816369	2	0.489657 0.481297
6	1	5	05	0	16000	94.007297	1.3778600	0	2	0	1	2	0 1
7	1	6	06	0	16000	99.883486	-28.723198	0	2	10.52197059	0.4435828877	2	18.918459 0.444711
8	1	7	07	0	16000	86.180070	-45.908423	0	2	12.53667474	0.4240481389	2	28.411274 0.421525
9	1	8	08	0	16000	91.699165	-66.505287	0	2	16.78408614	0.3897415818	2	35.049303 0.387539
10	1	9	09	0	16000	85.591641	-95.344627	0	2	-12.39452343	0.4289130669	2	-38.673472 0.452371
11	1	10	10	0	16000	85.591641	-95.344627	0	2	-12.39452343	0.4289130669	2	-38.673472 0.452371
12	1	11	11	0	16000	85.591641	-95.344627	0	2	-12.39452343	0.4289130669	2	-38.673472 0.452371
13	1	12	12	0	16000	85.591641	-95.344627	0	2	-12.39452343	0.4289130669	2	-38.673472 0.452371
14	1	13	13	0	16000	85.591641	-95.344627	0	2	-12.39452343	0.4289130669	2	-38.673472 0.452371
15	1	14	14	0	16000	85.591641	-95.344627	0	2	-12.39452343	0.4289130669	2	-38.673472 0.452371
16	1	15	15	0	16000	85.591641	-95.344627	0	2	-12.39452343	0.4289130669	2	-38.673472 0.452371

KEYWORD | value(s) | unit of measure | comment

```
#Format: | KEYWORD | value(s) | Unit of measure | Comment |
#Comments are ignored unit of measure # means none
#-----#
DETECTORS_PRESENT
MCP NO # EXOTIC Mcps are present YES/NO
SAURONFW NO # Sauron Forward is present YES/NO
EUCLIDES NO # Euclides is present YES/NO
LABR NO # Labr is present YES/NO
OSCARUP NO # Oscar Up is present YES/NO
SPIDER NO # Spider is present YES/NO
OSCARDOWN NO # Oscar Down is present YES/NO
AGATA NO # Agata is present YES/NO
SAURONBW NO # Sauron Backward is present YES/NO
DANTE NO # Dante is present YES/NO
EXSSIDE NO # EXSSIDE Forward is present YES/NO
PRISMA NO # Prisma is present YES/NO
#-----#
REPLAY_CONF
ENABLED_HISTOS enabled_histos.conf # File name with list of enabled histograms
SUM_ALL_PATTERN runs # Sum of all hadded files pattern
TREE_NAME TreeMaster # Input tree name
SUM_FILE_PATTERN sum # Hadded file pattern
OUT_FILE_PATTERN run_ # Output file pattern
IN_FILE_PATTERN Tree_ # Input file pattern
REPLAY_DIR_PATTERN run_ # Replay directory pattern
IN_SUB_PATH ./Out/Analysis # Input sub path
CONF_PATH ./Conf # Replay conf folder path
OUT_PATH ./Out # Output path
IN_PATH ./Data # Input path
#-----#
REACTION_CONF
REACTION_POSITION 0.5 # Position of the reaction in the target 0->front 0.5->m
ENERGY 0 MeV Beam energy
TARGET 1 1 # Target ion A Z
BEAM 1 1 # Beam ion A Z
ION 1 1 # Fragment of interest for binary reaction calculation:
#-----#
TARGET_CONF
ELOSS_TABLE_TYPE 0 # Type of eloss table 0 -> SRIM, 1-> Geant4, 2 ->
DEG_DISTANCE 0 um Degradation distance in um
DEG_THICKNESS 0 mg/cm2 Degradation thickness in mg
ROTATIONZ 0 deg Target rotation on the Z axis in degrees
ROTATIONX 0 deg Target rotation on the X axis in degrees
TILT 0 deg Target tilt in degrees; Negative values for cl
THICKNESS 0 mg/cm2 Target thickness-density in mg/cm2 or units al
DEG_PRESENT NO # Degradation present YES/NO
DEG_MATERIAL none # Degradation material
MATERIAL none # Target material
DEG_POS AFTER # Degradation position BEFORE/AFTER
#-----#
```

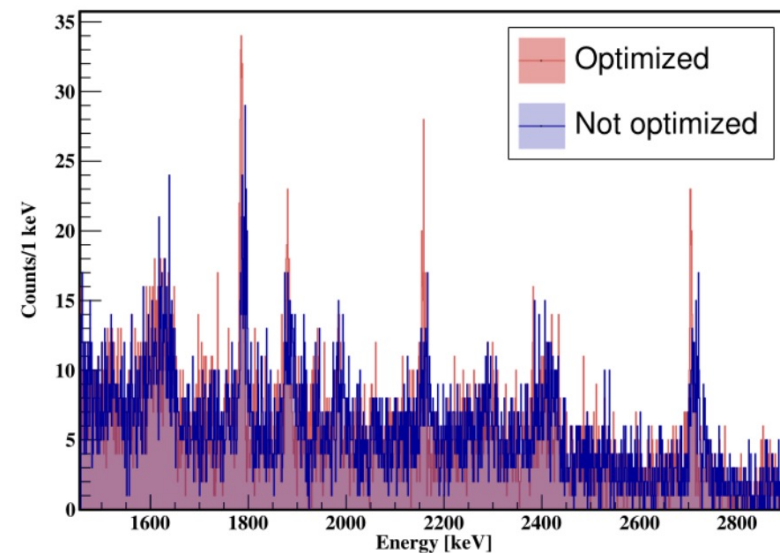
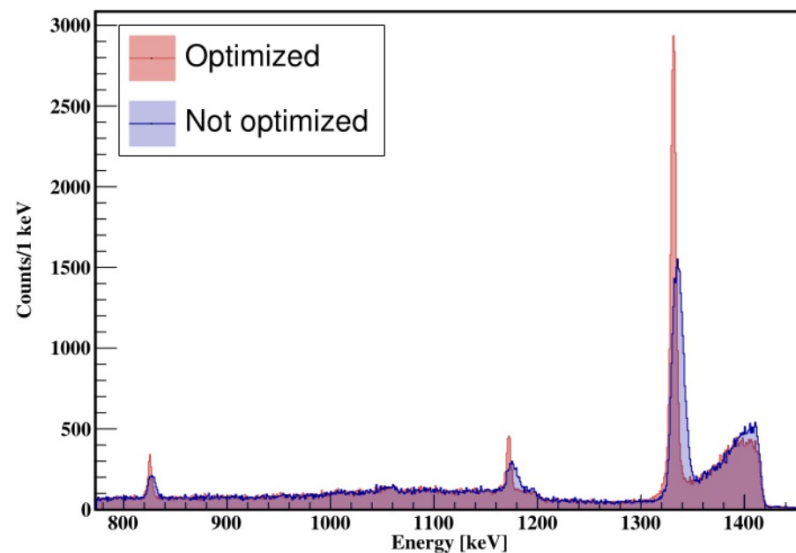
The UserSelector

- The code can be personalized for a specific experiment if something non-standard is needed
- The standard analysis can be accessed by the user and does not affect other experiments
- One can recall the configuration and specific code of the nearline analysis as by compiling the selector with an experiment-specific cmake option

h UserSelector.h 592 B

```
1  #pragma once
2  #include "AgataSelector.h"
3
4  class UserSelector : public AgataSelector {
5  public:
6      explicit UserSelector(const std::string& option
7                          : AgataSelector(options){};
8      Bool_t Process(Long64_t entry) override;
9      void SlaveBegin(TTree* tree) override;
10     void SlaveTerminate() override;
11
12 private:
13     // USER variable and histogram definition secti
14
15     // unsigned long long timeRef = 0;
16
17     struct UserHistograms {
18         std::vector<TObject*> ptrs;
19         TDirectory* dir{nullptr};
20         // TH1D *h_agataCR = nullptr;
21
22     } userHistograms;
23 };
24
```

The optimization procedure

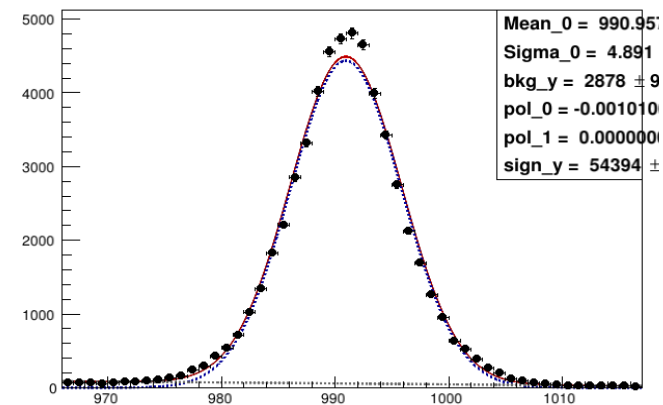
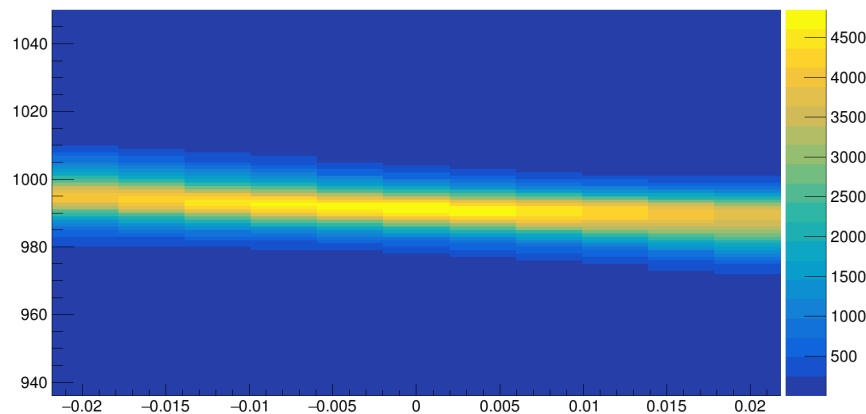


- Remarkable improvements are possible with the optimization but are experiment dependent.
- The selector contains a procedure to find the optimal parameters by running
RunSelector - -optimize 2

Optimization details

- The optimization has three options:
 - Scan a parameter within some limits
 - Fit + simultaneous variation of any parameter of the configuration file
 - Lookup table automatic optimization: two options are available and documented (developed by M. Balogh)

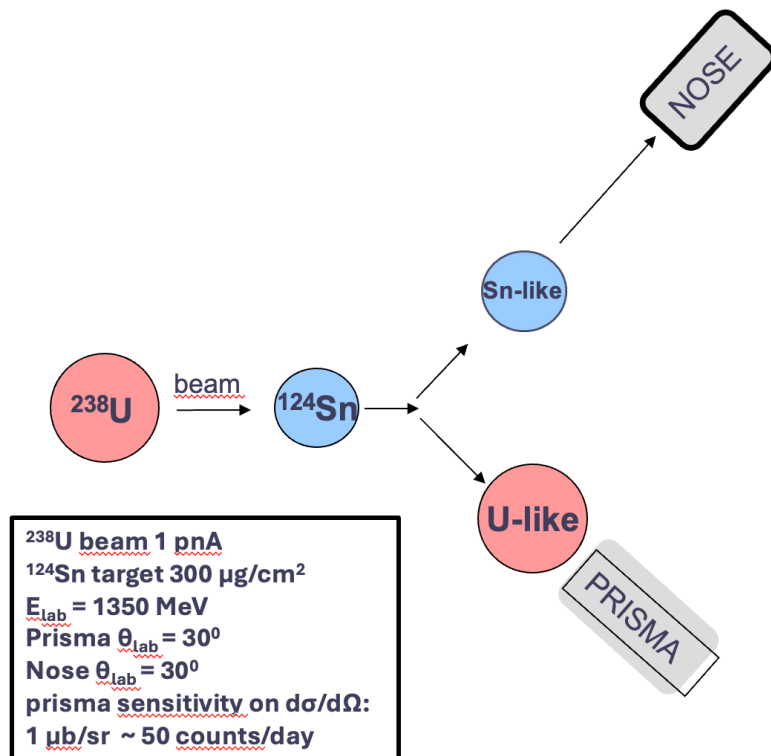
SPIDER_CONF_X_SHIFT_{EDC}



Some recent results from the
nearline analysis of the ^{238}U beams

Recent additions: the NOSE

- Kinematic coincidence between PRISMA and NOSE
- Bragg chamber (Z-id) and PPAC (X-Y and time)
- Especially useful to reject fission events

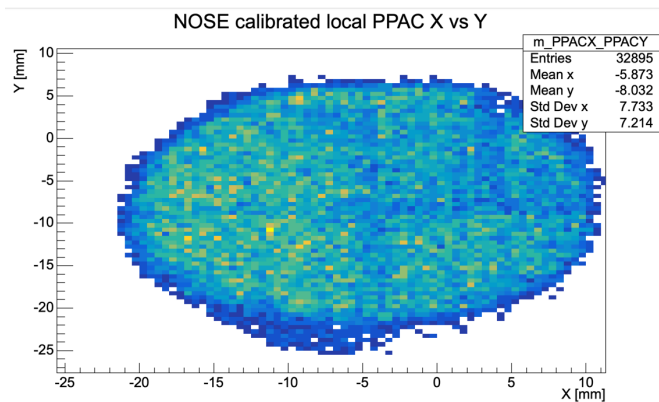


New implementation!

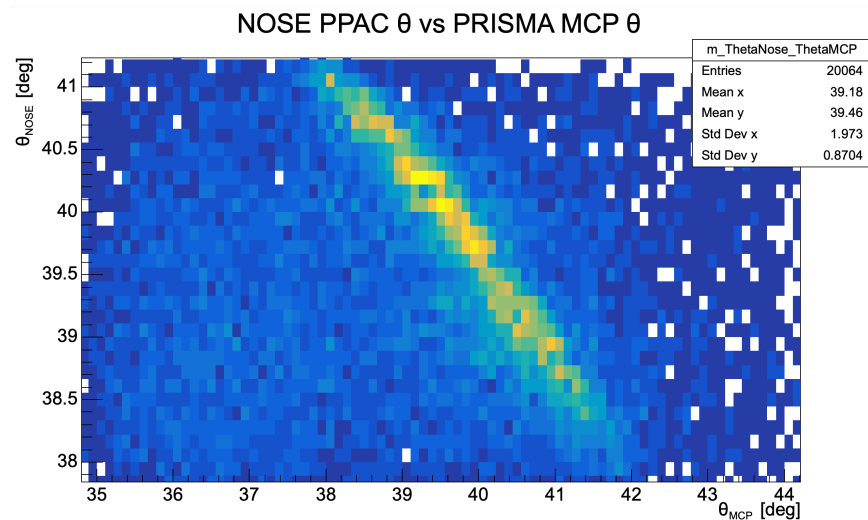
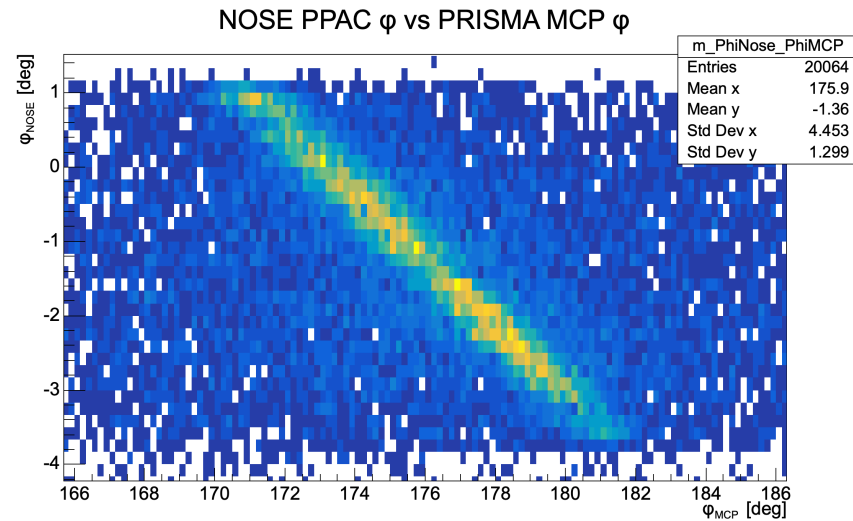
- NOSE analysis
- PRISMA-NOSE coincidence
- PRISMA-NOSE-AGATA coincidence

Courtesy of F. Galtarossa, L. Corradi and B. Sulignano

PRISMA-NOSE coincidence

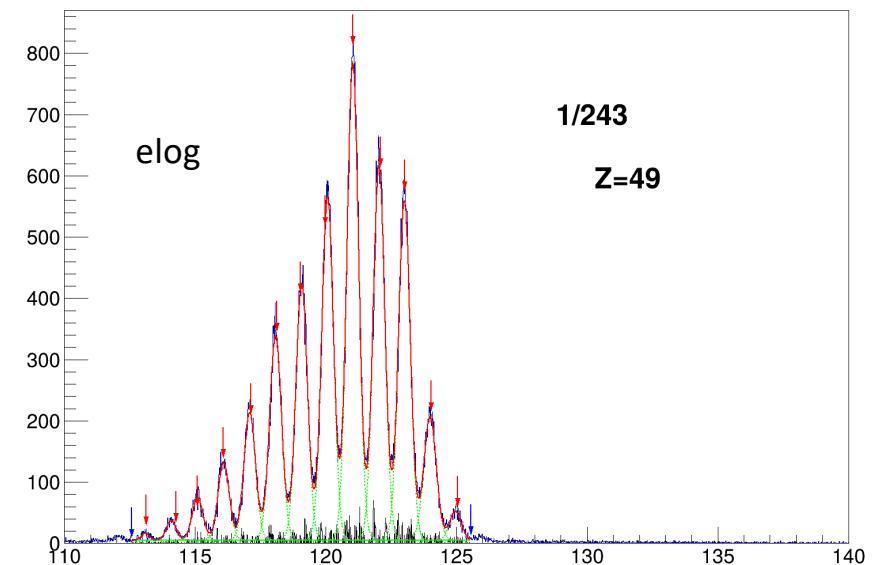


- The kinematics correlations are clear!
- The correlation of tof and energy allows the separation of heavy and light fragments



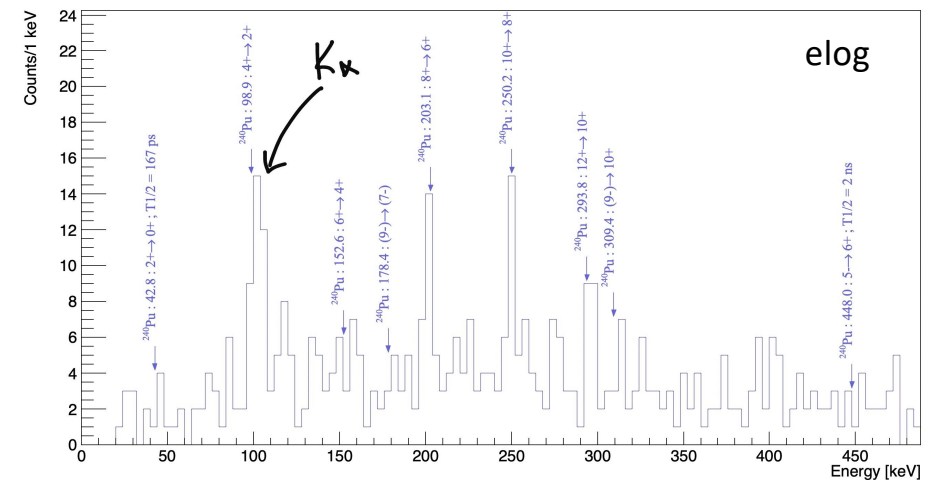
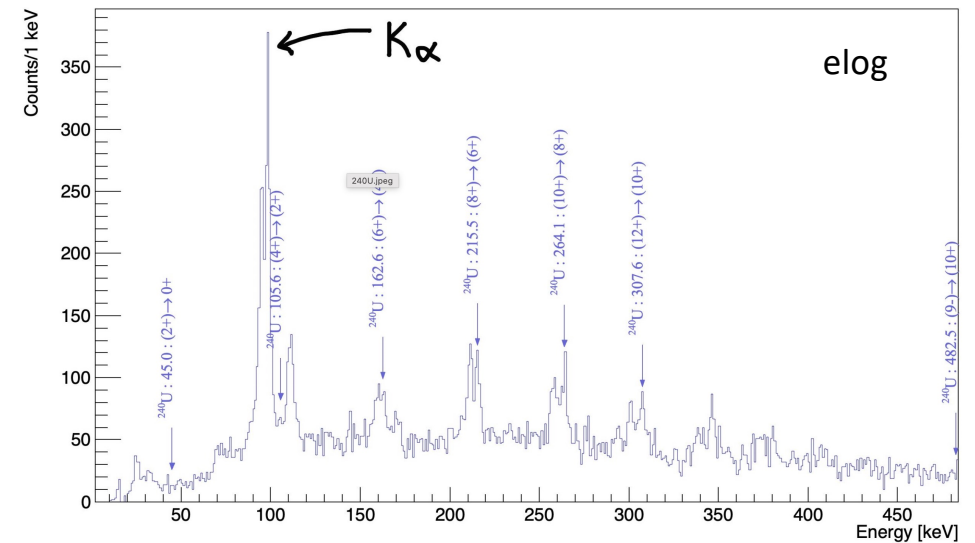
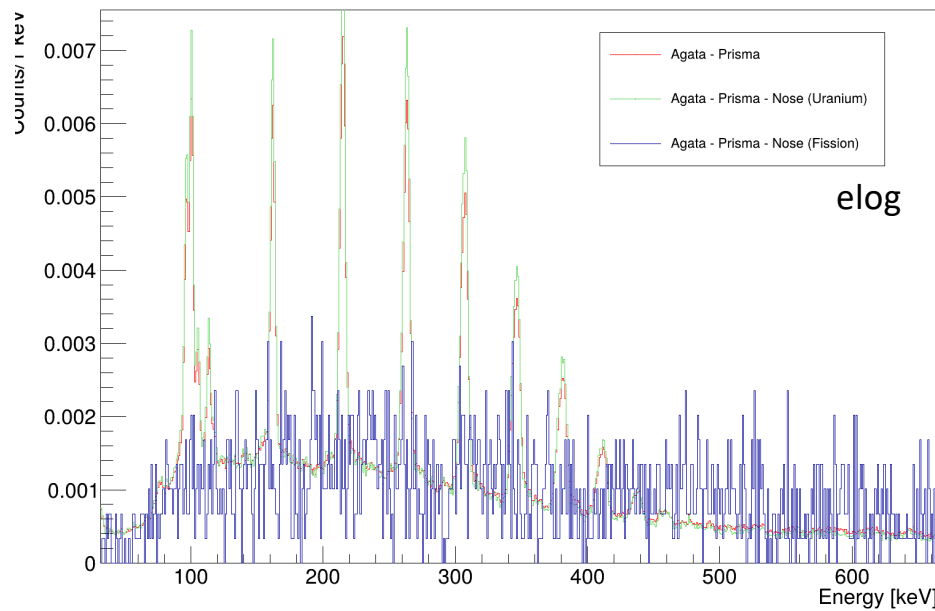
Improvements on the PRISMA analysis

- Recent improvements on the interactive procedure (gate-setting, calibration, thresholds, ..) thanks to L. Busak and F. Angelini
- It can now read directly raw data
- Faster and more streamlined procedure
- Great mass resolution already in the nearline



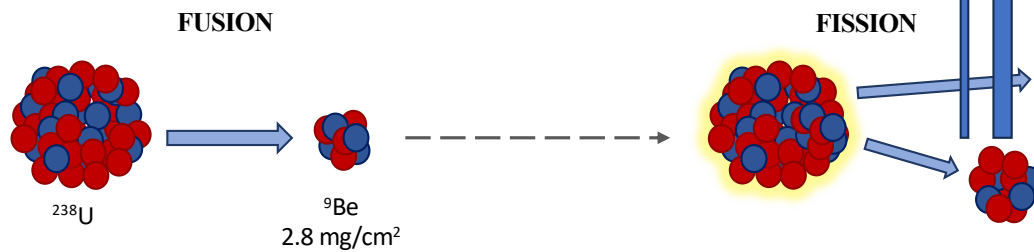
AGATA-PRISMA-NOSE coincidence

- Great fission rejection performance
- The background is reduced especially in for the transfer channels

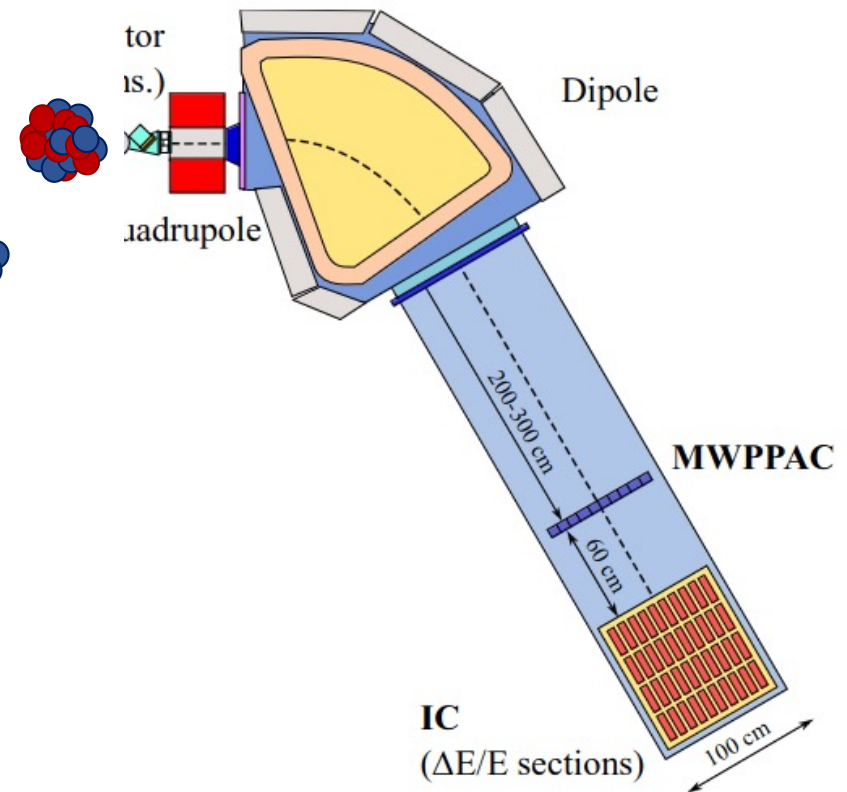
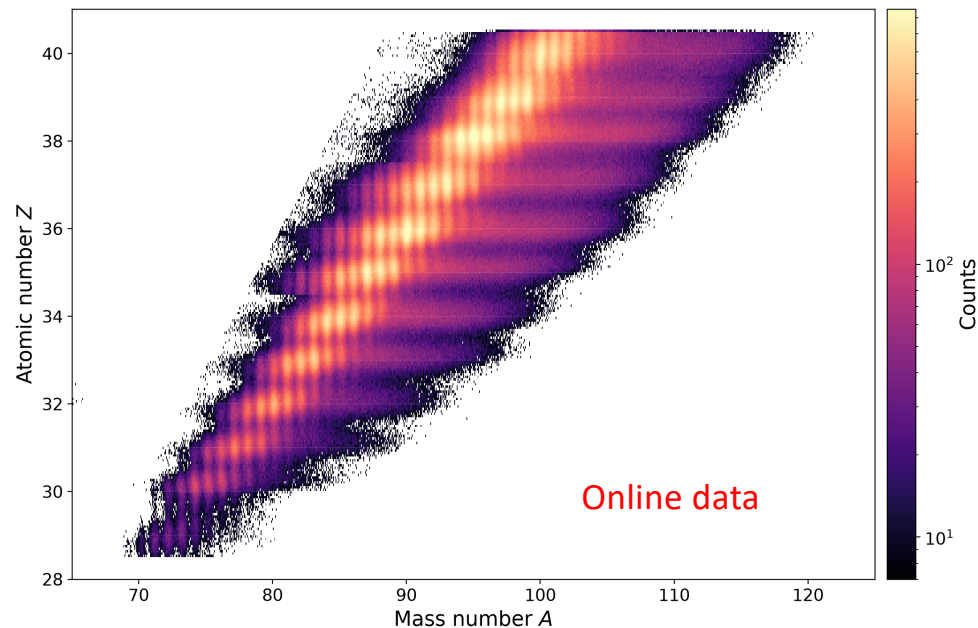


Fusion-fission and lifetime measurements

^{238}U beam at LNL!



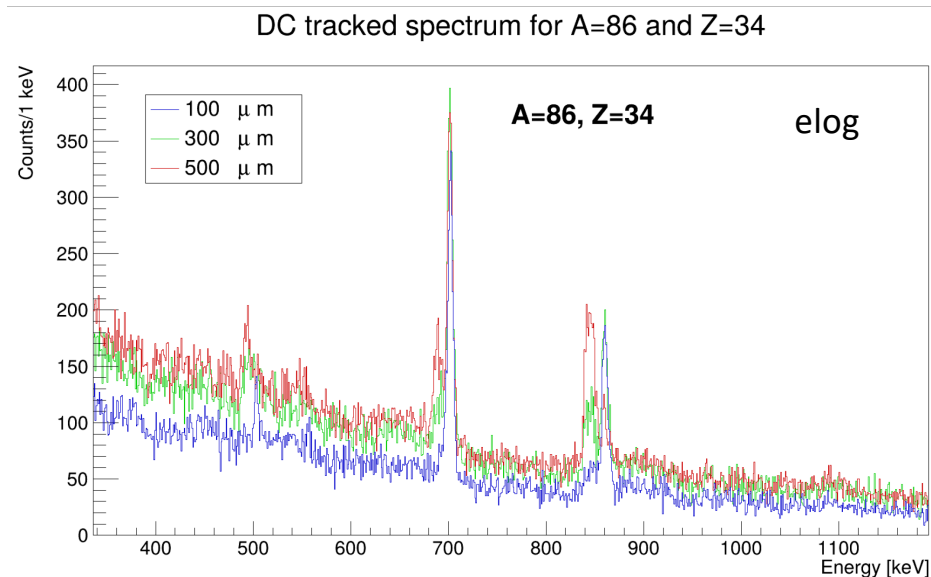
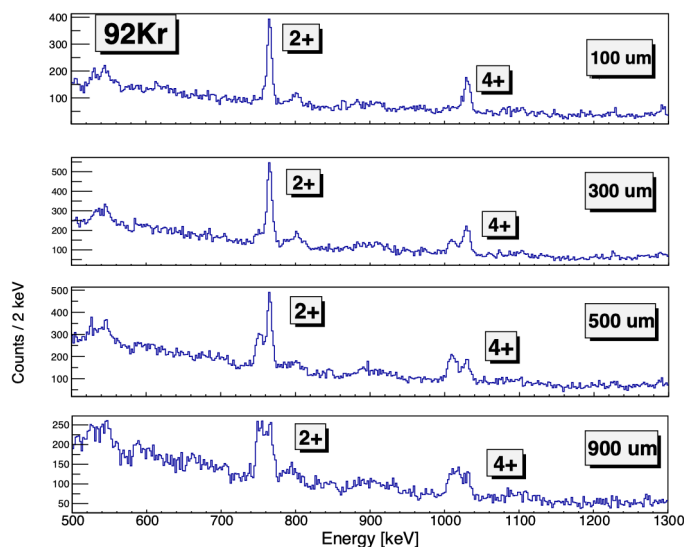
Fission yield identified by PRISMA



Courtesy of A. Gottardo, J.J. Valiente
Dobon, M. Camaño, F. Angelini, D.
Stramaccioni

AGATA-PRISMA coincidences

Thanks to the online analysis of F. Angelini, L. Busak, M. Del Fabbro, B. Gongora, E. Pilotto, B. Greaves, and many other nearline experts



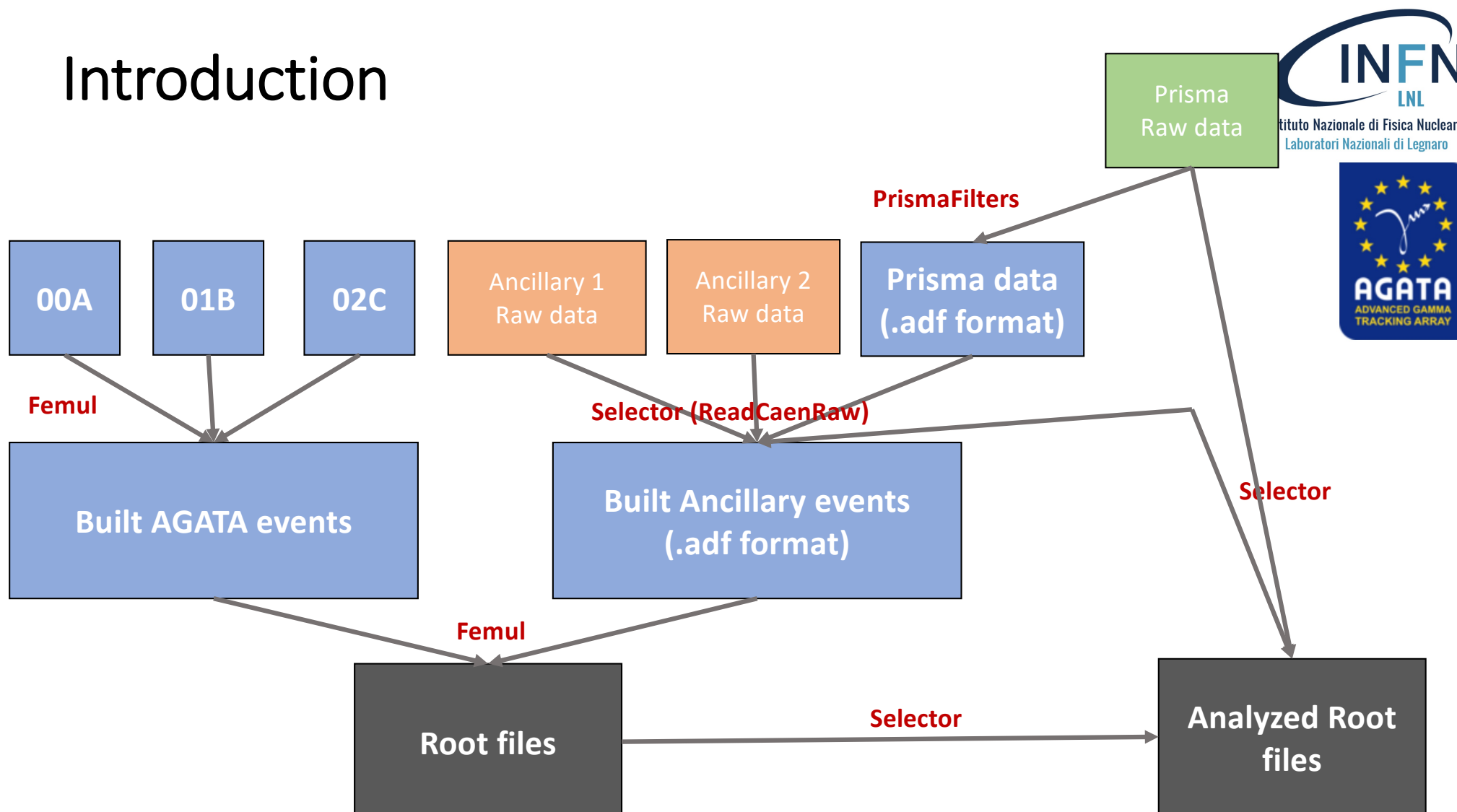
Perspectives

- We plan to switch to DAEMULE (developed by Guillaume Baulieu) for the ZD campaign in the nearline
- The analysis of NEDA and PARIS will be integrated and developed with the existing codebase

**Thanks to all people that have contributed with
solving bugs, improvements and new
functionalities!**

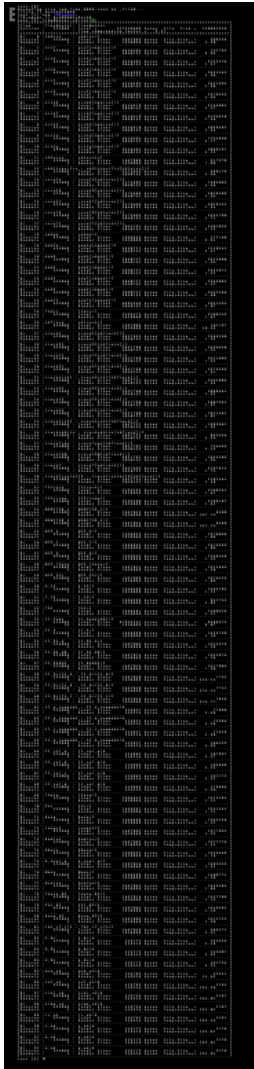


Introduction



The output

- The selector produces ROOT files containing:
 - Histograms (each analysis is contained within a folder):
 - Single detector analysis
 - Coincidence analysis
 - TTrees of:
 - High level data of a single detector. For instance, Spider provides things such as excitation energy or angles
 - Doppler correction based on the analysis of the agata+ancillary coincidence
- The analysis code handles kinematic reconstruction, energy loss, event selection, time and energy gates, optimization procedures
- It can be distributed on multiple machines if compiled with MPI



The starting point

- Femul produces a root file containing MANY leaves
- The analysis procedure is common to all experiments and there is little benefit of repeating the same steps over and over
- **The users can profit from a collaborative project for a quicker analysis**
- The code was created for a quick near-line analysis and has since evolved with more refinements with full analysis capabilities



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