

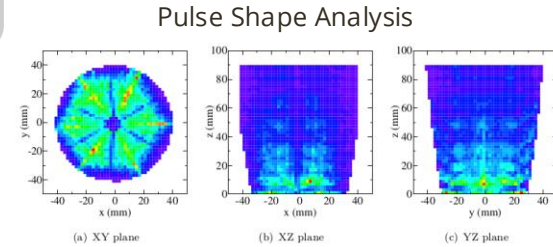
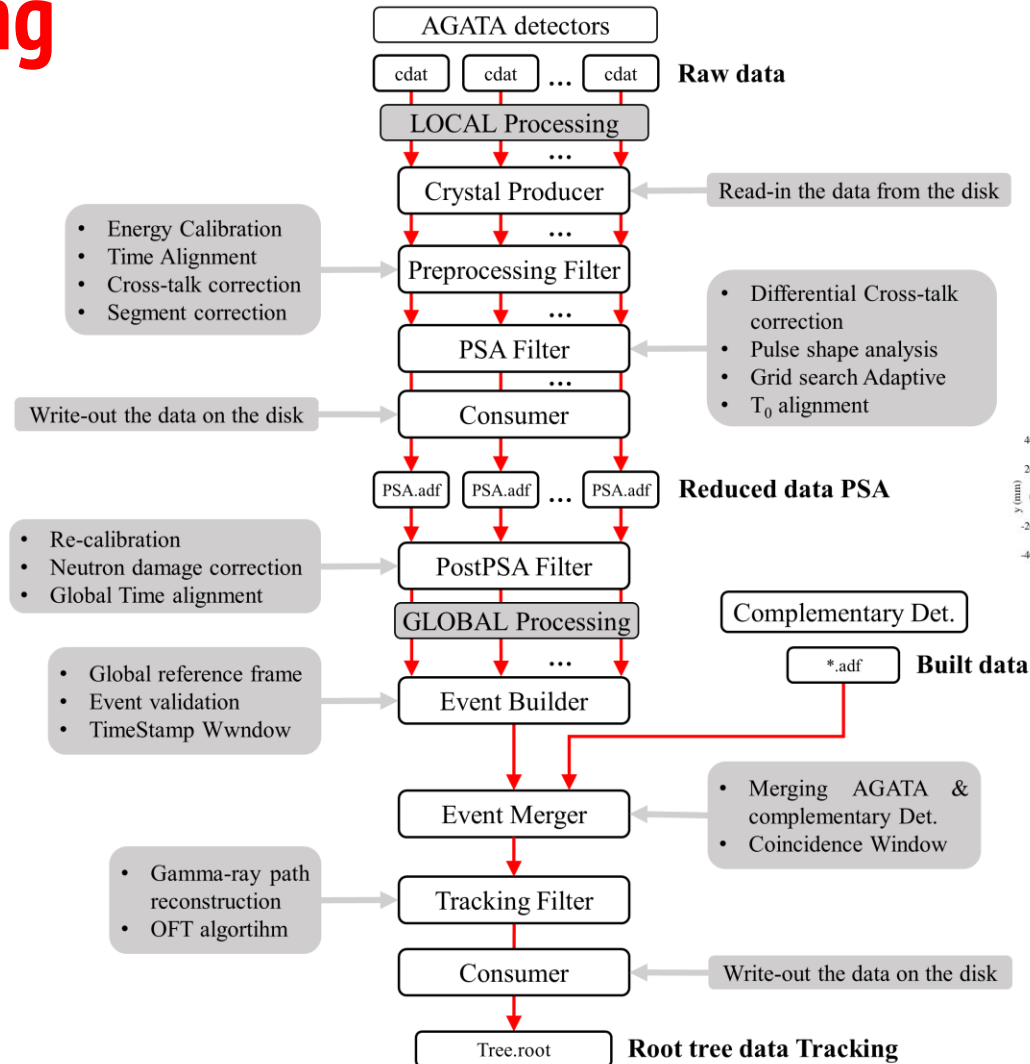
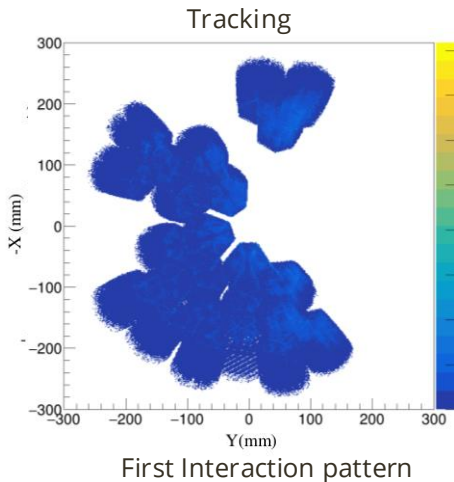
AGATA Analysis Workshop 2025

PostPSA Calibration

R.M. Pérez-Vidal

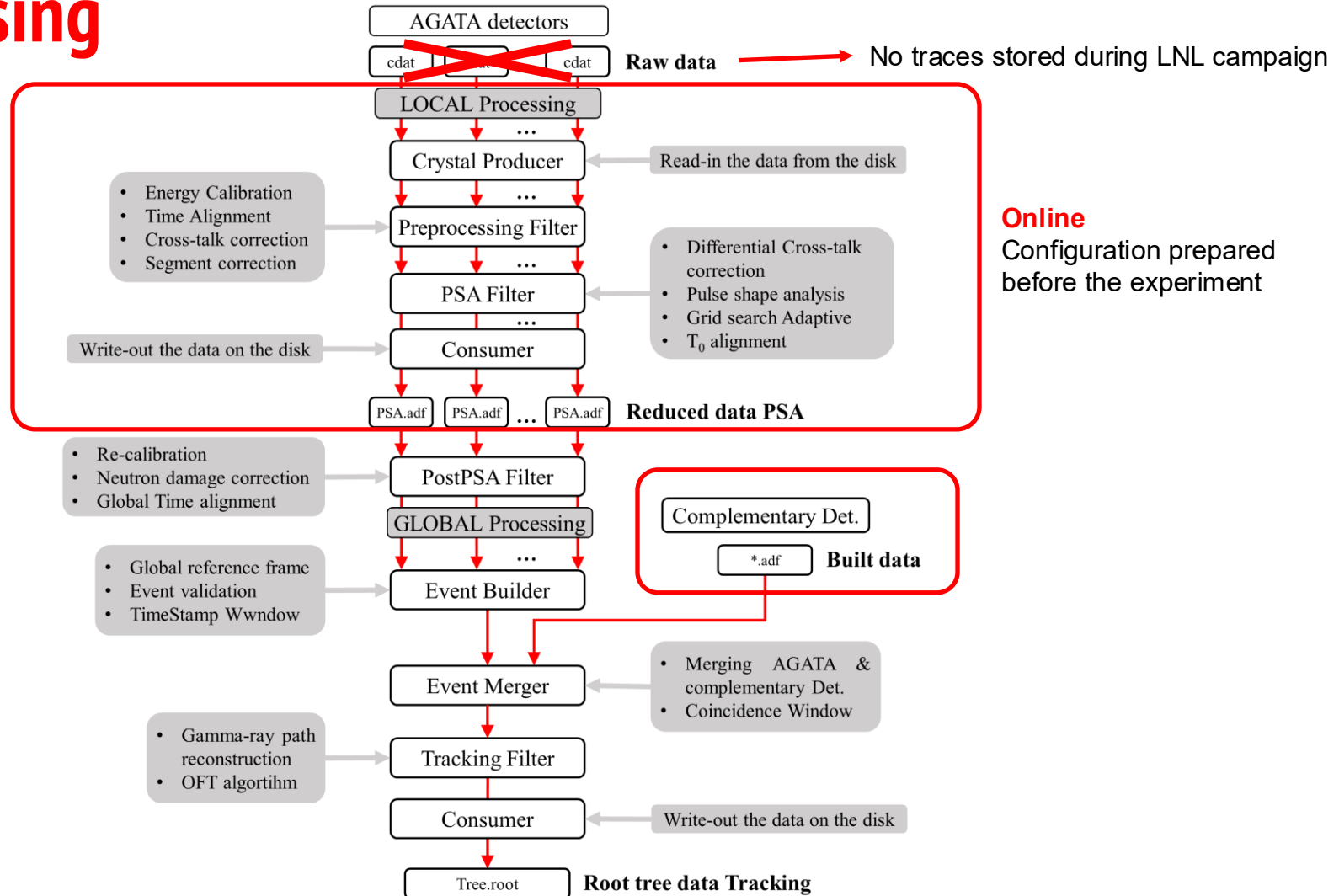
14/01/2025, Lyon

Narval actors



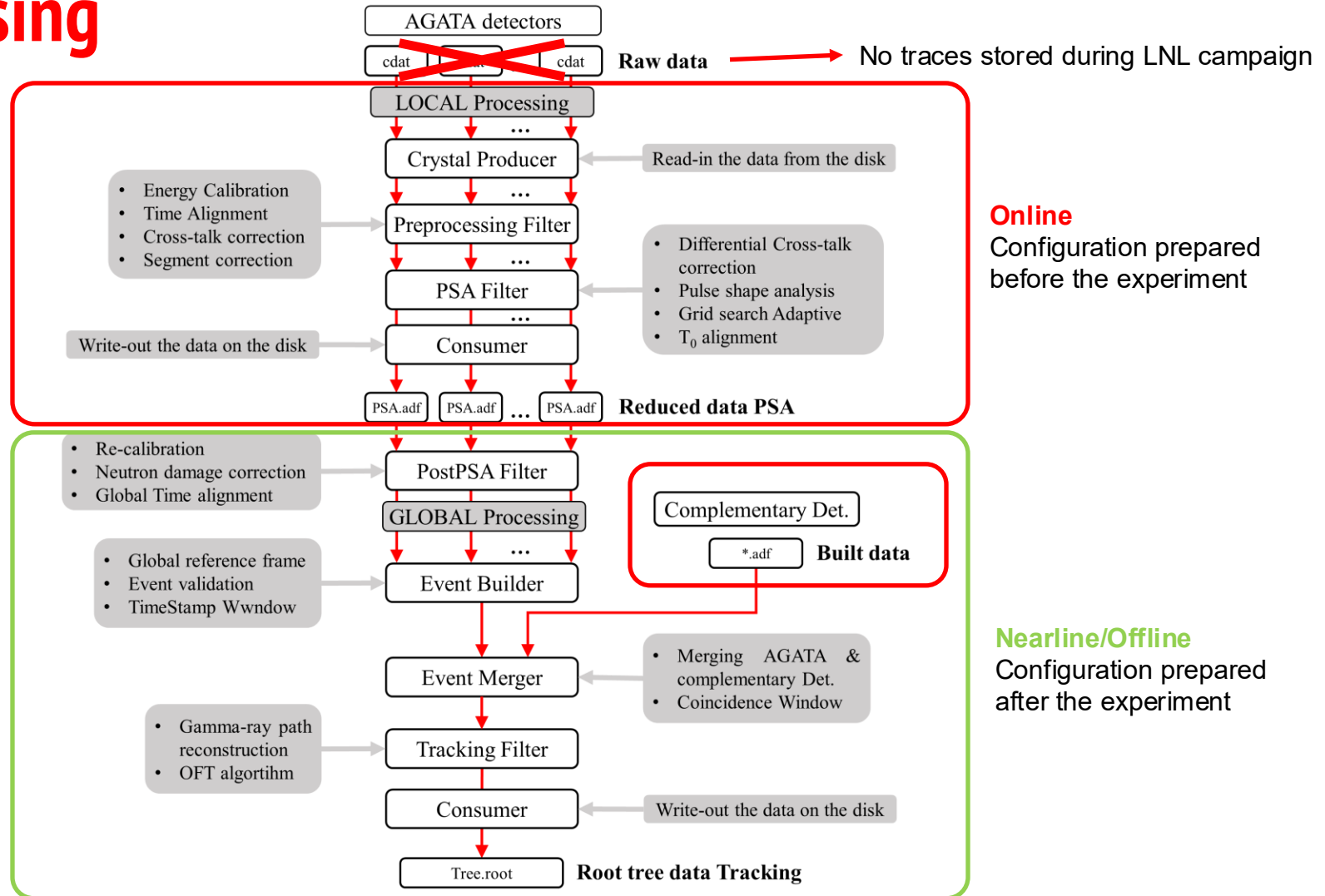
Data Processing

Narval actors



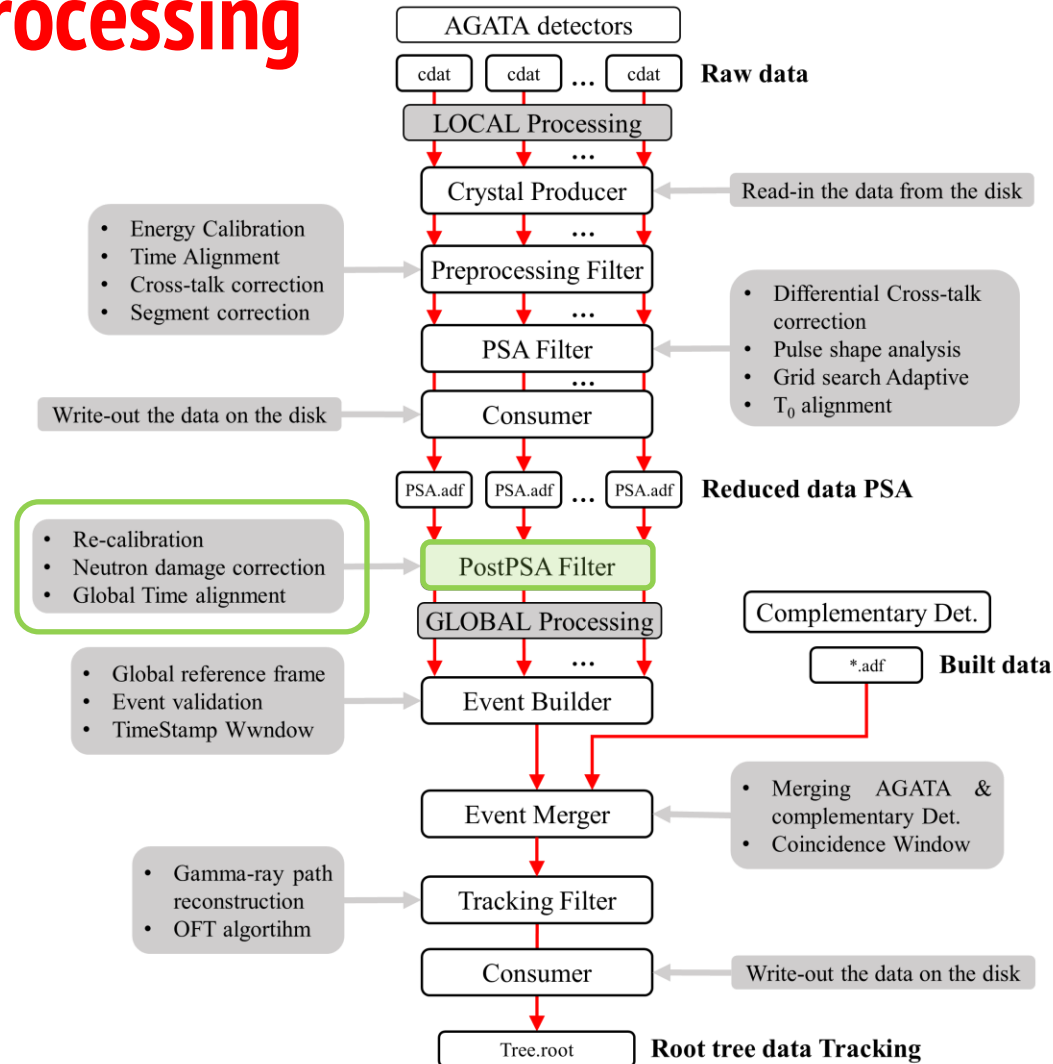
Data Processing

Narval actors



Local Level Processing

Narval actors



Local Level Processing

PostPSA Filter

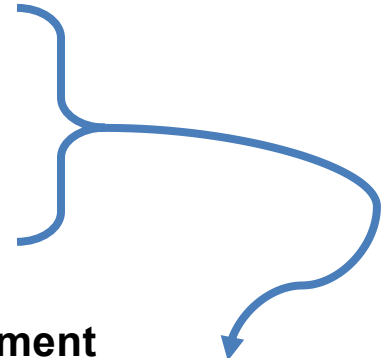
- Performs:

- Recovery (partial) of neutron damage using info from PSA: **Trapping.cal**
- Realibration over time (core): **TimeEvoCC.conf**
- Final energy calibrations with offset: **RecalEnergy2.cal**, **gen_conf.py**
- Force segments to core (optional) : **gen_conf.py**
- Global Time alignment: **gen_conf.py**

- **Configuration for this actor can be done by users during/after the experiment**

- Generates one file:

- Post__5-40-16384-UI__Ener.spec → **File in Out/00A e.g.**



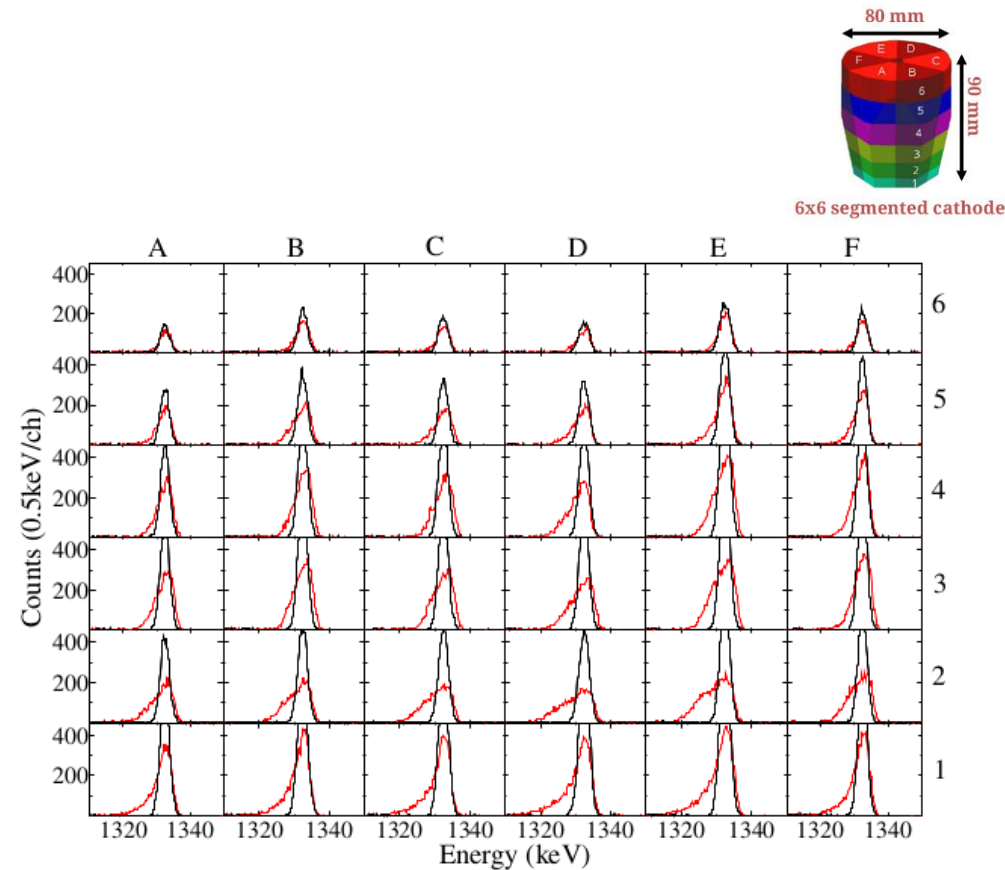
**Files in Conf/00A e.g.
gen_conf.py its outside the
configuration directory**

PostPSA Filter

1. Neutron Damage correction

- **Fast neutrons** are well known to produce specific lattice defects in germanium crystals which act as **efficient hole traps**.
- Reduction in the charge collection efficiency of the detectors observable by **a low energy tailing** on the energy line shape
- AGATA crystals are n-type HPGe detectors, which are found to be less sensitive to the neutron radiation in terms of central contact signals
- The energy deficit can be **corrected using the position information obtained by the PSA**

SortPsaHits program applies to the PSA hits a grid of **correction parameters for the electron and hole trapping**, determining the optimum set parameters for all the detector channels **that minimizes the FWHM and the tail on the left** side of the energy peaks in the spectra.



PostPSA Filter

1. Neutron Damage correction

Replay to generate Psa__0-16-F__Hits.fdat files
femul key in PSAFilter: "WritePSAHits",

What is needed:

- Long 60Co run
- PSA hits file : Data/{crystalID}/Psa__0-16-F__Hits.fdat
- Conf File: **Trapping.cal, gen_conf.py**
- Auxiliary files: Pso__2-4-40-2048-UI__Ener.spec
- Programs/scripts:

femul key in PostPSAFilter:
"Trapping Trapping.cal",
Add it in gen_conf.py

SortPsaHits: get the optimum lambdaE lambdaH parameters and generates Pso__2-4-40-2048-UI__Ener.spec

```
SortPsaHits -f Psa__0-16-F__Hits.fdat -best 1300 1350 -bpar 1 10000 0
```

```
SortPsaHits -f ../Data/{crystalID}/Psa__0-16-F__Hits.fdat -gain 5 -offs 5000 -fcal Trapping.cal
```

RecalEnergy: generate calibration coefficients for the different columns of Trapping.conf

```
RecalEnergy -spe Pso__2-4-40-2048-UI__Ener.spec -num 40 -sub 0 -offs -5000 -gain 5 -noTR -dwa 25 2 | tee Recal_SG_orig.txt
```

colupdate.py: add these coefficients to the different columns of Trapping.conf

```
./colupdate.py -c 1 13 Trapping.cal Recal_SG_orig.txt -o Trapping.cal
```

The Trapping.cal file has 36 lines, one per segment:

3 steps process

1. RecalEnergy, SortPSAHits (Check) [Optional]
2. SortPSAHits (Check)
3. RecalEnergy, SortPSAHits (Check)

#SG	gainSG_orig	gainCC_orig	lambdaE	lambdaH	gainSG_corr	gainCC_corr
0	1.	1.	51.6	6.6	1.	1.
1	1.	1.	269.3	6.6	1.	1.
2	1.	1.	51.6	6.6	1.	1.
...						
35	1.	1.	104.4	8.9	1.	1.

More details in [AGATA_LL_P_UsersGuide](#)

PostPSA Filter

1. Neutron Damage correction

What is needed:

- Long 60Co run
- PSA hits file : Data/{crystalID}/**Psa__0-16-F__Hits.fdat**
- Conf File: **Trapping.cal, gen_conf.py**
- Auxiliary files: Pso__2-4-40-2048-UI__Ener.spec
- Programs/scripts:

Replay to generate Psa__0-16-F__Hits.fdat files
femul key in PSAFilter: "WritePSAHits",

femul key in PostPSAFilter:
"Trapping Trapping.cal",
Add it in gen_conf.py

NEW!

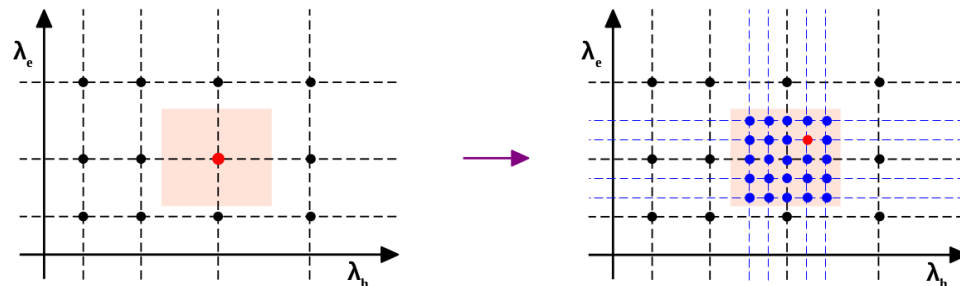
Neutron damage correction improvements:

- For SG+CC, we employ a normalization of the whole grid of FOMs (such that the highest FOM has value 1) before calculating the average
 - Implementation of an Adaptive Grid – Search
- “-size” option:** variable grid size, default is kept at nH=100, nE=50
- “-algo” option:** user must choose number of iterations and which spectra to optimize (0=SG, 1=CC or 2=SG+CC). Default values are 1 2
- “zoom” option:** user can specify the magnification factor M between iterations. Default is 0.25

Example of command:

```
SortPsaHits -f Psa__0-16-F__Hits.fdat -best 1300 1350 -bpar 1 10000 0 -size 50 30 -algo 3 2
```

More details in [AGATA_LL_P_UsersGuide](#)



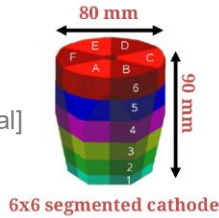
Elia Pilotto Agata Week 2024

PostPSA Filter

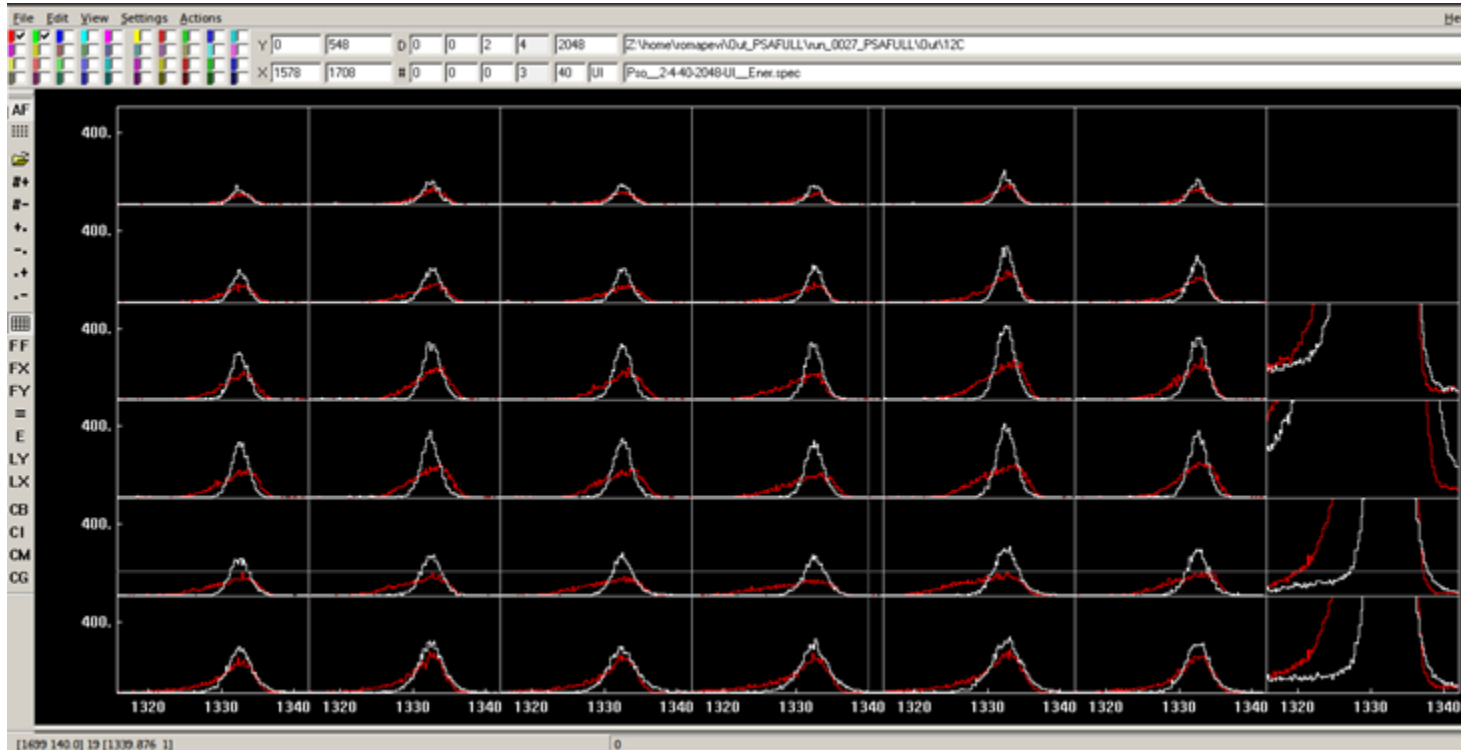
1. Neutron Damage correction

3 steps process

1. RecalEnergy, SortPSAHits (Check) [Optional]
2. SortPSAHits (Check)
3. RecalEnergy, SortPSAHits (Check)



Pso_2-4-40-2048-UI_Ener.spec [0][1][all] before (red) and [0][3][all] after (white) the neutron correction



[0-SG,1-CC]

[0-orig,1-orig+recal,2-corr,3-corr+recal]

[0-39]:

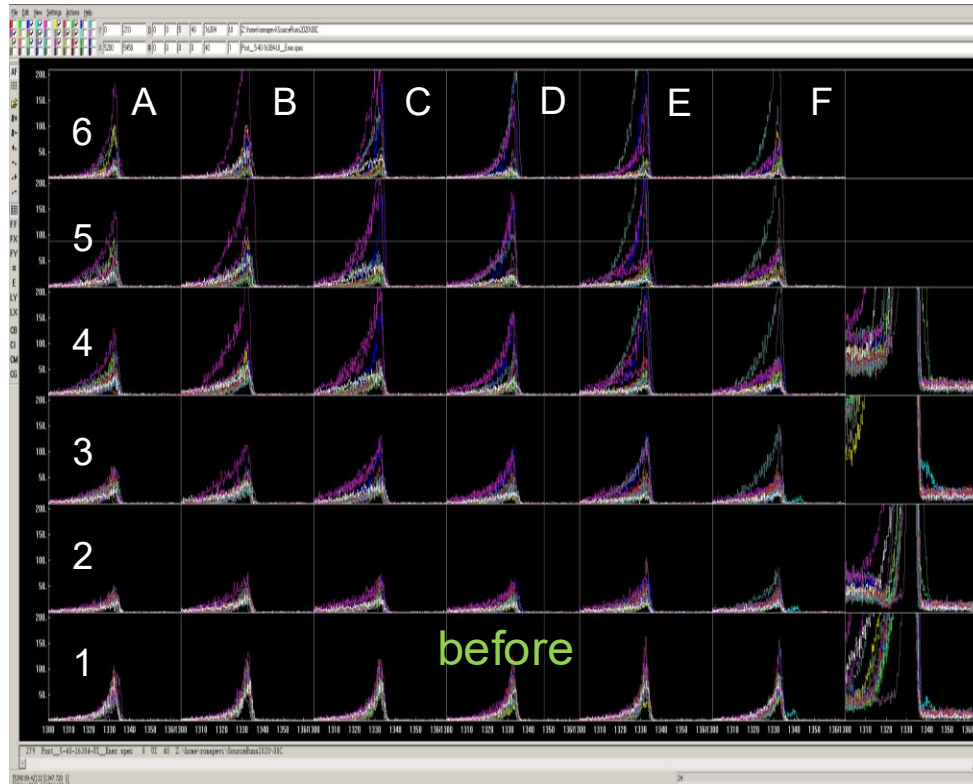
- 0-35 segments
- 36 Or of segments/cores
- 37 Or of segments/cores M=1
- 38 Sum of segments/cores
- 39 Average SumSegs+SumCC

#SG	gainSG_orig	gainCC_orig	lambdaE	lambdaH	gainSG_corr	gainCC_corr
0	1.	1.	51.6	6.6	1.	1.
1	1.	1.	269.3	6.6	1.	1.
2	1.	1.	51.6	6.6	1.	1.
...						
35	1.	1.	104.4	8.9	1.	1.

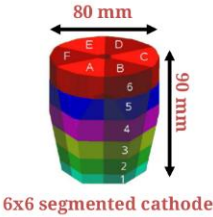
PostPSA Filter

1. Neutron Damage correction

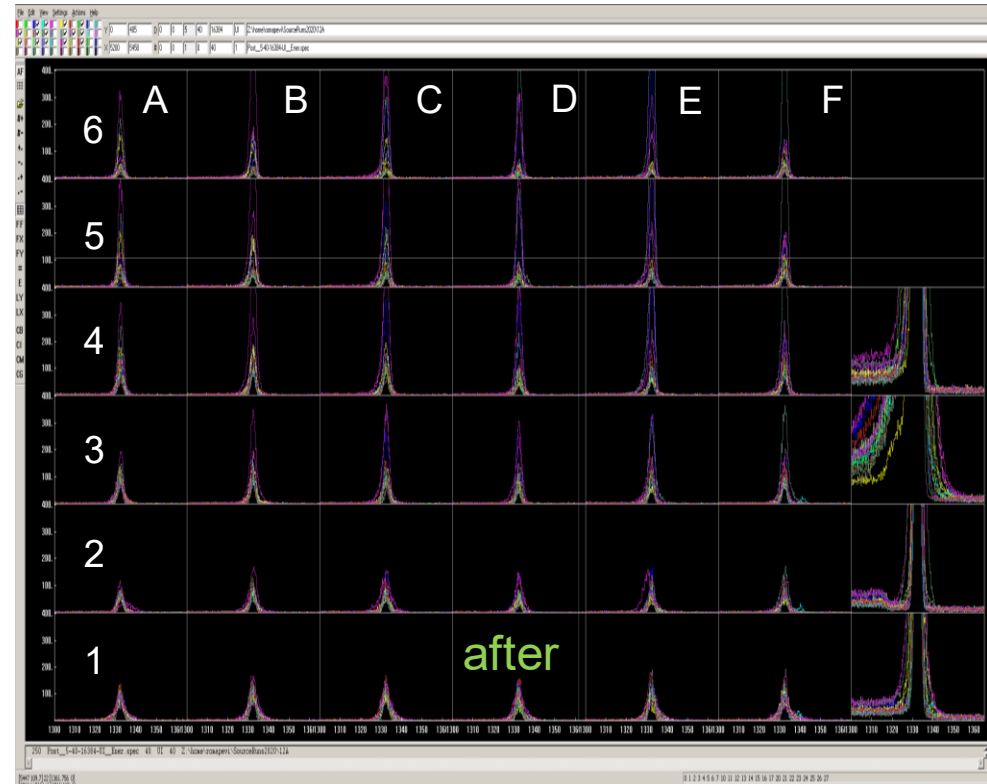
Post__5-40-16384-UI__Ener.spec [0] [0-39]



Verification with femul replay



Post__5-40-16384-UI__Ener.spec [1] [0-39]



PostPSA Filter

2. Recalibration

The recalibration of segments and cores is a several steps process and one must be careful of the possible redundant calibration done by the PostPSA filter actor:

- 0) Recalibration of Segments from file RecalEnergy1
- 1) Recalibration of Segments and Cores from the Trapping file

Filling histograms libraries 0 to 3 in PostSpecEner

- 2) Correction for energy drifting over time (Core energy)
- 3) Recalibration of Segments from file RecalEnergy2
- 4) Core recalibration from RecalCC and Segments recalibration from RecalSG
- 6) Force segments to core

Filling histograms library 4 in PostSpecEner

PostPSA Filter

2. Recalibration

The recalibration of segments and cores is a several steps process and one must be careful of the possible redundant calibration done by the PostPSA filter actor:

~~Recalibration of Segments from file RecalEnergy1~~ ➡ Typically not used

~~Recalibration of Segments and Cores from the Trapping file~~ ➡ performed in the neutron damage correction

Filling histograms libraries 0 to 3 in PostSpecEner

0) Correction for energy drifting over time (Core energy) **NEW!**

1) Recalibration of Segments from file RecalEnergy2

2) Core recalibration from RecalCC and Segments recalibration from RecalSG

3) Force segments to core

Filling histograms library 4 in PostSpecEner

PostPSA Filter

Replay to generate ROOT files
femul Topology_Global_Tree.conf

2.0 Recalibration: Correction for energy drifting over time

What is needed:

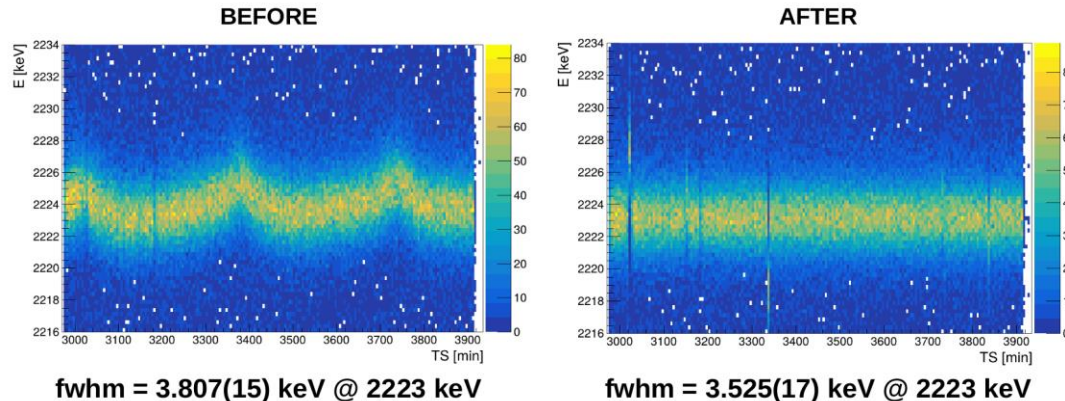
- Long run with high energy transitions
- Root file : Out/Analysis/**Tree_0000.root**
- Conf File: **TimeEvoCC.conf**, **gen_conf.py**
- Programs/scripts:

NEW!

Cross – correlation Correction Method (CCM)

[Matus Balogh NIM paper](#)
[GitHub code](#)

- Create a matrix Core energy versus time and divide in slices
- For each a projection, calculate product between a reference spectrum and the shifted projection
- Perform a scan for each slice



- It has been observed that especially when using the lower energy gain of the core contact gain oscillations of the order of 1/1000 can occur
- Pseudo – periodic with 6h period (AGATA filling)
- More pronounced in detectors placed in the lower part of Agata, pointing upwards
- **To be corrected before the recalibration with offset of the core (RecalCC)**

The TimeEvoCC.conf file must contain three columns with the following format:

#TS_start	TS_stop	gain
6150000000000	6210000000000	0.999948755522396
6210000000000	6270000000000	0.999887585155303
...		

femul key in PostPSAFilter:
"TimeEvoCCFile TimeEvoCC.conf",",
Add it in gen_conf.py

PostPSA Filter

2.1 Recalibration: RecalEnergy2

Verification with femul replay each step

3 steps process

1. RecalEnergy2 (Check)
2. RecalCC and RecalSG (Check)
3. ForceSegmentstoCore (Check)[Optional]

What is needed:

- Long 152Eu run
- PostPSA file : Out/{crystalID}/Post__5-40-16384-UI__Ener.spec
- Conf File: **RecalEnergy2.cal, gen_conf.py**
- Auxiliary files: Recal2.txt
- Programs/scripts:

RecalEnergy: generate calibration coefficients

The RecalEnergy2.cal file has 36 lines, one per segment:

#segm	%d(id)	%f(offset)	%f(egain)
segm	0	0.216	1.001157
segm	1	7.040	0.996854
segm	2	-0.732	1.003805
...			
segm	35	-1.083	1.002038

```
RecalEnergy -spe Out_norecal/{crystalID}/Post__5-40-16384-UI__Ener.spec -sub 160 -num 36 -gain 4 -poly1 -152Eu
```

#	indx	#spec	#pks	#ok	rEnergy	FW05	FW01	Area	Position	Width	Ampli	WTML	WTMR	slope*gain	rChi2%	offs1*g	slope1*g	rChi2%
	0	40	32	10	1408.15	5.032	12.662	4294	5624.78	16.5	171	4.320	1.823	1.001387	1.74	0.216	1.001157	0.42
	1	41	18	8	1409.43	9.080	18.768	2425	5614.94	36.0	59	2.329	1.823	1.004054	999.99	7.040	0.996854	91.65
	2	42	24	10	1407.05	7.637	17.111	3068	5611.22	29.1	85	2.861	1.823	1.003023	52.70	-0.732	1.003805	41.66
	3	43	25	9	1407.38	5.256	13.774	3854	5624.43	15.7	144	5.210	1.823	1.000901	29.62	-1.073	1.001038	2.44
	4	44	20	9	1408.21	4.658	12.739	2799	5621.39	12.1	116	6.562	1.823	1.002036	6.53	0.213	1.001812	6.24
	5	45	14	9	1408.15	4.126	10.131	1331	5623.55	14.1	65	3.909	1.823	1.001609	37.60	-0.389	1.002013	38.46
	6	46	29	10	1407.69	5.470	13.087	4393	5626.02	19.5	165	3.546	1.823	1.000845	9.79	0.293	1.000532	8.18
	7	47	22	10	1408.45	6.756	17.530	2612	5619.43	20.6	76	4.964	1.823	1.002556	21.86	0.213	1.002329	23.10
	8	48	22	10	1406.20	7.643	17.149	3859	5615.23	29.2	107	2.873	1.823	1.001704	58.23	-0.356	1.002085	61.32
	9	49	27	5	1408.70	7.257	18.956	4738	3839.23	17.0	186	3.888	2.197	1.467693	919.64	3.908	1.461898	601.93

colupdate.py: add these coefficients to the 3rd and 4th columns of RecalEnergy2.cal

```
./colupdate.py -c 2 14 RecalEnergy2.cal Recal2.txt -o RecalEnergy2.cal
```

```
./colupdate.py -c 3 15 RecalEnergy2.cal Recal2.txt -o RecalEnergy2.cal
```

femul key in PostPSAFilter:

"RecalEnergy2 RecalEnergy2.cal",
Add it in gen_conf.py

PostPSA Filter

2.2 Recalibration: RecalCC & RecalSG

Verification with femul replay each step

3 steps process

1. RecalEnergy2 (Check)
2. RecalCC and RecalSG (Check)
3. ForceSegmentstoCore (Check)[Optional]

What is needed:

- Long 152Eu run
- PostPSA file : Out/{crystalID}/**Post__5-40-16384-UI__Ener.spec**
- Conf File: **gen_conf.py**
- Auxiliary files: Recal2.txt
- Programs/scripts:

RecalEnergy: generate calibration coefficients

- For the **core** recalibration with offset

```
RecalEnergy -spe Out_norecal/{crystalID}/Post__5-40-16384-UI__Ener.spec -sub 199 -num 1 -gain 4 -poly1 -152Eu
```

#	indx	#spec	#pks	#ok	rEnergy	FW05	FW01	Area	Position	Width	Ampli	WTML	WTMR	slope*gain	rChi2%	offs1*g	slope1*g	rChi2%
#	0	199	90	10	1408.00	3.048	6.491	214459	5629.83	12.1	15148	2.432	1.869	1.000385	0.32	-0.050	1.000438	0.28

- For the **sum of segments** recalibration with ofset (**one gain and offset for all the segments, optional**)

```
RecalEnergy -spe Out_norecal/{crystalID}/Post__5-40-16384-UI__Ener.spec -sub 198 -num 1 -gain 4 -poly1 -152Eu
```

#	indx	#spec	#pks	#ok	rEnergy	FW05	FW01	Area	Position	Width	Ampli	WTML	WTMR	slope*gain	rChi2%	offs1*g	slope1*g	rChi2%
#	0	198	67	10	1408.17	5.674	14.718	179916	5639.15	20.0	6174	2.245	3.643	0.998854	7.87	0.454	0.998371	2.04

Add these coefficients to gen_conf.py

femul key in PostPSAFilter:
"RecalCC offset egain",
"RecalSG offset egain",

More details in [AGATA_LLQ_UsersGuide](#)

PostPSA Filter

2.3 Recalibration: Force Segments to core

3 steps process

Verification with femul replay each step

1. RecalEnergy2 (Check)
2. RecalCC and RecalSG (Check)
3. ForceSegmentstoCore (Check)[Optional]

- Energy of segments scaled in such a way that their sum equals Energy of Core
- Optional/Experiment dependent

What is needed:

- Source and In-beam data
- PostPSA file and Track files : Out/{crystalID}/**Post__5-40-16384-UI__Ener.spec**
Out/Merger/**Track__2-24-16384-UI__EC.spec**
Out/Merger/**Track__2-15-16384-UI__EE.spec**
- Conf File: **gen_conf.py**

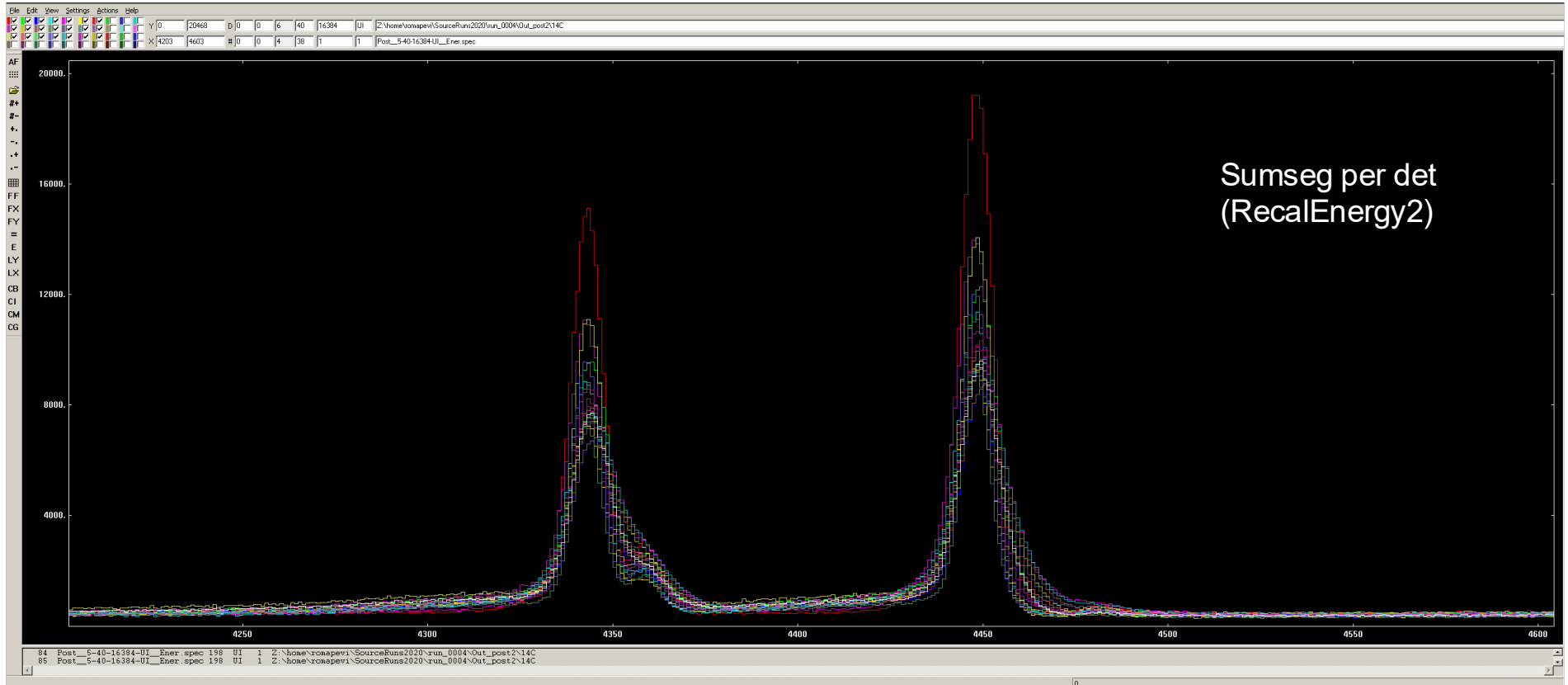
femul key in PostPSAFilter:
"ForceSegmentstoCore",
Add it in gen_conf.py

PostPSA Filter

Verification with femul replay each step

2. Recalibration

Post__5-40-16384-UI__Ener.spec [4] [38]

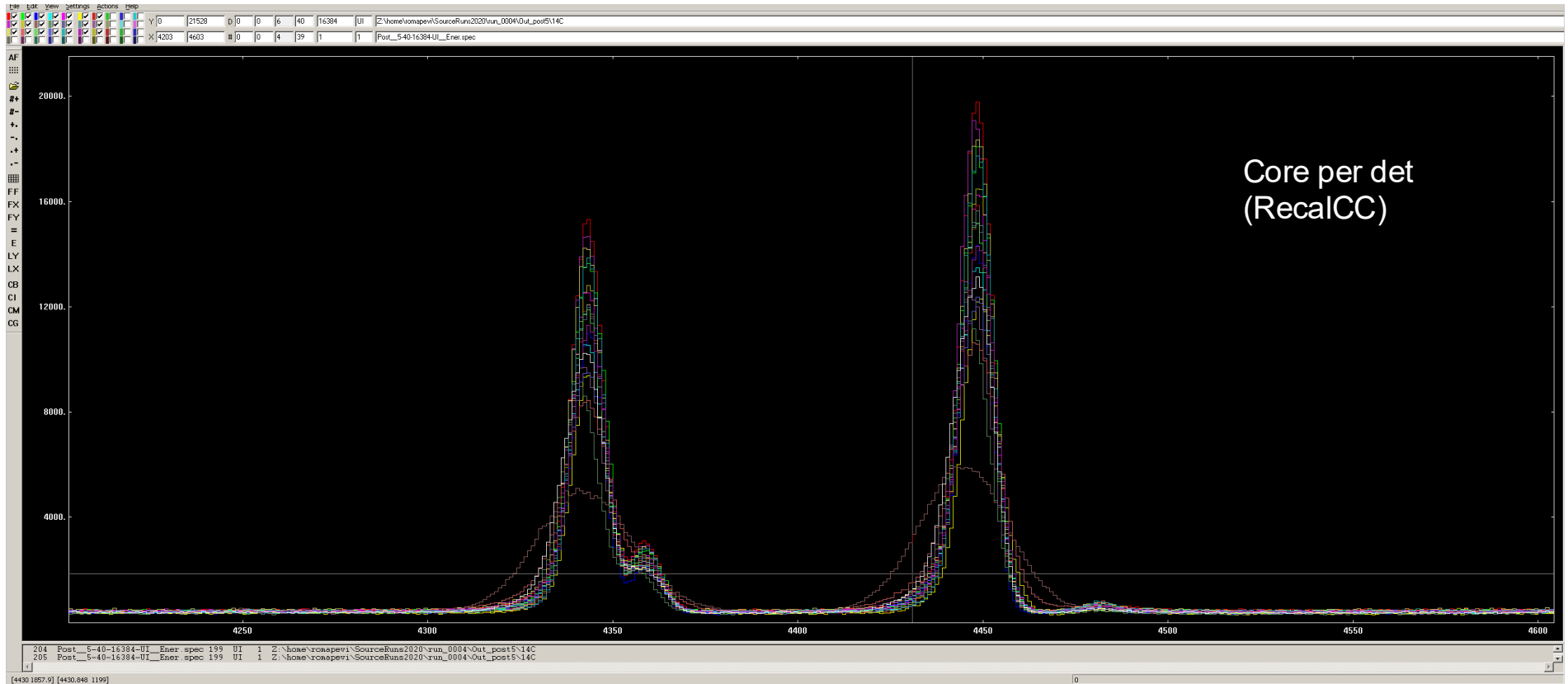


PostPSA Filter

Verification with femul replay each step

2. Recalibration

Post__5-40-16384-UI__Ener.spec [4] [39]

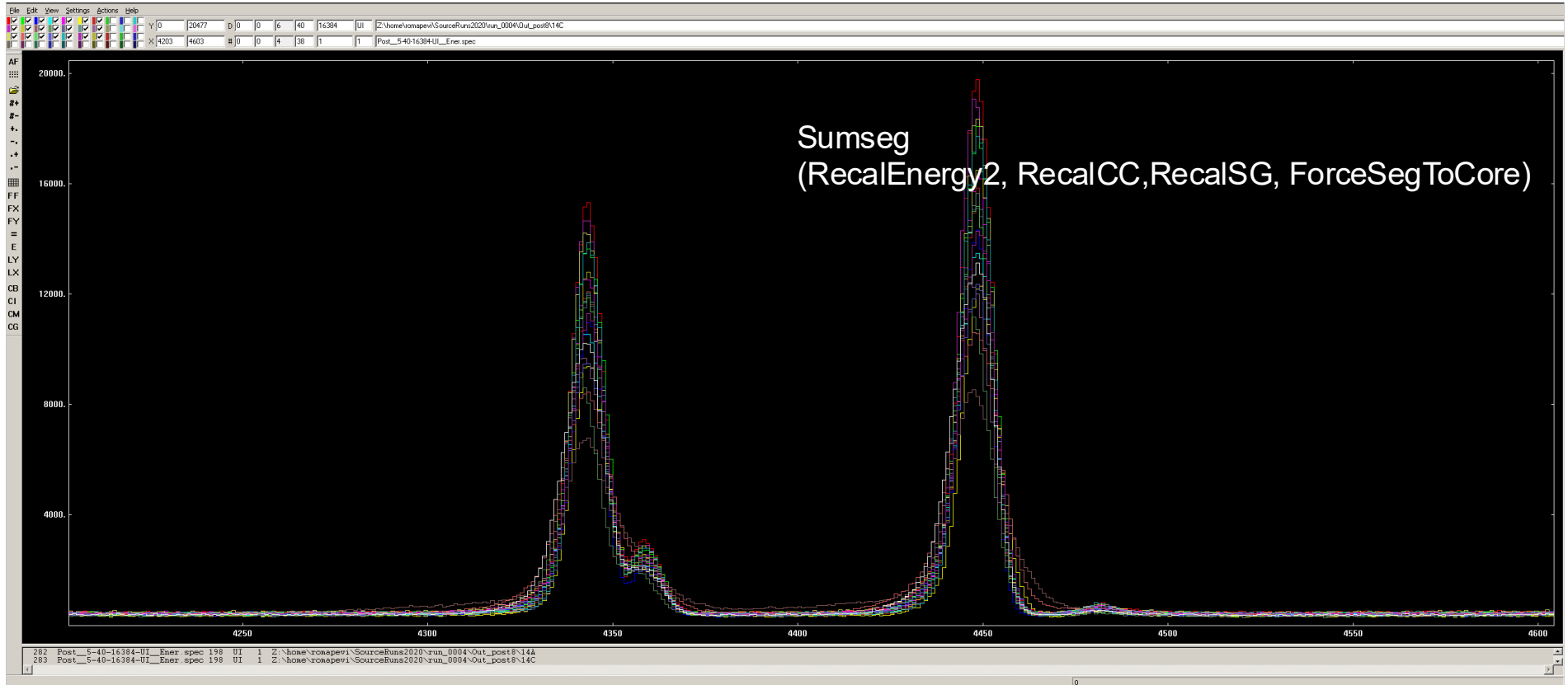


PostPSA Filter

Verification with femul replay each step

2. Recalibration

Post__5-40-16384-UI__Ener.spec [4] [38]



PostPSA Filter

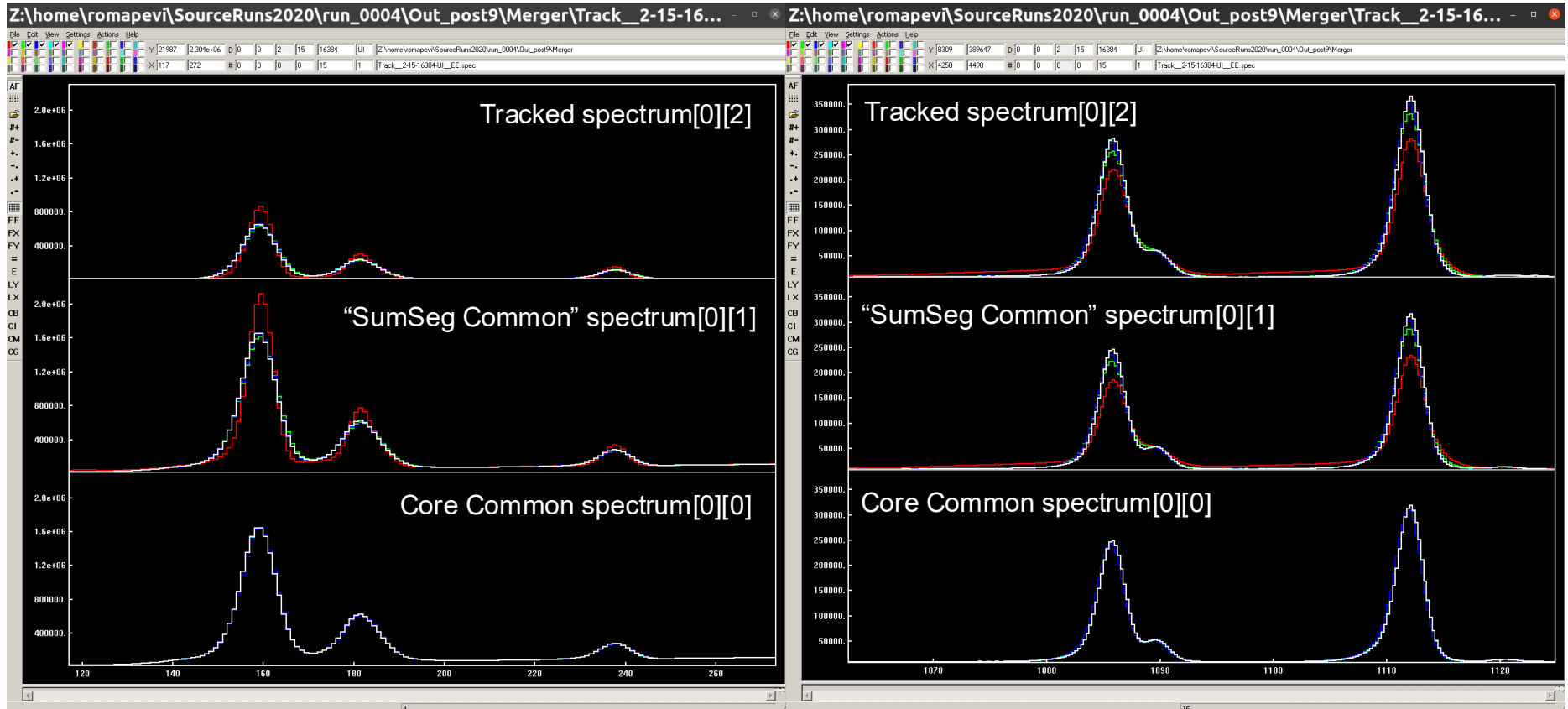
2. Recalibration

Track__2-15-16384-UI__EE.spec [0][0-2]

No ForceSeg, RecalEnergy2

ForceSeg, RecalEnergy2, RecalCC RecalSG

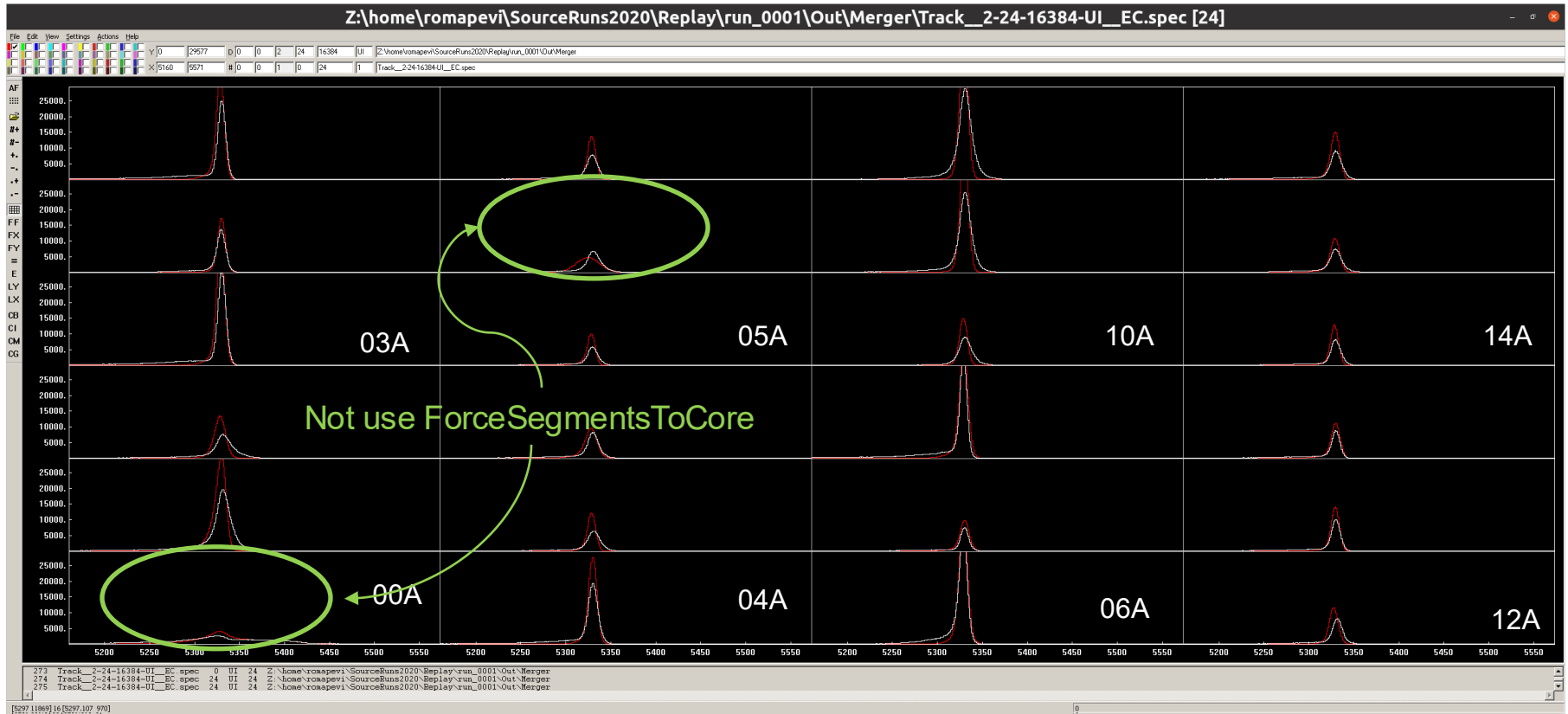
ForceSeg, RecalEnergy2 no offset, RecalCC RecalSG



PostPSA Filter

2. Recalibration

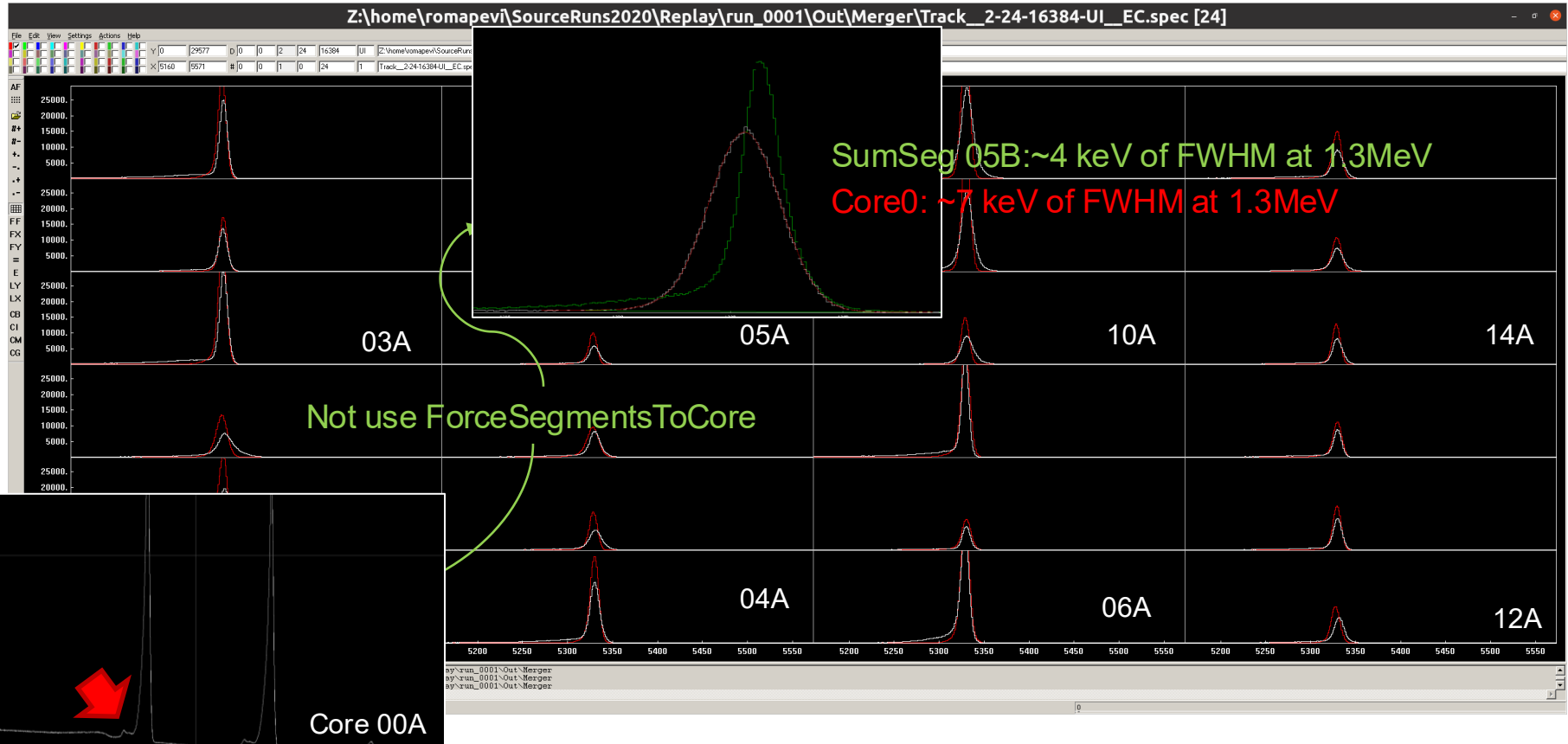
Track__2-24-16384-UI__EC.spec



PostPSA Filter

2. Recalibration

Track__2-24-16384-UI__EC.spec



PostPSA Filter

3. Global time alignment

To be done every time that there is a GTS alignment

2 steps process

1. Replay without coeff
2. Replay with coeff

What is needed:

- Any run
- Spectra file : Data/Merger/Track__35-35-1000-UI__TT.spec
- Conf File: **gen_conf.py**
- Auxiliary files: recalT.dat
- Programs/scripts:

RecalEnergy: generate shift coefficients for the N*N time spectra

```
RecalEnergy -spe ../Out/Merger/Track__${N}-${N}-1000-UI__TT.spec -T 500 -num ${NN} | tee recalT.dat
```

#	indx	#spec	#pks	#ok	rEnergy	FW05	FW01	Area	Position	Width	Ampli	WTML	WTMR	shift*gain
#	0	0	1	1	0.00	0.000	0.000	0	0.00	0.0	0	0.000	0.000	0.000
	1	1	1	1	503.37	19.629	44.382	9422	500.85	19.5	400	2.382	2.171	-0.850
	2	2	1	1	501.04	16.180	36.094	11042	500.47	15.8	574	2.604	1.955	-0.473
	3	3	1	1	502.67	16.667	36.949	10726	499.84	16.4	543	2.558	1.958	0.164
	4	4	1	1	503.28	17.366	40.384	8202	499.65	17.1	388	2.484	2.231	0.350
	...													

solveTT.py: find the best shift combination.

```
${pathSoftware}/solveTT.py -f recalT.dat -n ${N} -c 13 -p 500
```

Shifts that minimize Chi2

```
0.001  
-0.181  
0.004  
-0.087  
0.040  
-0.239  
...
```

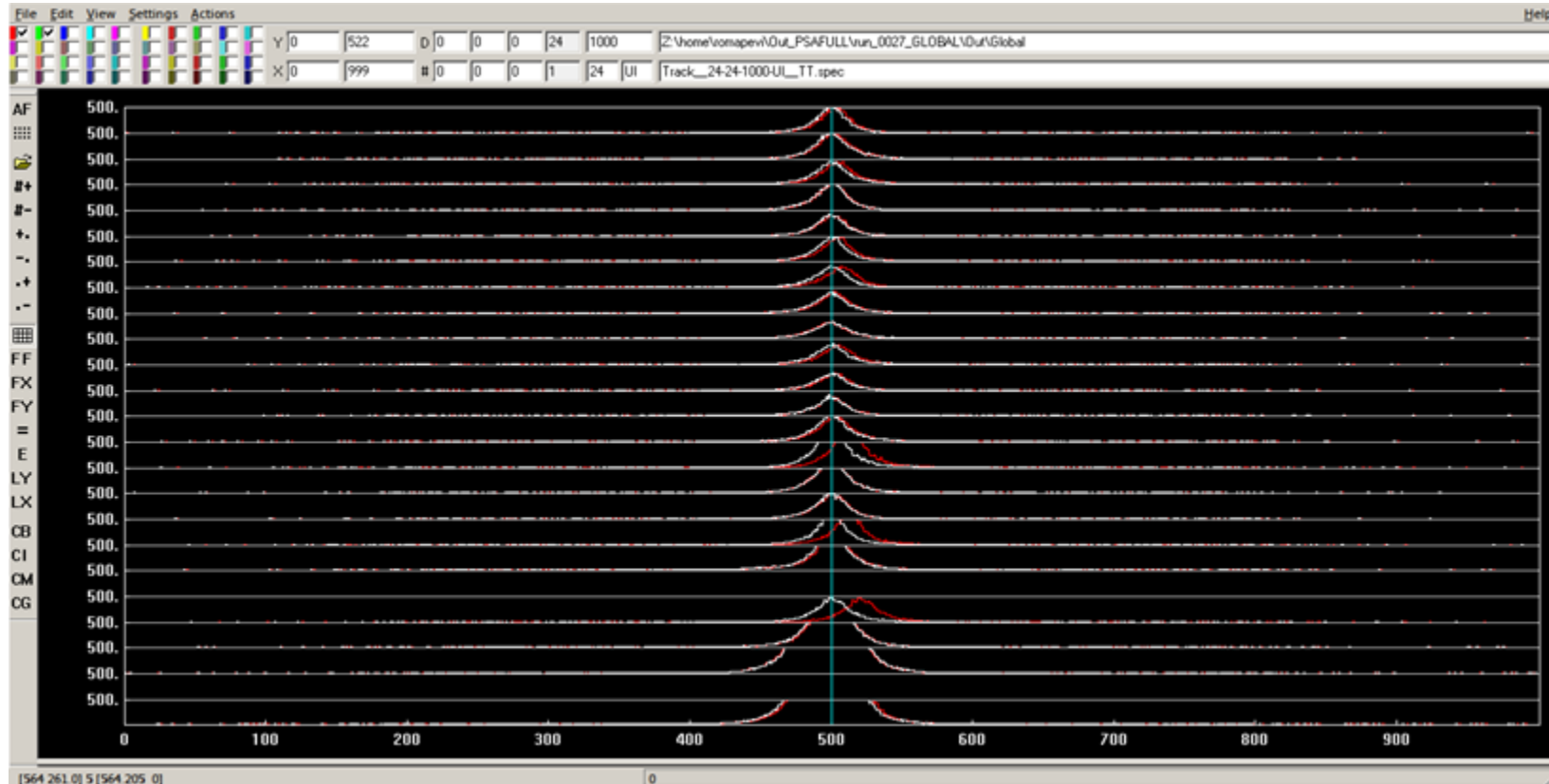
femul key in PostPSAFilter:
"TimeShiftCC coeff",

PostPSA Filter

Verification with femul replay

3. Global Time Alignment

Track__24-24-1000-UI__TT.spec example detector 00B [0][all] red before, white after time alignment

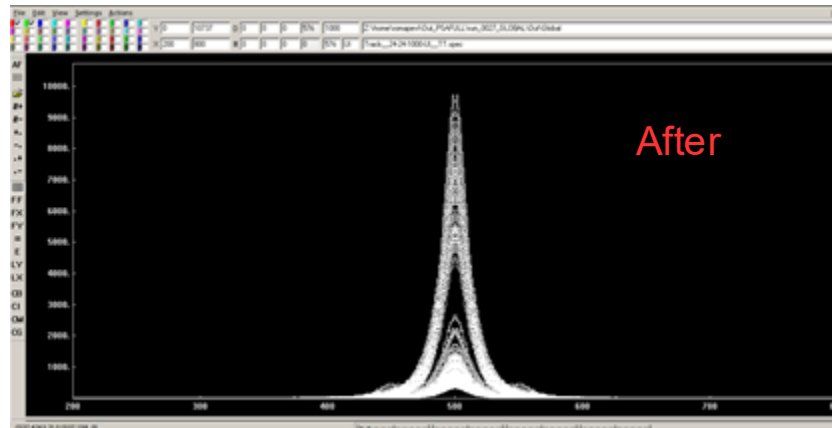
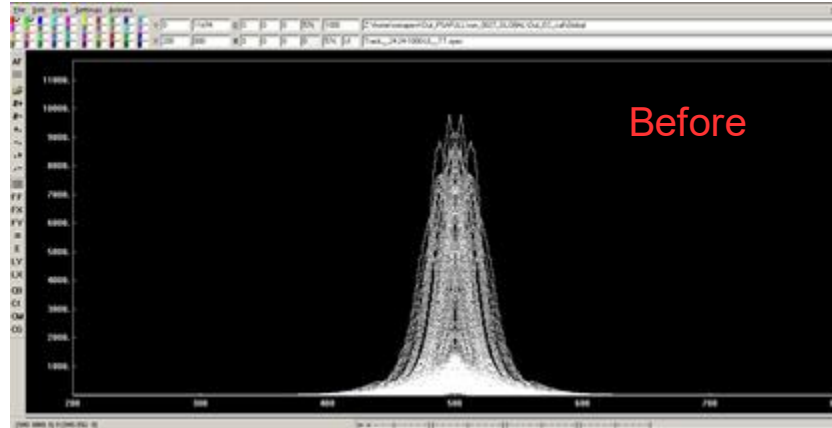


PostPSA Filter

Verification with femul replay

3. Global Time Alignment

Track__24-24-1000-UI__TT.spec all [all][all]

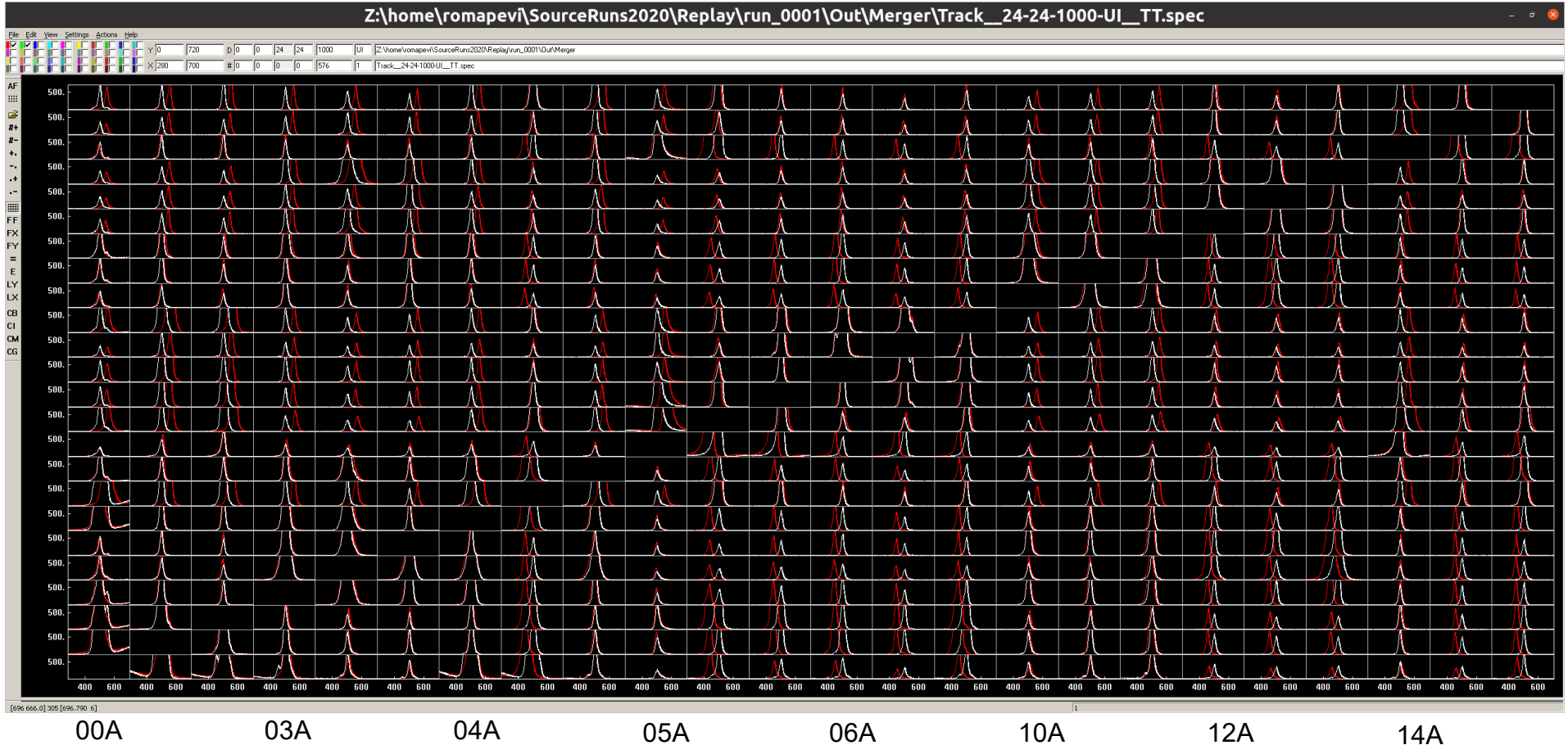


PostPSA Filter

Verification with femul replay

3. Global Time Alignment

. Track__24-24-1000-UI__TT.spec all [all][all] red before, white after time alignment



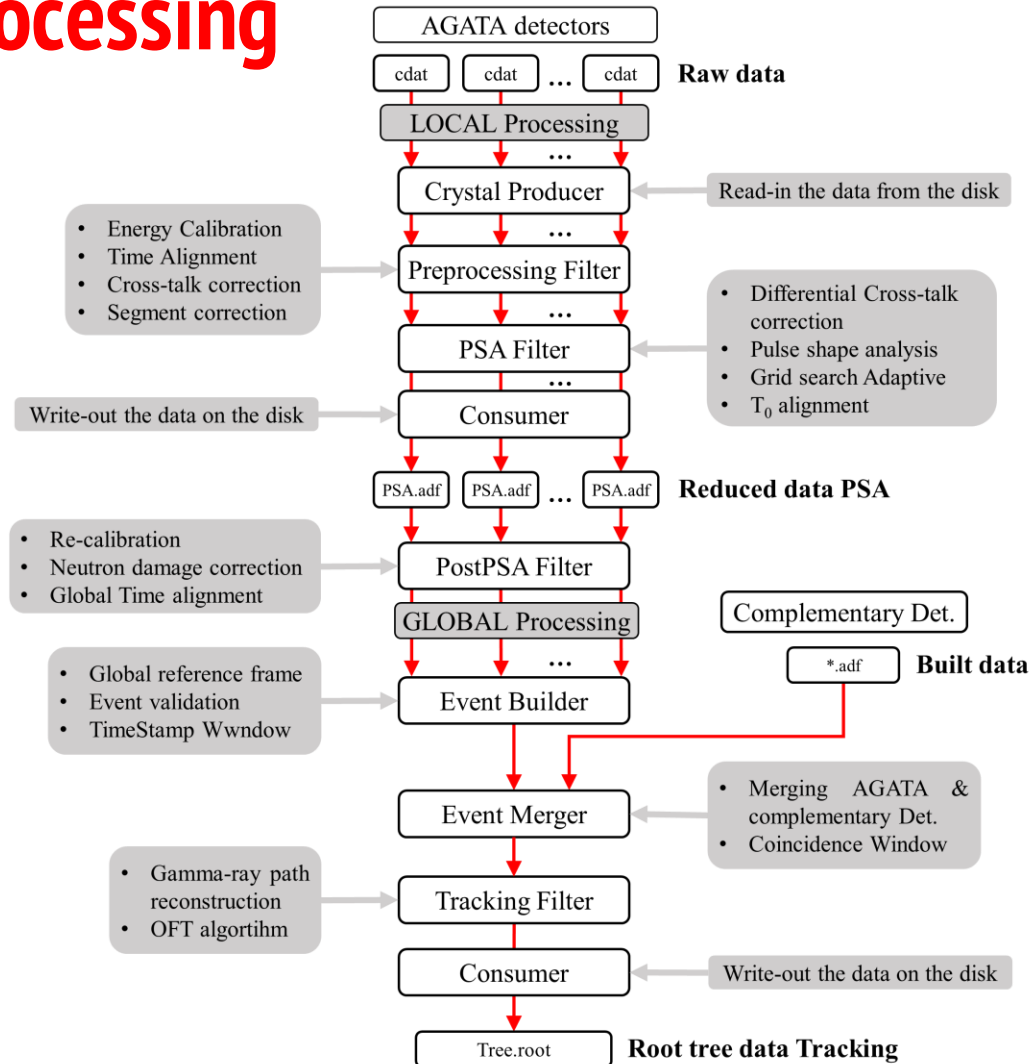
Local Level Processing

PostPSA Filter

- Follow the order given here.
- Be careful of the possible redundant calibration done by the PostPSA filter actor.
- The PostPSA is the last chance to have properly calibrated segments.
- The calibration offset can only be set at this level of the analysis.
- ForceSegToCore! final correction, only when the core resolution is good.
- Keep track of the GTS alignment for the Global Time alignment (important to reduce random coincidences)

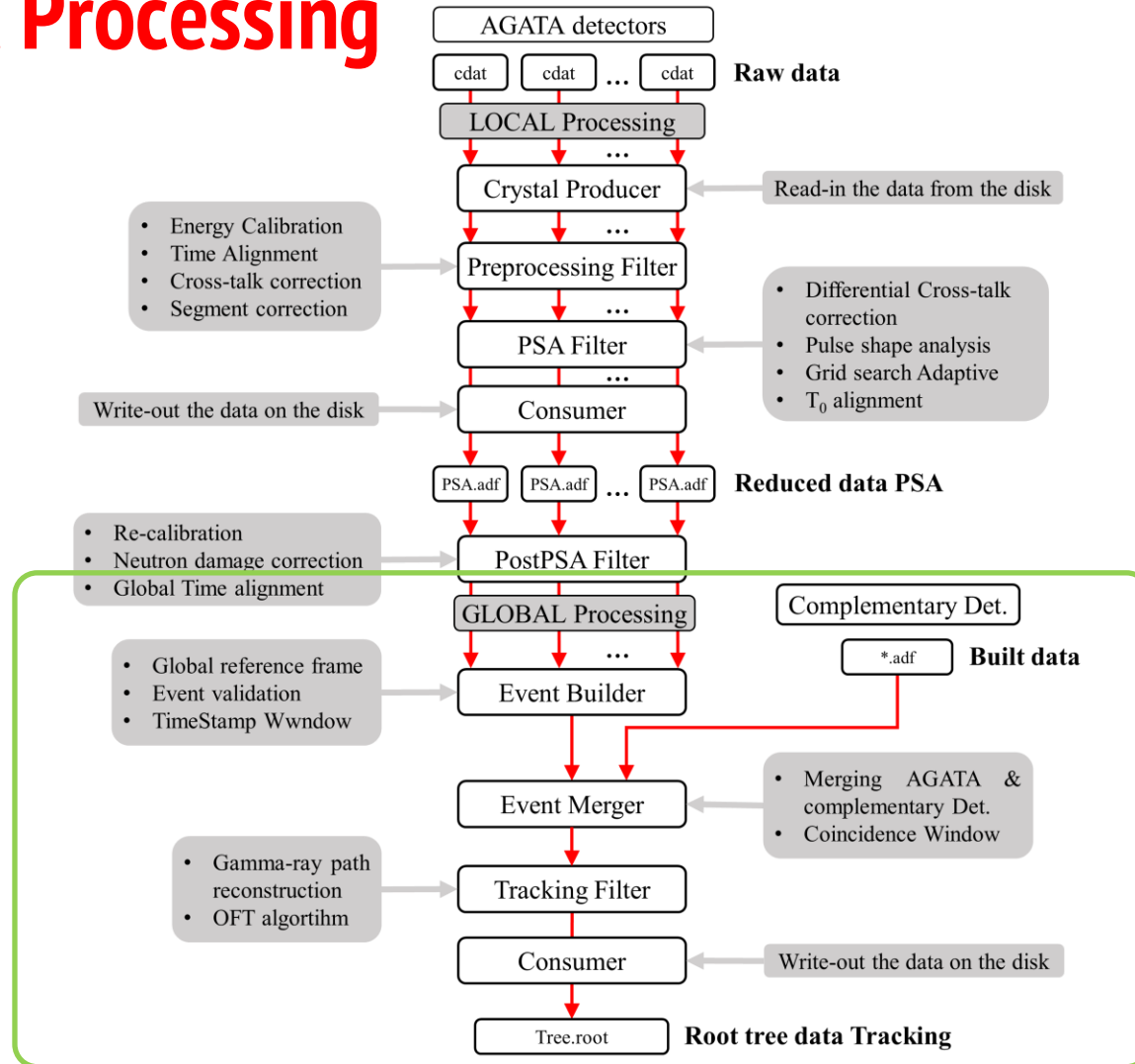
Local Level Processing

Narval actors



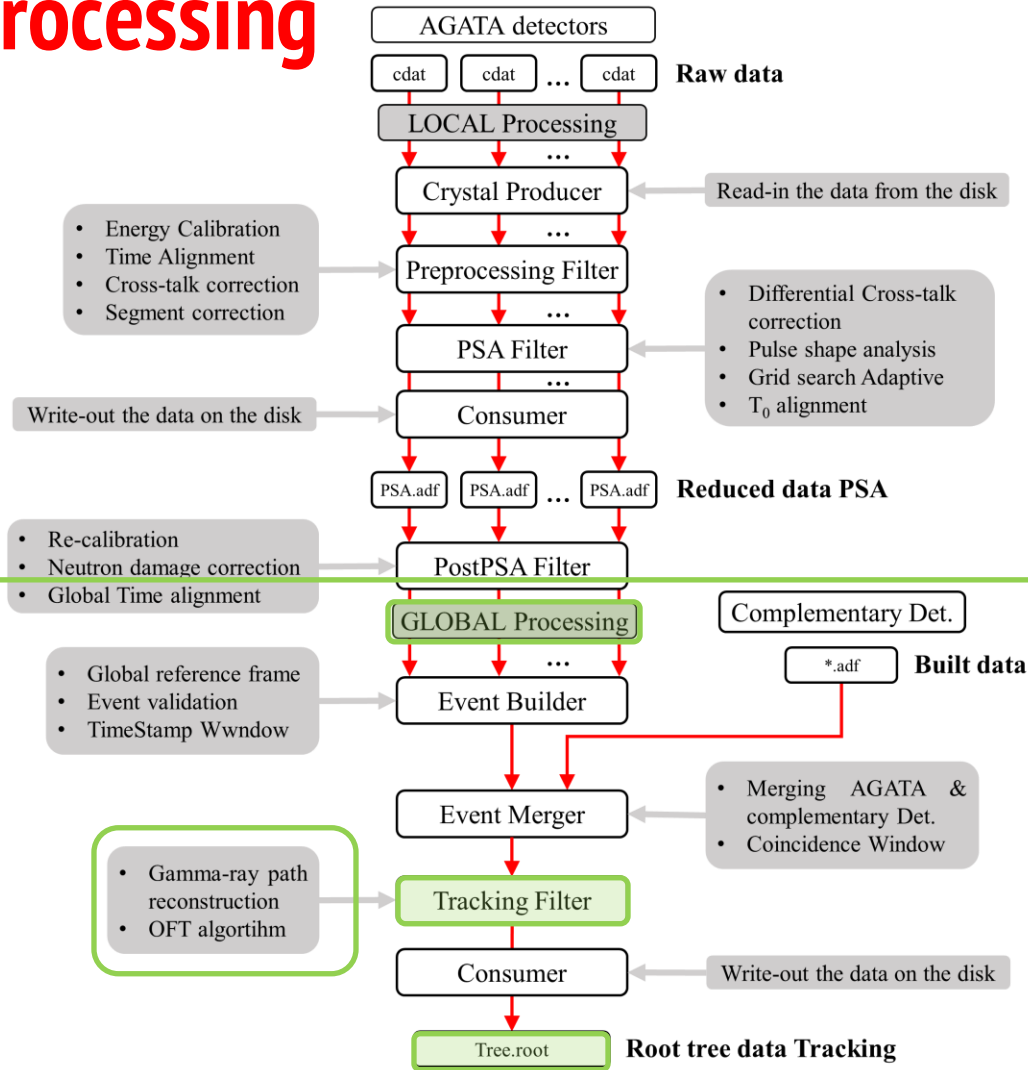
Global Level Processing

Narval actors



Global Level Processing

Narval actors



Global data replay

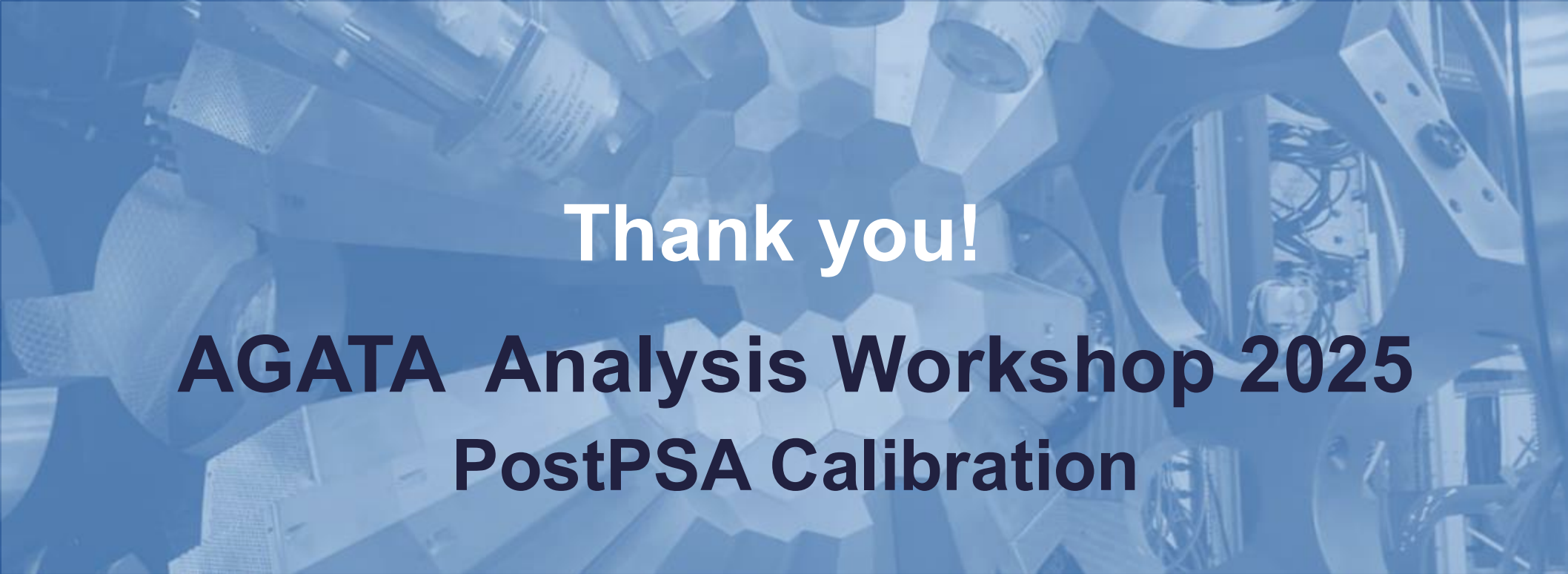
Speaker: **Jérémie Dudouet**
Wednesday

Tracking

Speaker: **Araceli López Martens**
Friday

Selector

Speaker: **Daniele Brugnara**
Wednesday



Thank you!

AGATA Analysis Workshop 2025
PostPSA Calibration

R.M. Pérez-Vidal

14/01/2025, Lyon

Questions?