

AGATA Local Level Processing (adapted from E. Clément and D. Bazzacco)

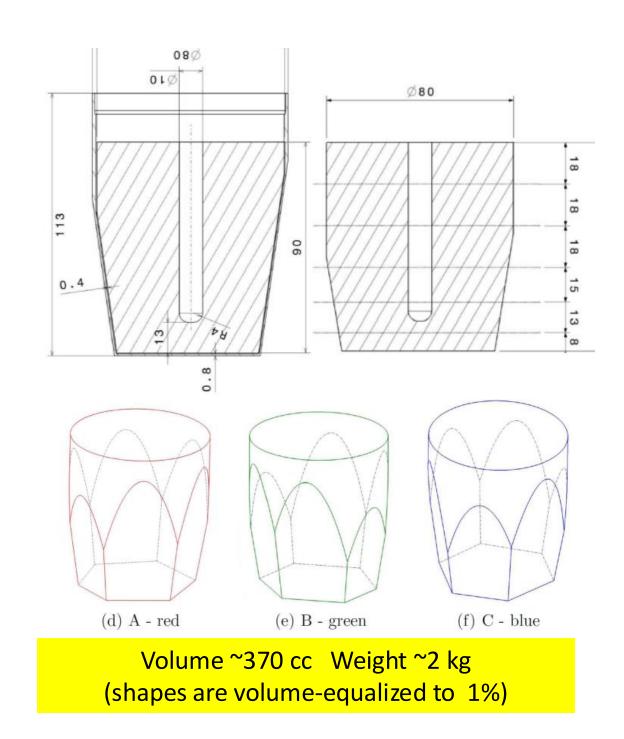
Jérémie Dudouet
on behalf of the Data Analysis Working Group

AGATA Analysis School: 13/01/2025, Lyon

Topics

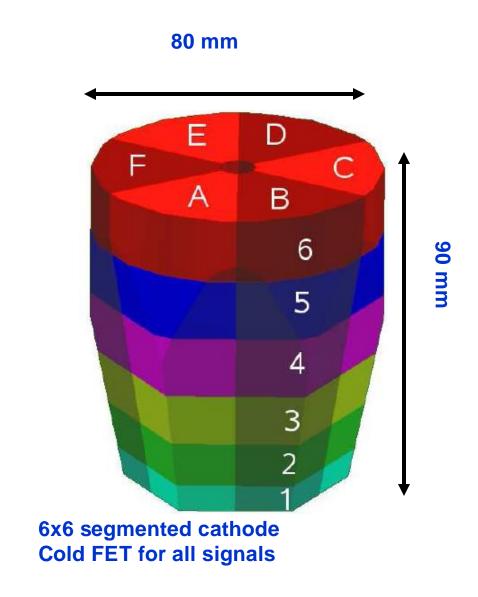
- General structure of data acquisition
- Data flow **ADF**
- Data processing model
- Emulators
- Narval (ADA/C/C++) and actors

AGATA detectors





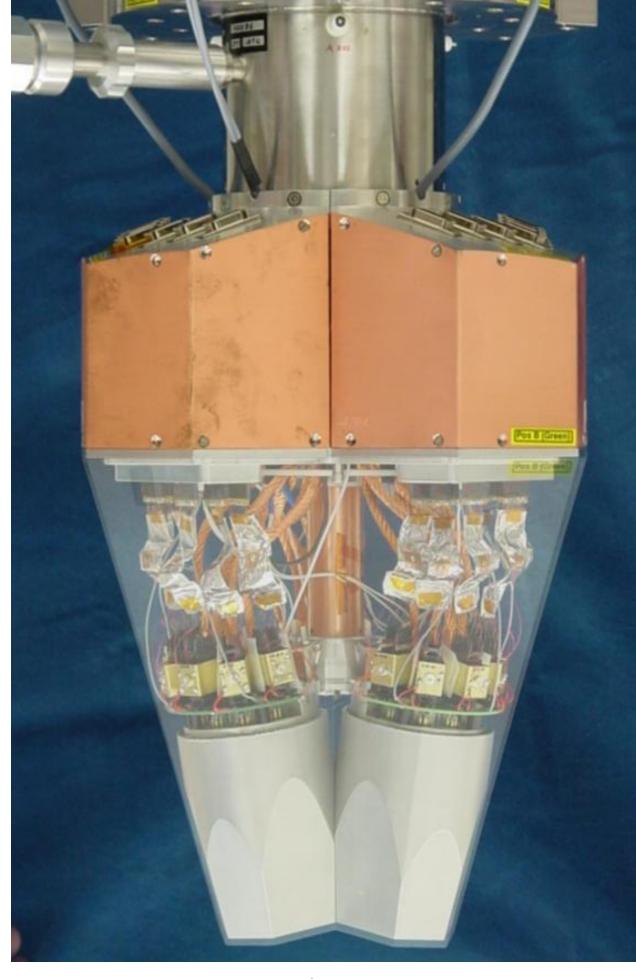




Energy resolution
Core: 2.35 keV
Segments: 2.10 keV
(FWHM @ 1332 keV)

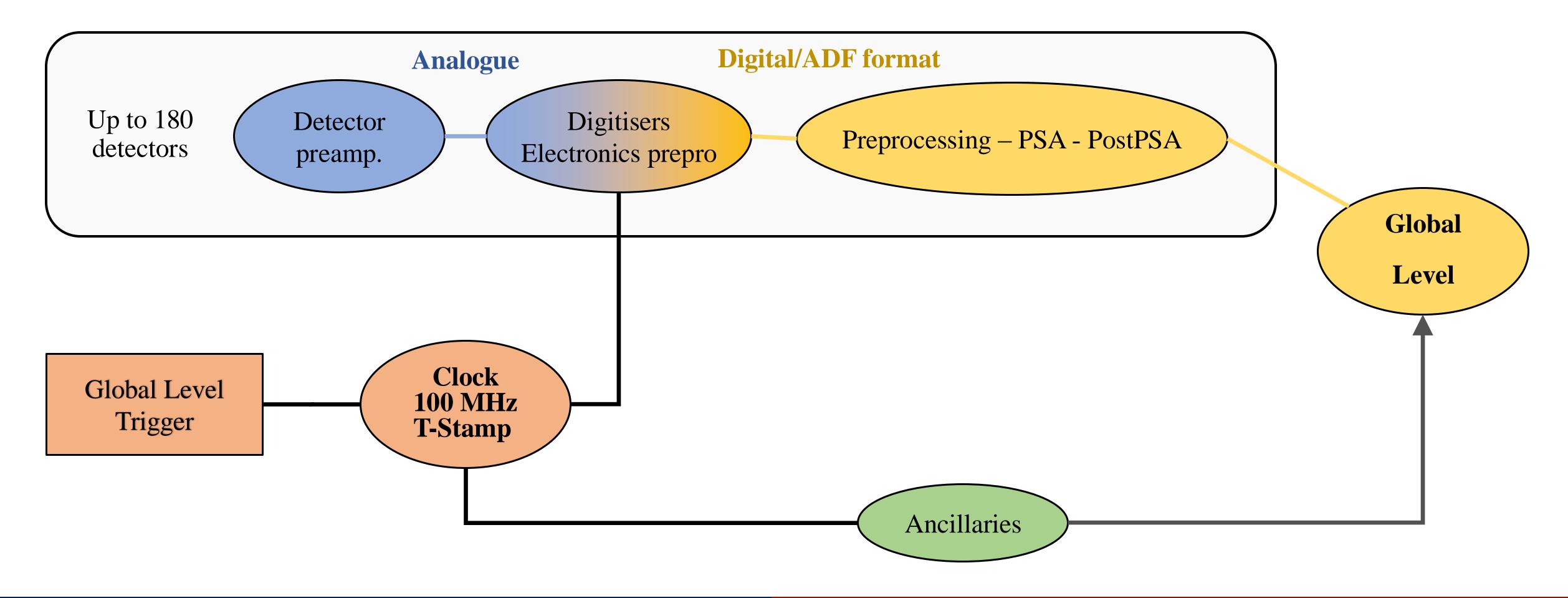
A. Wiens et al. NIM A 618 (2010) 223D. Lersch et al. NIM A 640(2011) 133

38 high-resolution signals / detector



AGATA Triple/Double Cryostat

Structure of Electronics and DAQ



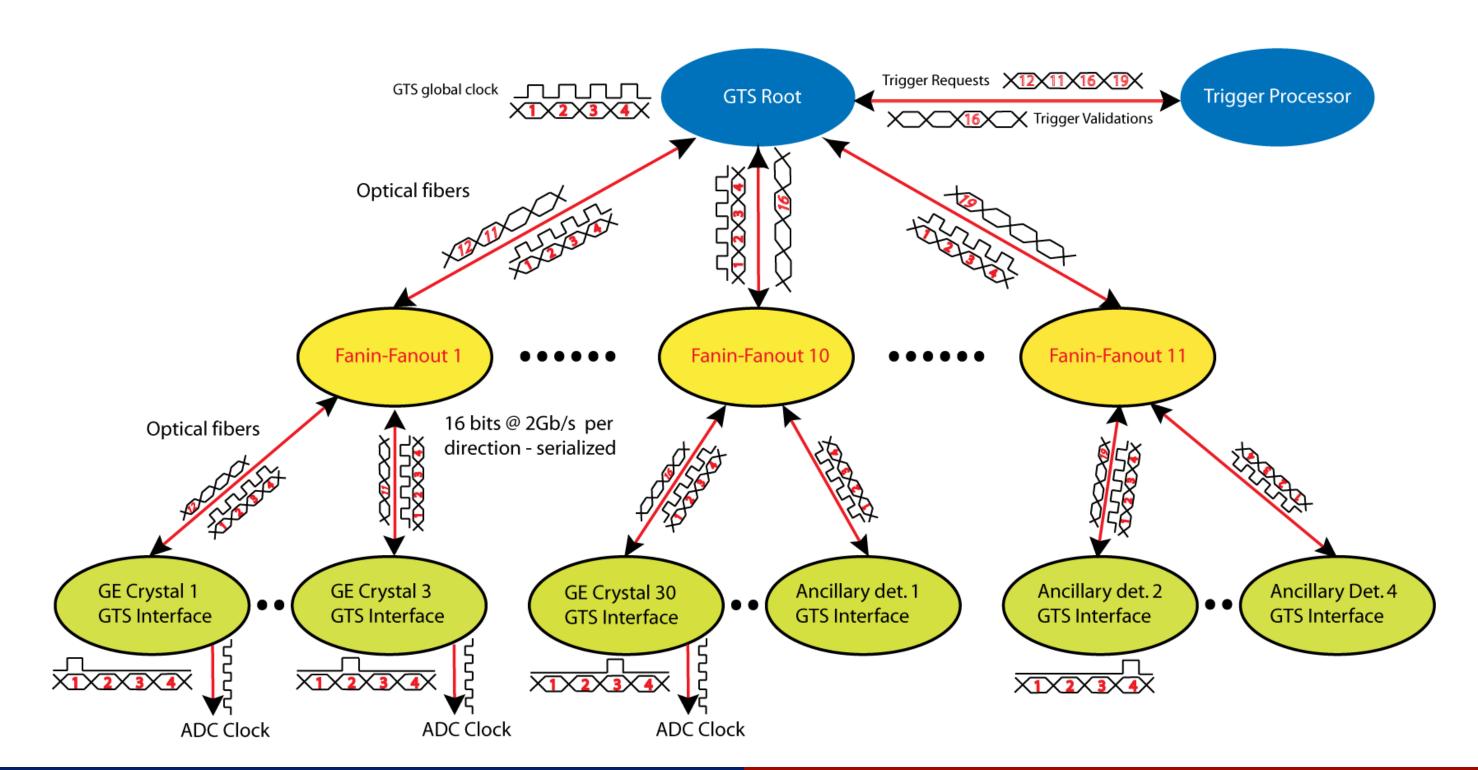
GTS: the system coordinator

All detectors operated on the same 100 MHz clock

Upwards: Trigger requests, consisting of address (8 bit) and timestamp (16 bit)

max request rate 10 MHz total, 1 MHz/detector

Downwards: Validations/rejections, consisting of 48 bit timestamp + 24 bit event number



The AGATA Data Format: ADF

ADF Frame:



The key contains:

- Data lenght
- Data type
- Timestamp
- Event number

The Data contains either:

- data (energies, hits...)
- adf frames

The AGATA Data Format: ADF

The ADF.conf configuration file

Frame: Agata Frame: Agata Frame: Agata Frame: Agata	<pre>data:crystal data:ccrystal data:ranc0 data:ranc1 data:ranc2 data:psa</pre>	4 4 4 4	0 0 0 0 0	Agata Agata Agata Agata	<pre>data:crystal data:ccrystal data:ranc0 data:ranc1 data:ranc2 data:psa</pre>	65 65	5000 5000 5000	0 0
Frame: Agata Agata Agata	data:tracked event:data event:ranc event:data:crystal event:data:crystal event:data:psa event:data:ranc0 event:data:ranc1 event:data:ranc2 meta:vertex meta:sync meta:scan	4 4 4 4 4 4 4 4	000000000000000000000000000000000000000	Agata Agata Agata Agata Agata Agata Agata Agata Agata Agata	data:tracked event:data event:ranc event:data:crystal	1 4 4 4	0 0 0 0 0 0 0 0 0 0	

Each key and frame is associated to a version number

Architecture of the system

Local Level: where the individual detectors don't know of each other.

- Electronics and computing follow a model with minimum coupling among the individual detectors, which are operated independently as long as possible
- Electronics is almost completely digital, operated on the same 100 MHz clock
- Data processing (in the electronics and in the front-end computers) is the same for all detectors and proceeds in parallel
- Every chunk of produced data is tagged with a 48 bit number (time stamp) giving the absolute time (with a precision of 10 ns) since the last hard-reset of the system → roll around takes place every 32.5 days.

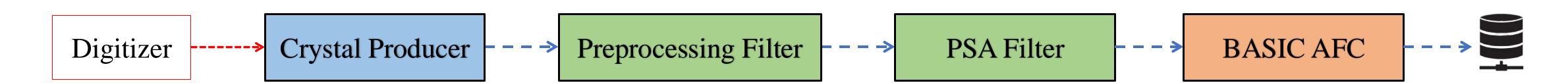
Global Level: where the detectors do know of each other

- By means of the real time trigger
- Via the event builder and merger that assemble the event fragments (including the ancillaries) into complete events that are further processed
- In the Tracking and in the Physical analysis stages

Data processing: NARVAL/DCOD

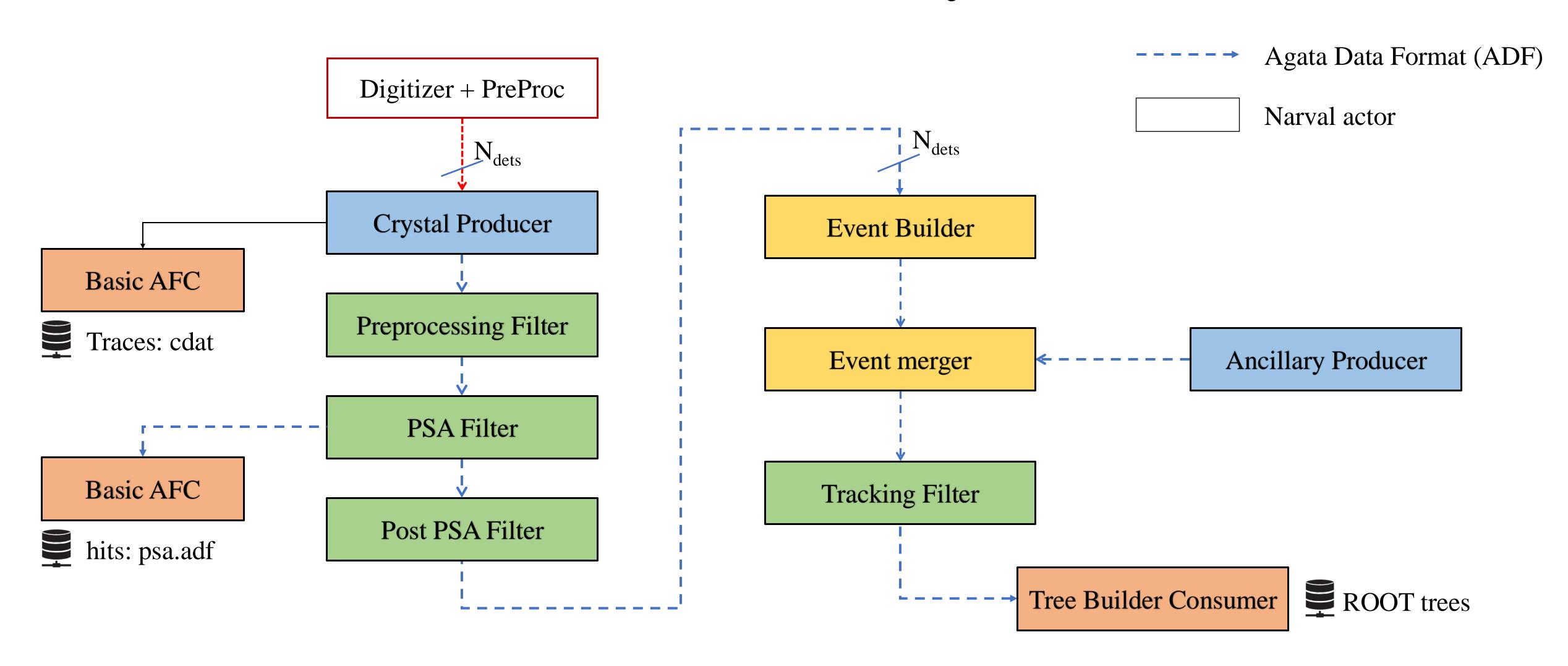
List of actors, applying successive operations on the data flow:

- Producer: Read data from outside (electronics/disk) to the NARVAL chain
- Filters: Process operations on the dataflow (calibrations/PSA/Tracking)
- Dispatchers: Merge multiple entries in the NARVAL chain (Event Builder/Event Merger)
- Consumer: Close the NARVAL chain (write data on disk)

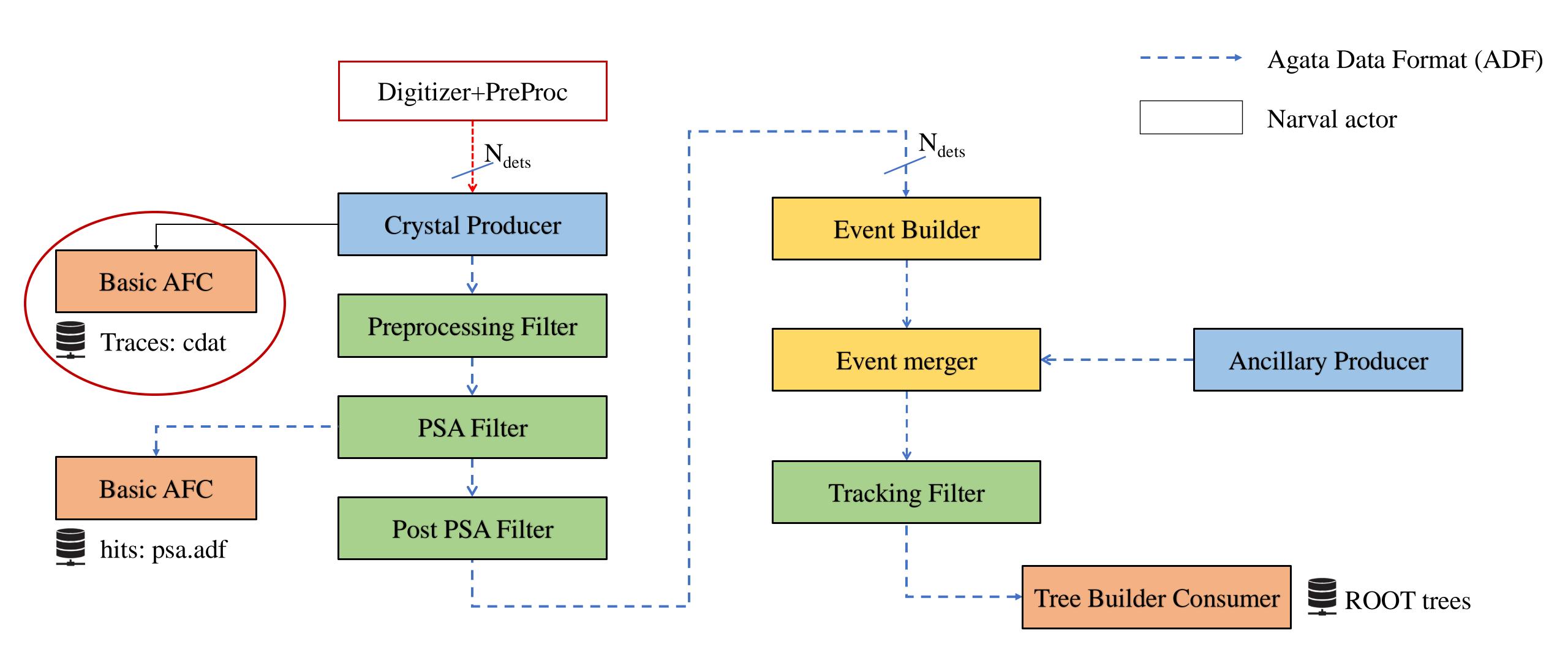


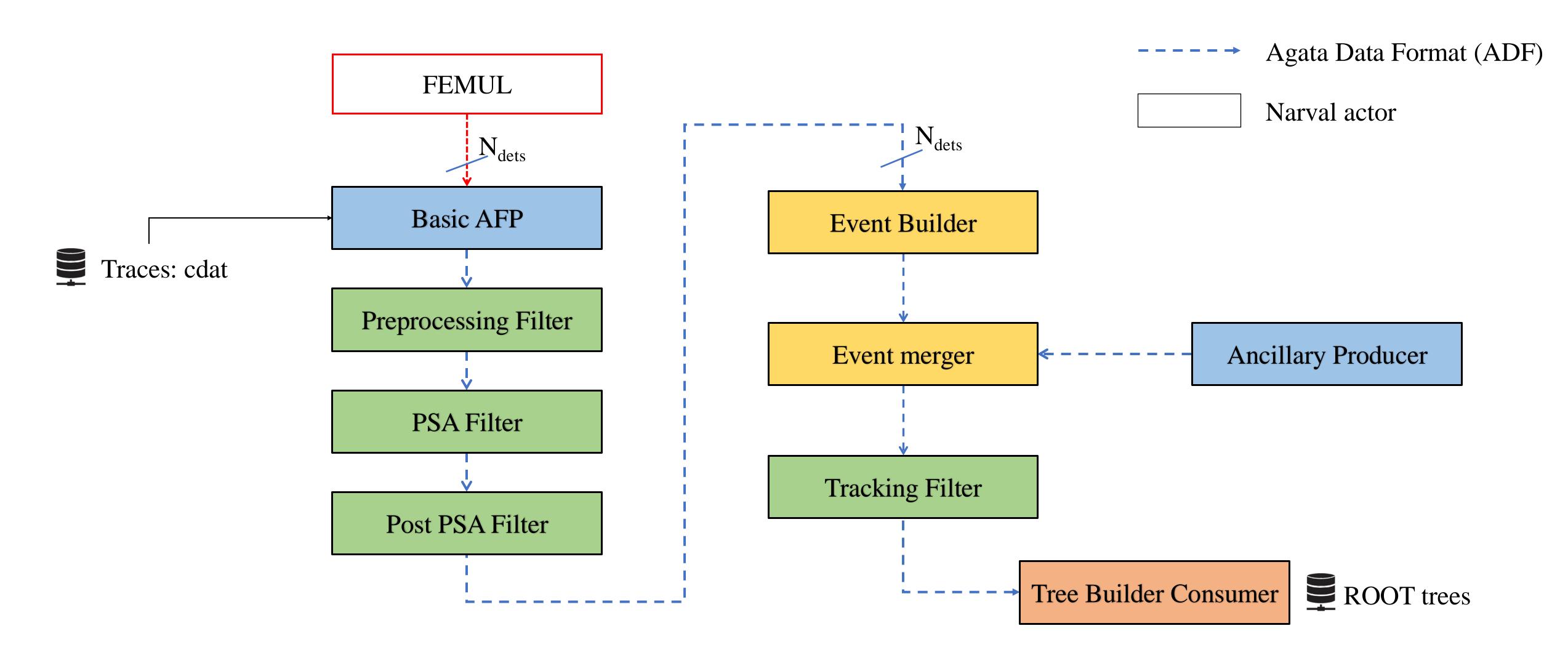
Standard actors chain for AGATA LLP

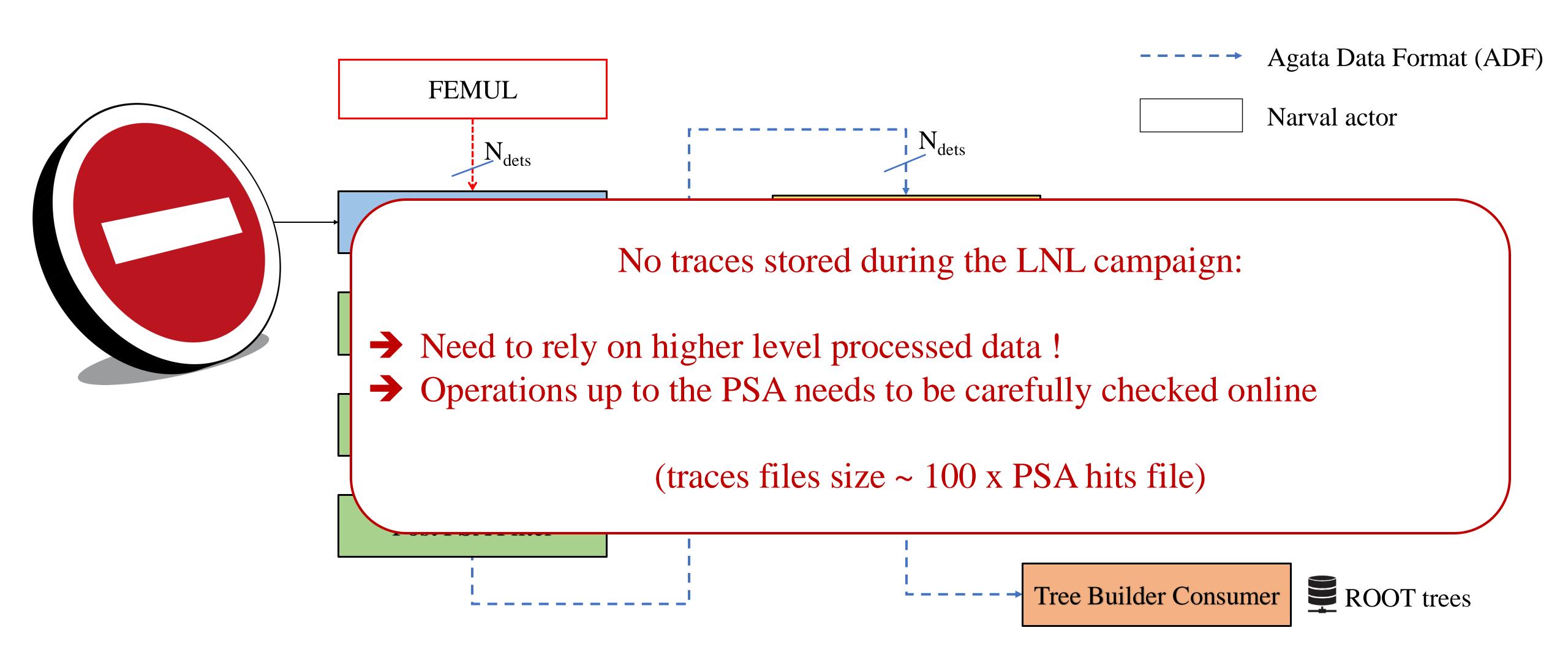
Architecture of the system

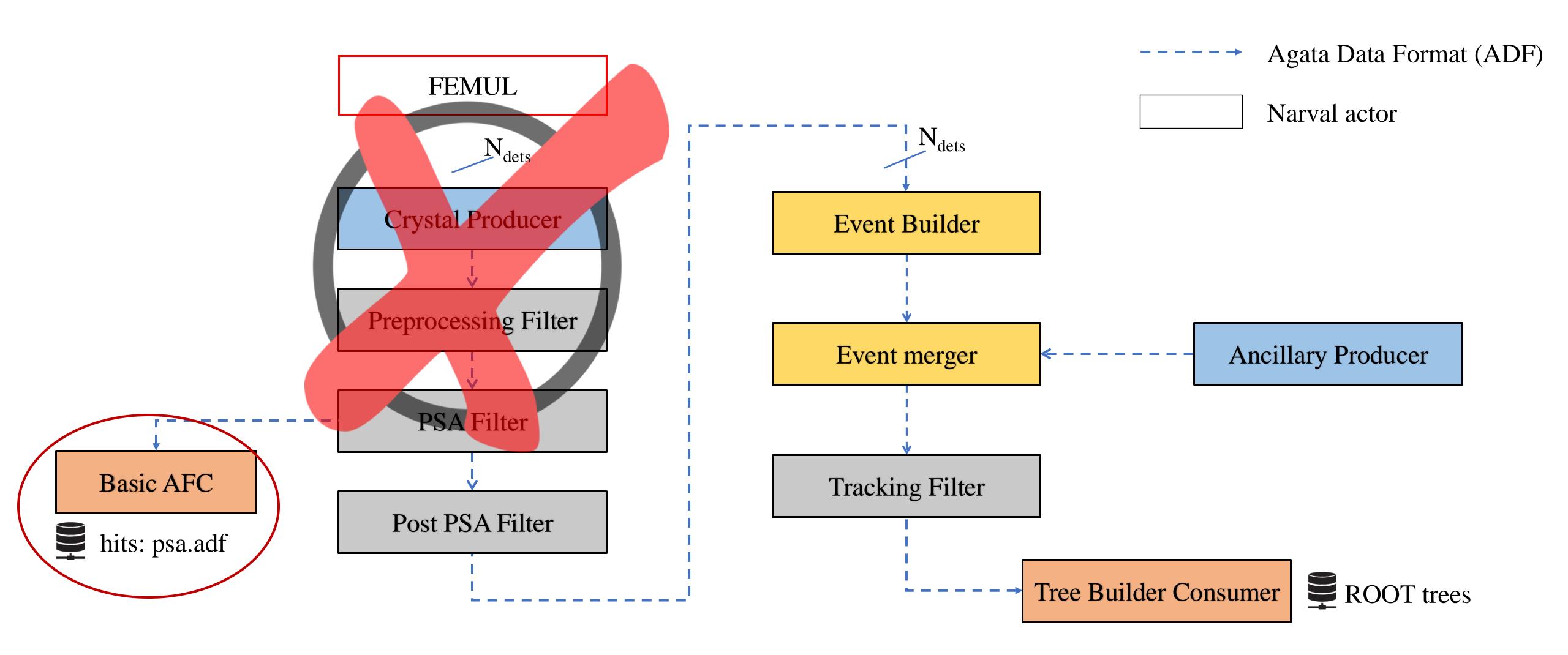


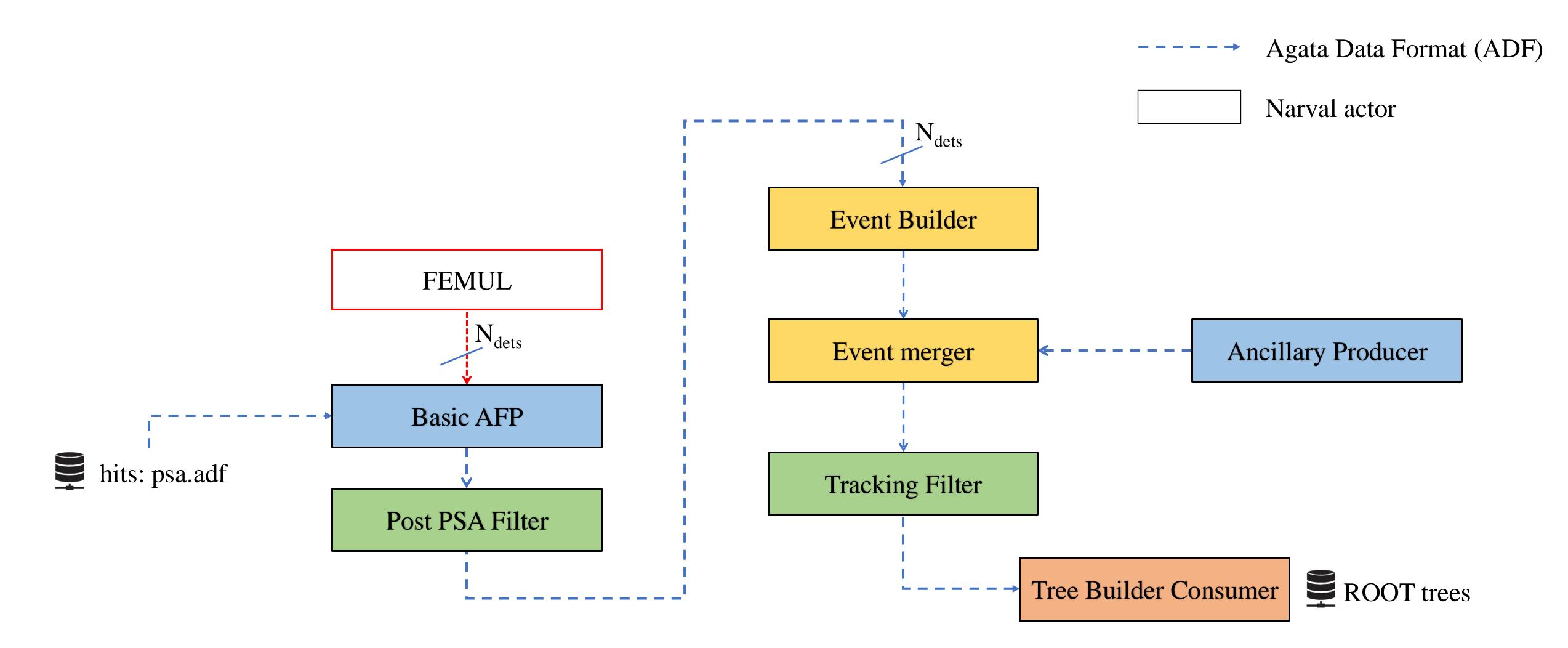
- Originally intended to help developing and debugging the user libraries which is very hard to do in a distributed processing environment like Narval.
- Gradually developed as a full emulation of the Narval framework with the limitation of being a single process running a specific machine (threads used to distribute the work on the available cores).
- Configuration (detectors, actors, ..) specified by a "topology" file.
- Generation of configuration parameters via gen_conf.py
- **→** Online == Offline processing











Structure of analysis directories

The directory where you produce your data contains some standard sub-directories

(e.g. /agatadisks/exptname (EXXXX) /(Config EXXXX)/run_XXXX_date)

Conf: Configuration of actors, calibrations, ... for each detector

→ 00A, 00B, 00C ... Ancillaries, Global, Merger

with minimal differences between online and offline

Data: Data and spectra produced during the experiment

→ Online writes data here

→ Offline replay takes data from here

Out: Data and spectra produced during data replay

→ Offline writes data here

Typical configuration directories

Conf/12A

- CrystalProducer.conf
- CrystalProducerATCA.conf
- PreprocessingFilter.conf
- PreprocessingFilterPSA.conf
- PSAFilter.conf
- PostPSAFilter.conf
- xdir_1325-1340.cal
- xinv_1325-1340.cal
- BasicAFC.conf
- BasicAFP.conf

Conf/Builder

• EventBuilder.conf

Conf/Merger

- EventMerger.conf
- TrackingFilter.conf
- CrystalPositionLookUpTable
- TreeBuilder.conf

Binary spectra

- Simple C-style multidimensional (max 6) arrays written mostly in binary format
- For historical reasons the format is not recorded in the file
 - → Often written as part of the file name:
 Prod__4-38-32768-UI__Ampli.spec is a file dump of an array defined as:
 unsigned integer Ampli[4][38][32768],
 containing 4*38 = 152 spectra of 32768 channels
- The viewers TkT and Mat can decode and interpret the format.
- Other programs (e.g. RecalEnergy) can interpret the spectrum length and type but the user have to specify the number of spectra to act upon.

Some Useful programs

The number of channels (38 x number of detectors) to be calibrated and checked at each analysis level is too large to be done one by one: **automatic tools and procedures are distributed**

- → TkT spectrum viewer: to plot any spectrum produced all along the chain of actors
- → RecalEnergy: Analysis of spectra looking for peaks (energy and time calibrations)
- → SortPsaHits: Sort of PSA hits (special format) to determine neutron damage correction parameters
- → gen_conf.py: Unified procedure to produce configuration files for all actors
- → solveTT.py: Optimize time alignment of "equal" detectors (Global time alignment)

Local actors



CrystalProducer

- Readout of electronics, or get raw data from file
- Save original data to be able to replay experiment



PreprocessingFilter

- Energy calibrations (no offset)
- local time alignment of segments and core
- Cross talk correction
- Amplitude calibration and time alignment of traces
- Improved pile-up rejection



PSAFilter

- Decomposition of calibrated experimental traces by comparison with a calculated signal basis
- In principle more than one algorithm available but only one used in practice



PostPSAFilter

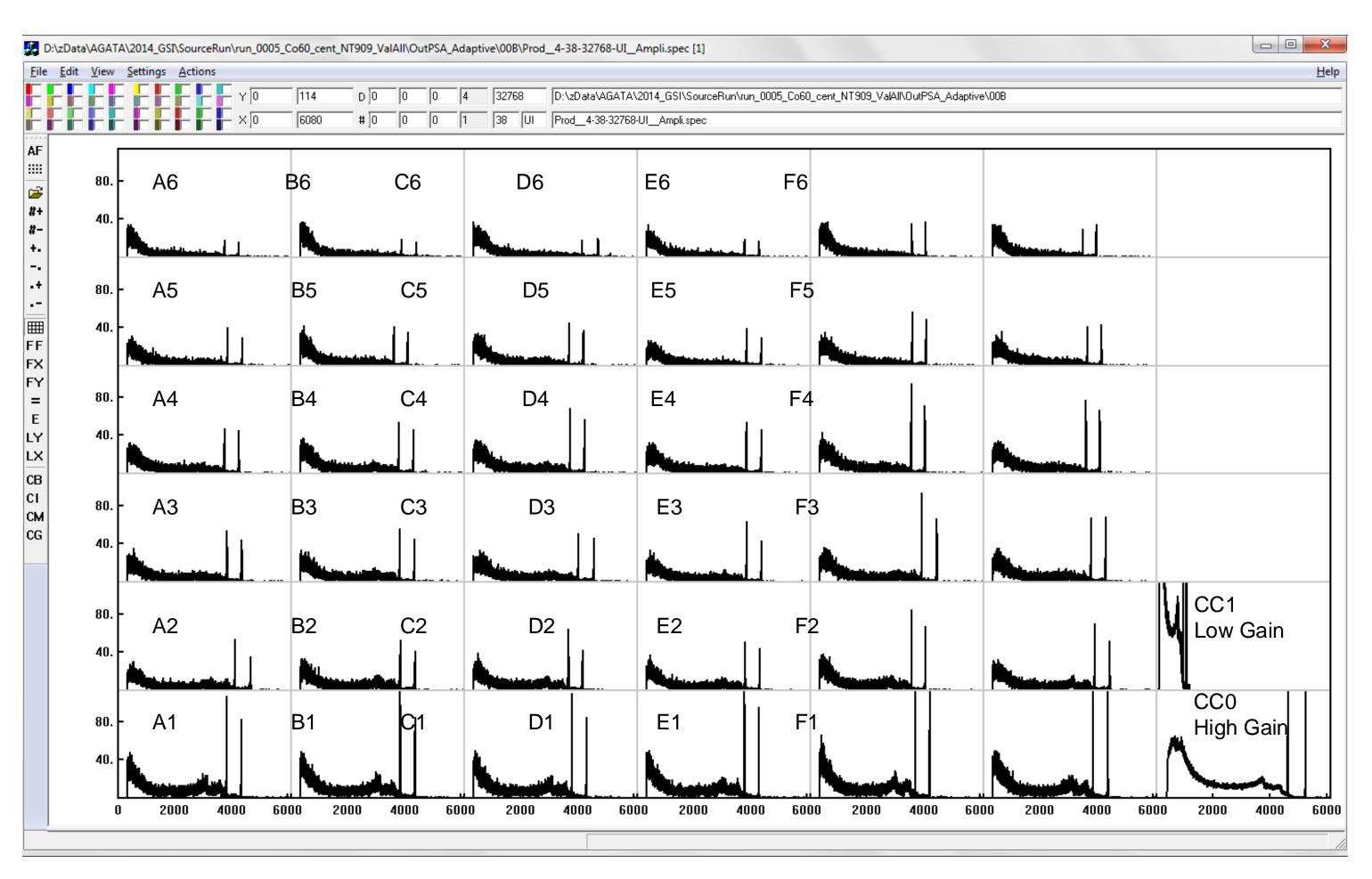
- Neutron deficit corrections
- Recalibrations of energy and time (including offsets)
- Smearing of positions (not recommended)

Crystal Producer

What does it read and write?

- Raw event: data from GGP electronics
- Traces length (100 samples of 10ns)
- Can write original data on disk (in compressed format)
 - > data files typically split after 1 M events ~ 3.6 GB
 - ✓ event_mezzdata.cdat.0000
 - ✓ event mezzdata.cdat.0001 ...
- Generate raw spectra for amplitudes and baselines
- Format data into a "data:crystal" adf frame

Prod__4-38-32768-UI_Ampli.spec [1]-38-32768 used for energy calibrations



Preprocessing Filter

Performs:

- Energy calibrations and cross talk corrections
- Analysis of traces
 - Calculation of T0 from the core signal
 - Time alignment of the segments
 - Energy and time calibration of the trace (amplitude and time shift)
 - Define the segment multiplicity (number of segments with a net charge)

• After Preprocessing:

- energies are stored in units of keV
- times are in units of samples (10 ns) (but time calibration parameters are in ns)
- positions are given in mm, when they show up after the PSA

Preprocessing Filter

Calibrations:

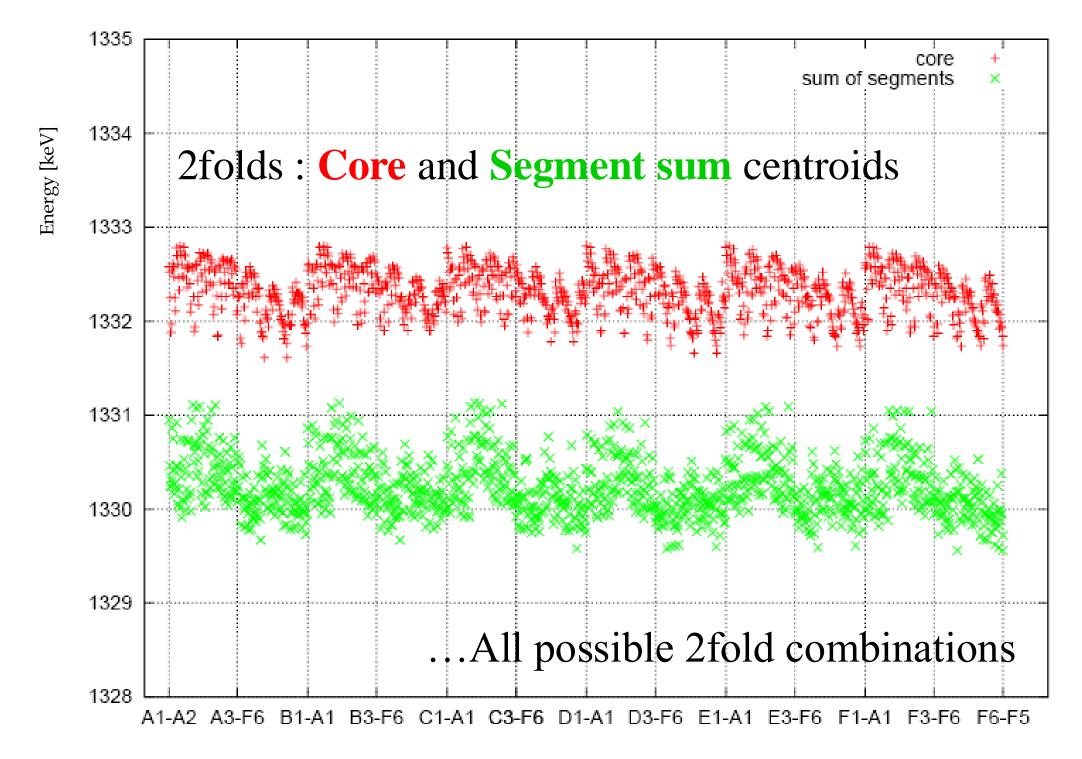
- Energy
 - Gain-only, no offset coefficient needed because of the way the amplitude is generated in the preprocessing electronics.
- Cross-talk
 - 36*36=1296 coefficients to correct capacitive coupling correlations between segments and core
 - Used also to recover up to one broken or missing segment per crystal
- Local Time
 - 36 coefficients to align segments to the core (great influence on the performance of PSA)

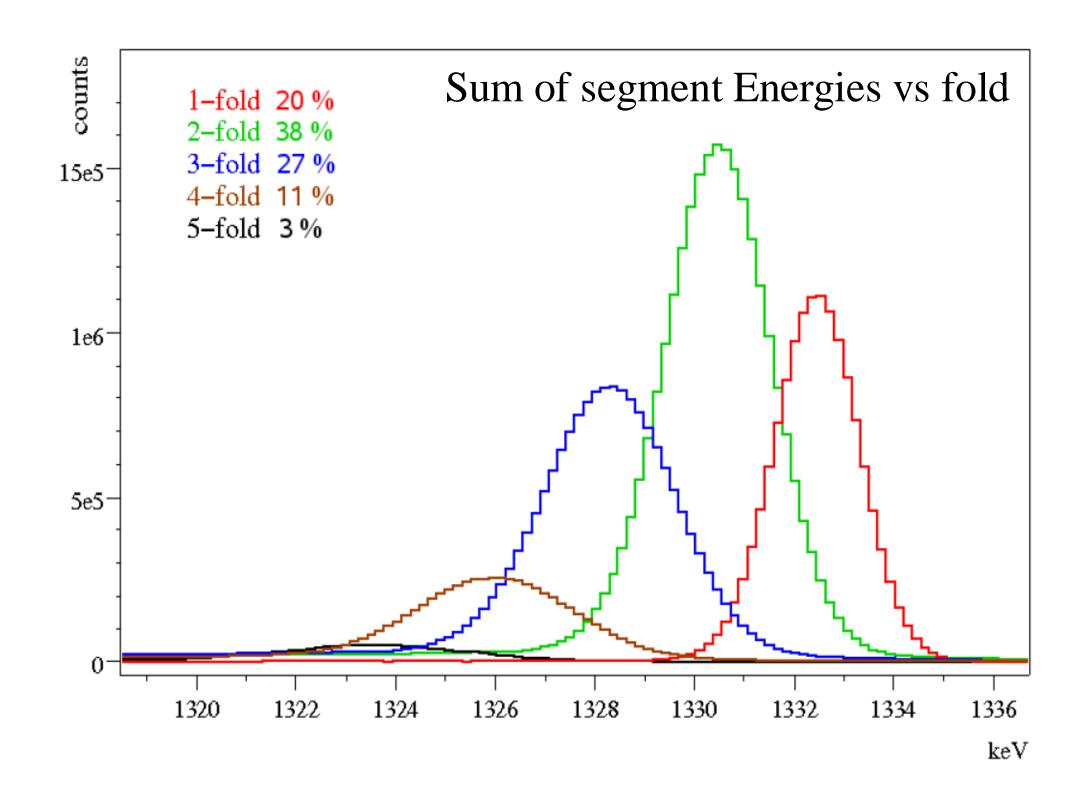
PreprocessingFilterPSA.conf

#segm/core	%d(id)	%f(tfall)	%f(trise)	%f(egain)	%f(emink)	%f(tmove)
segm	0	4600	1000	0.162979	10.0	9.891
segm	1	4600	1000	0.170079	10.0	10.463
segm	2	4600	1000	0.163820	10.0	10.361
segm	3	4600	1000	0.169401	10.0	9.485
segm	4	4600	1000	0.158867	10.0	7.971
segm	5	4600	1000	0.155504	10.0	10.077
segm	6	4600	1000	0.170291	10.0	9.050
segm	7	4600	1000	0.165092	10.0	9.263
segm	8	4600	1000	0.145804	10.0	7.420
segm	9	4600	1000	0.168806	10.0	8.448
segm	10	4600	1000	0.143493	10.0	6.188
segm	11	4600	1000	0.159609	10.0	10.510
segm	12	4600	1000	0.153815	10.0	10.251
segm	13	4600	1000	0.155996	10.0	9.448
segm	14	4600	1000	0.168760	10.0	9.537
segm	15	4600	1000	0.175860	10.0	8.866
segm	16	4600	1000	0.185031	10.0	13.873
segm	17	4600	1000	0.157300	10.0	10.564
segm	18	4600	1000	0.169683	10.0	9.836
segm	19	4600	1000	0.168100	10.0	9.683
segm	20	4600	1000	0.170233	10.0	9.677
segm	21	4600	1000	0.174663	10.0	9.472
segm	22	4600	1000	0.174109	10.0	8.942
segm	23	4600	1000	0.165021	10.0	11.498
segm	24	4600	1000	0.152862	10.0	10.267
segm	25	4600	1000	0.169911	10.0	11.067
segm	26	4600	1000	0.165142	10.0	9.910
segm	27	4600	1000	0.159595	10.0	9.393
segm	28	4600	1000	0.168353	10.0	8.347
segm	29	4600	1000	0.167807	10.0	11.519
segm	30	4600	1000	0.163006	10.0	9.945
segm	31	4600	1000	0.159887	10.0	10.383
segm	32	4600	1000	0.155449	10.0	9.600
segm	33	4600	1000	0.143345	10.0	8.487
segm	34	4600	1000	0.150043	10.0	7.216
segm	35	4600	1000	0.176351	10.0	12.197
core	0	4200	1000	0.347055		25.000
core	1	4200	1000	0.069358	20.0	-0.285
tntf 209715	52					

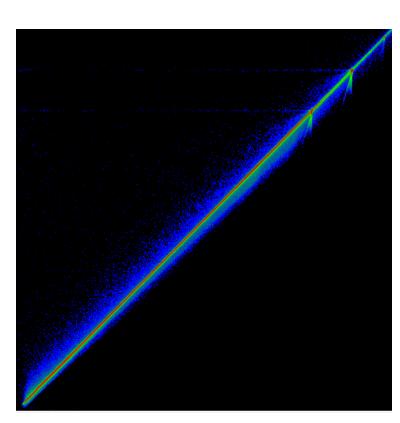
Cross-Talk correction: Motivations

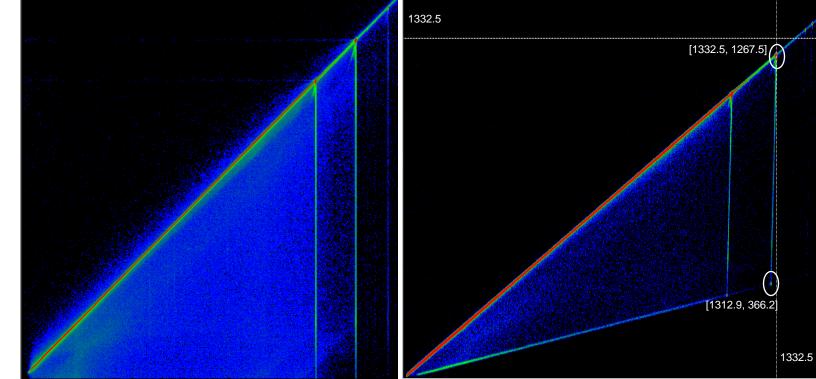
- Crosstalk is present in any segmented detector
- Creates strong energy shifts proportional to fold
- Tracking needs segment energies!





Missing/Broken segments



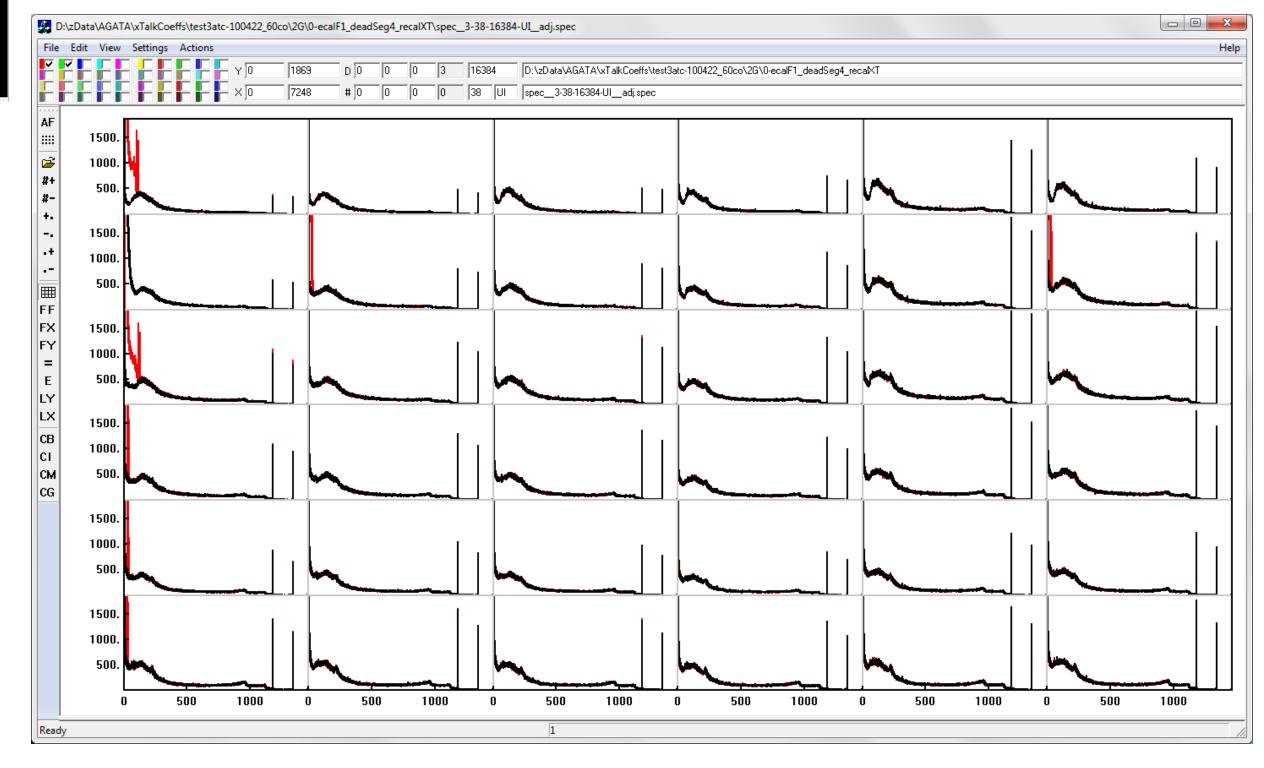


• Exploiting the redundancy:

$$\sum Segs == Core$$

- Compensate loss of energy in core
- Assign energy to missing segment

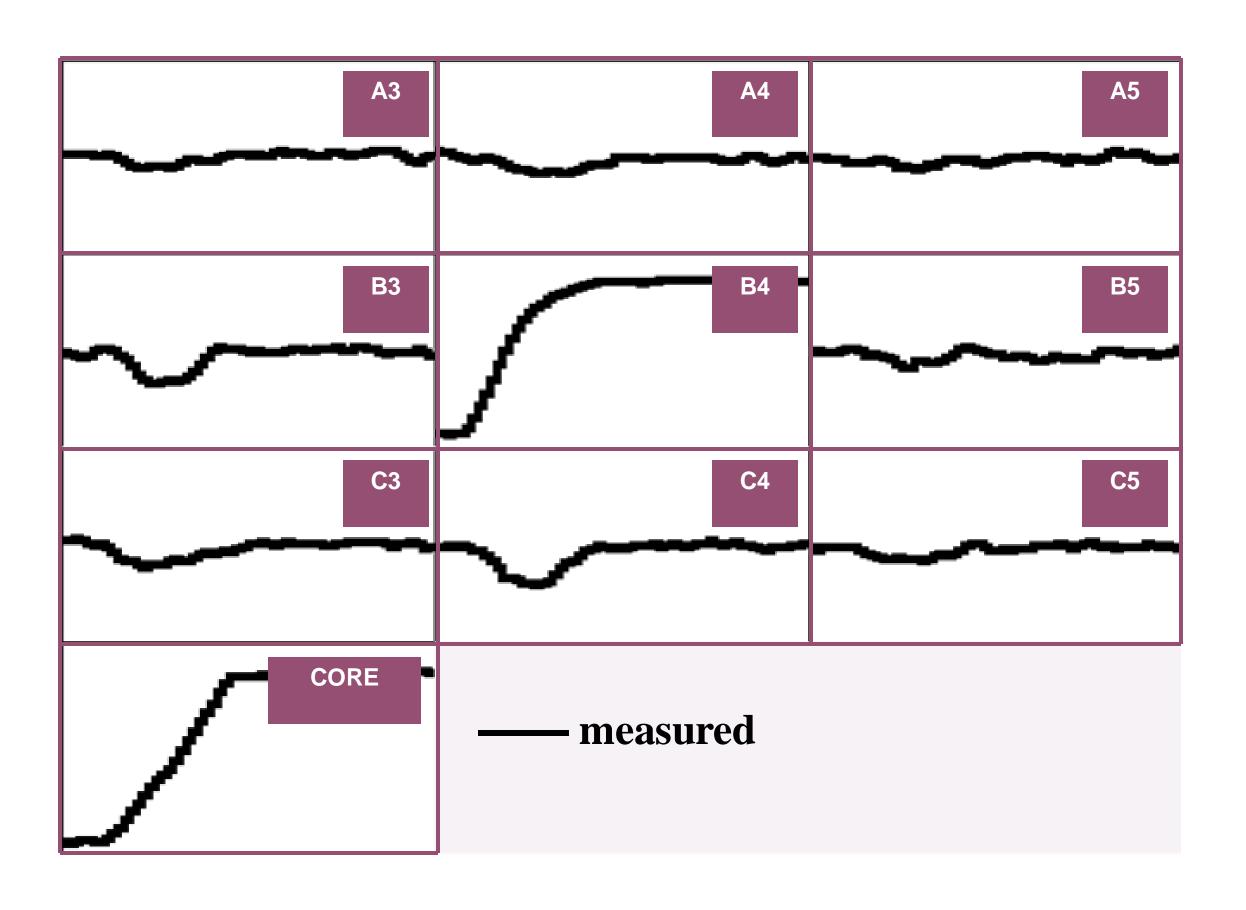
$$E_{\text{missing}} = E_{\text{core}} - Sum_{\text{Other}}$$



PSA Filter

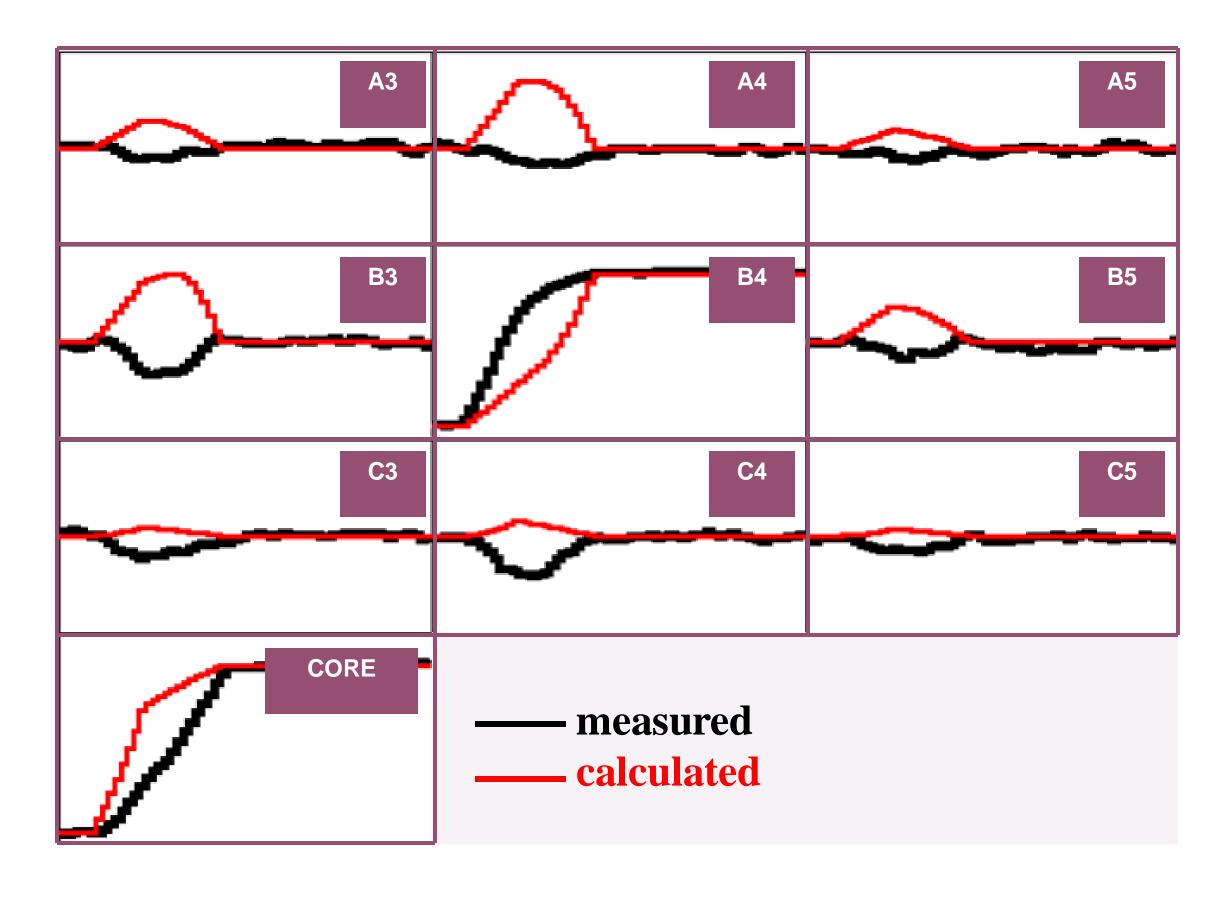
Performs:

- Read the simulated basis (ADL)
- Apply the preamp response function to the simulated traces
- Perform the signal decomposition:
 - > Implemented algorithm is the Grid Search
 - ✓ As a full grid search
 - ✓ As an adaptative coarse/fine search (AGS) => Default used online
- Reduces size of data by factor ~80
- Provides the parameters for the correction of neutron damage
- Is the critical point for the processing speed of online and offline analyses
 - > Takes ~ 95 % of total CPU time in the AGATA data processing

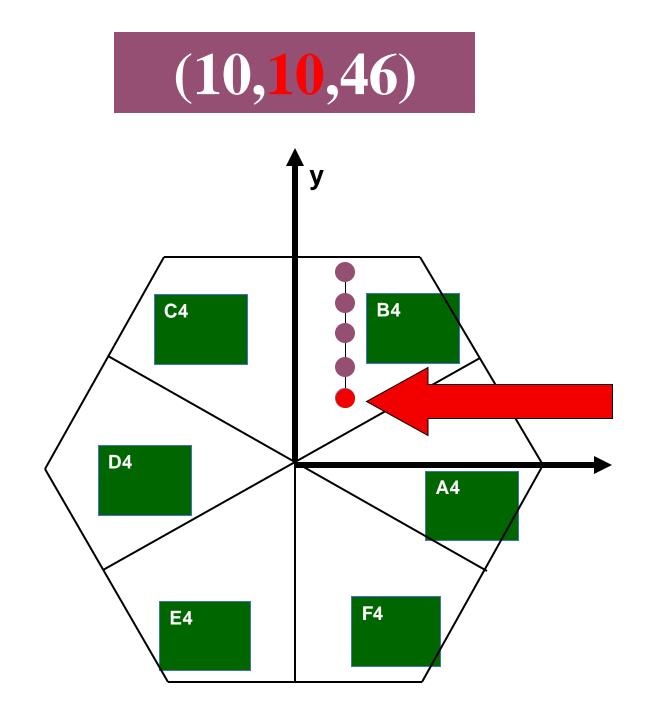


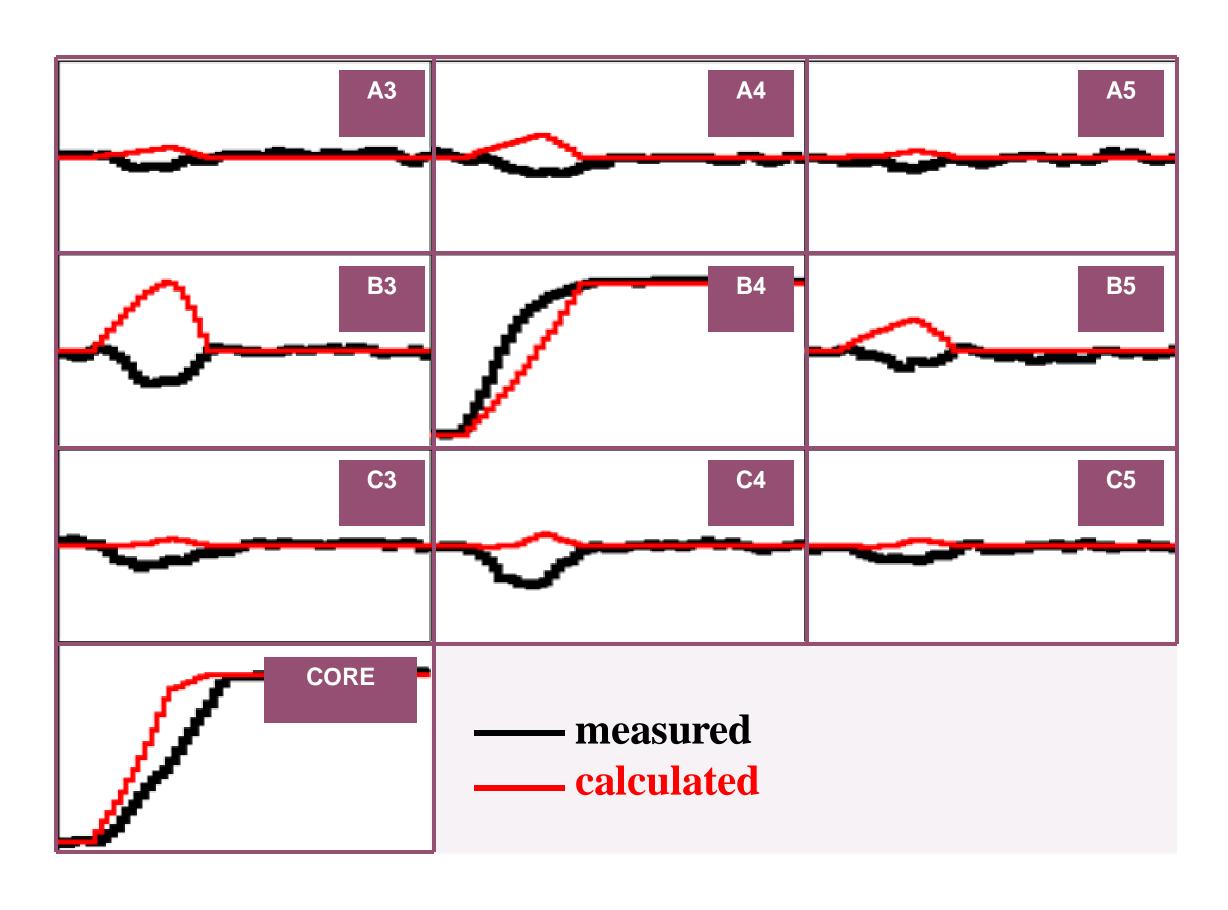
T B B C

791 keV deposited in segment B4

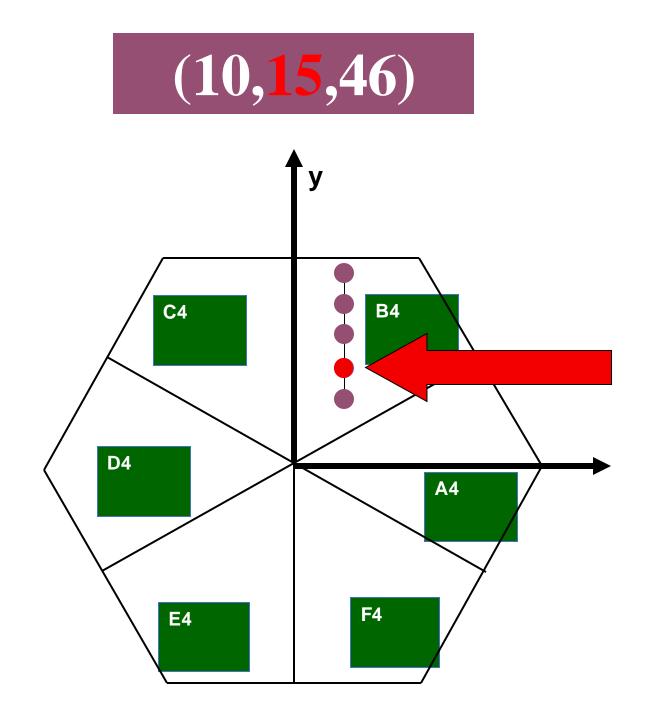


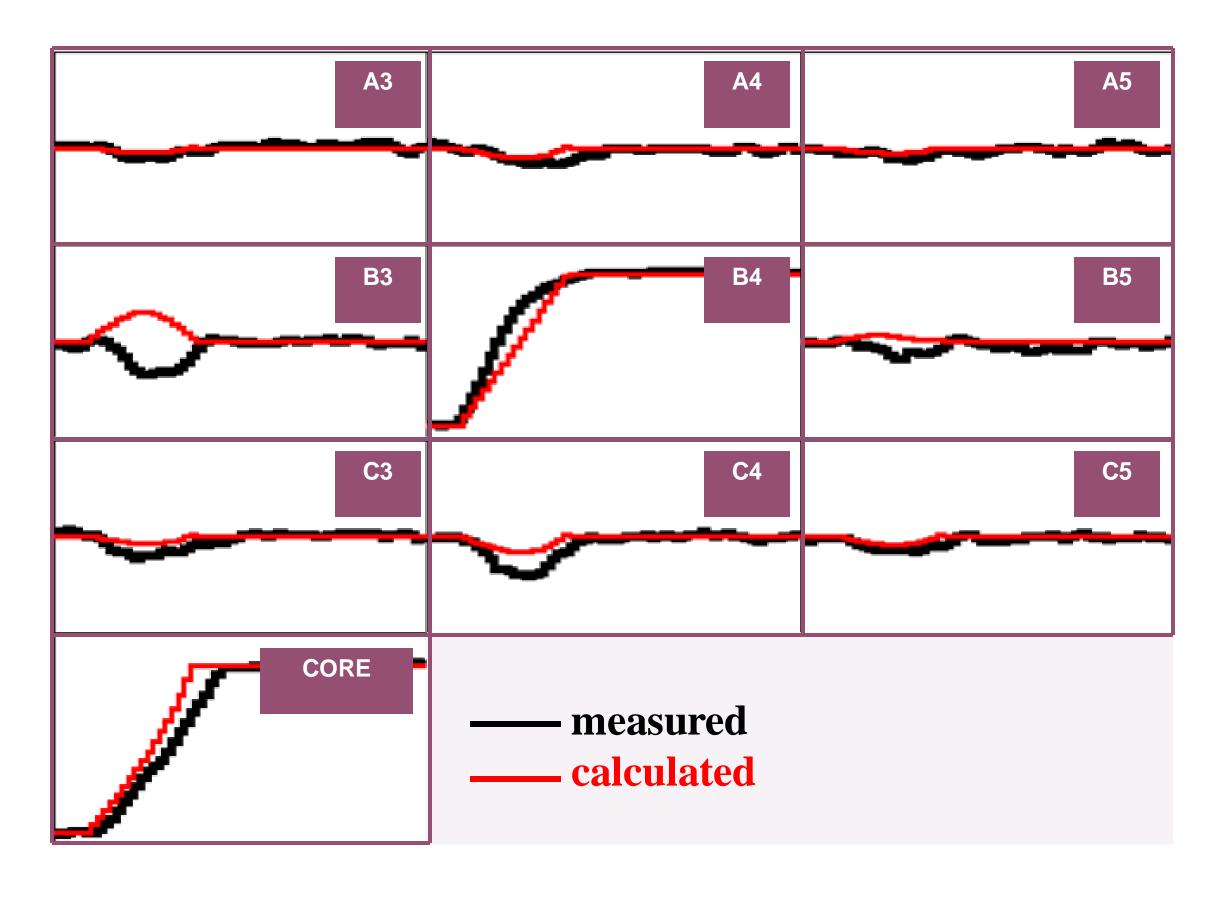
791 keV deposited in segment B4



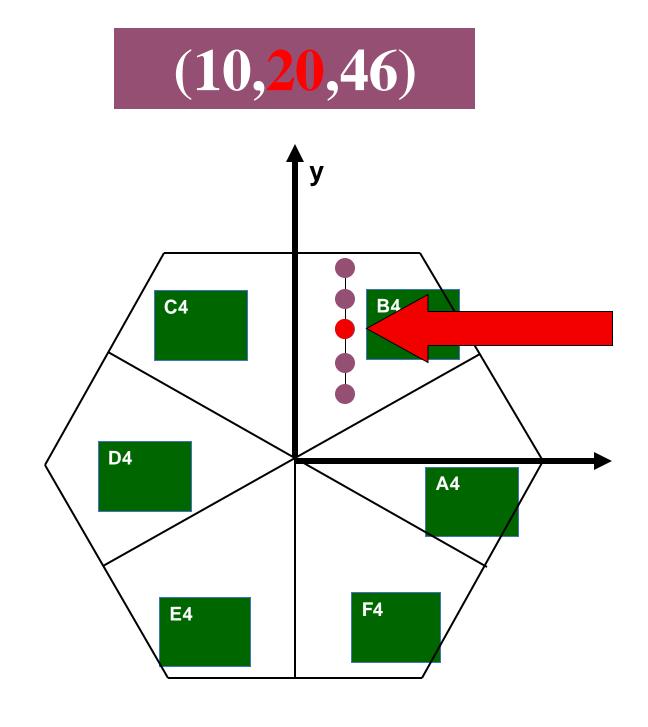


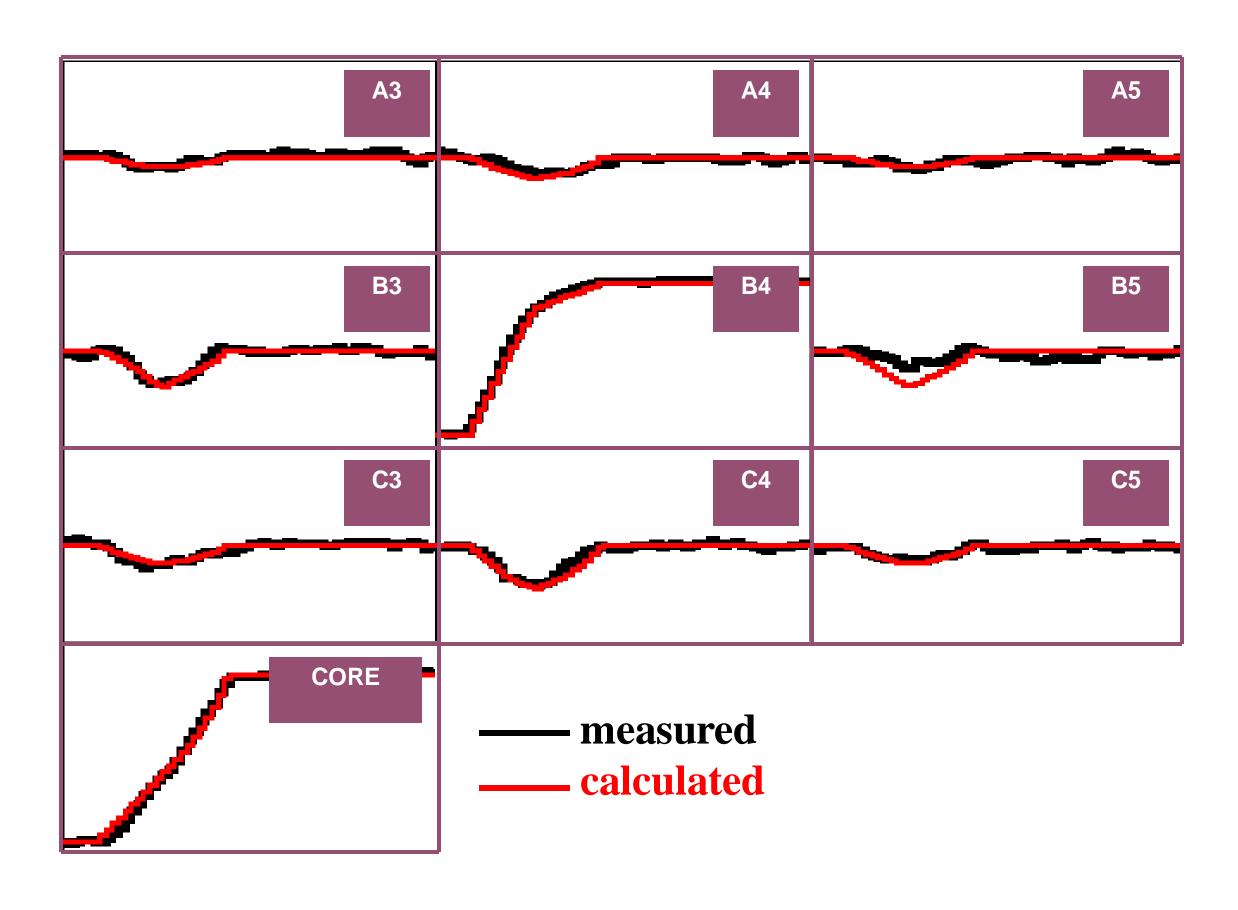
791 keV deposited in segment B4



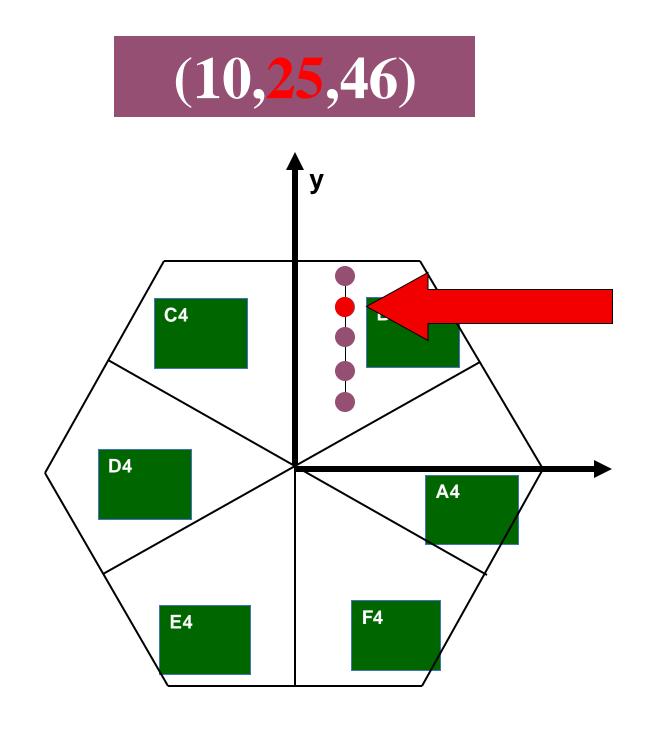


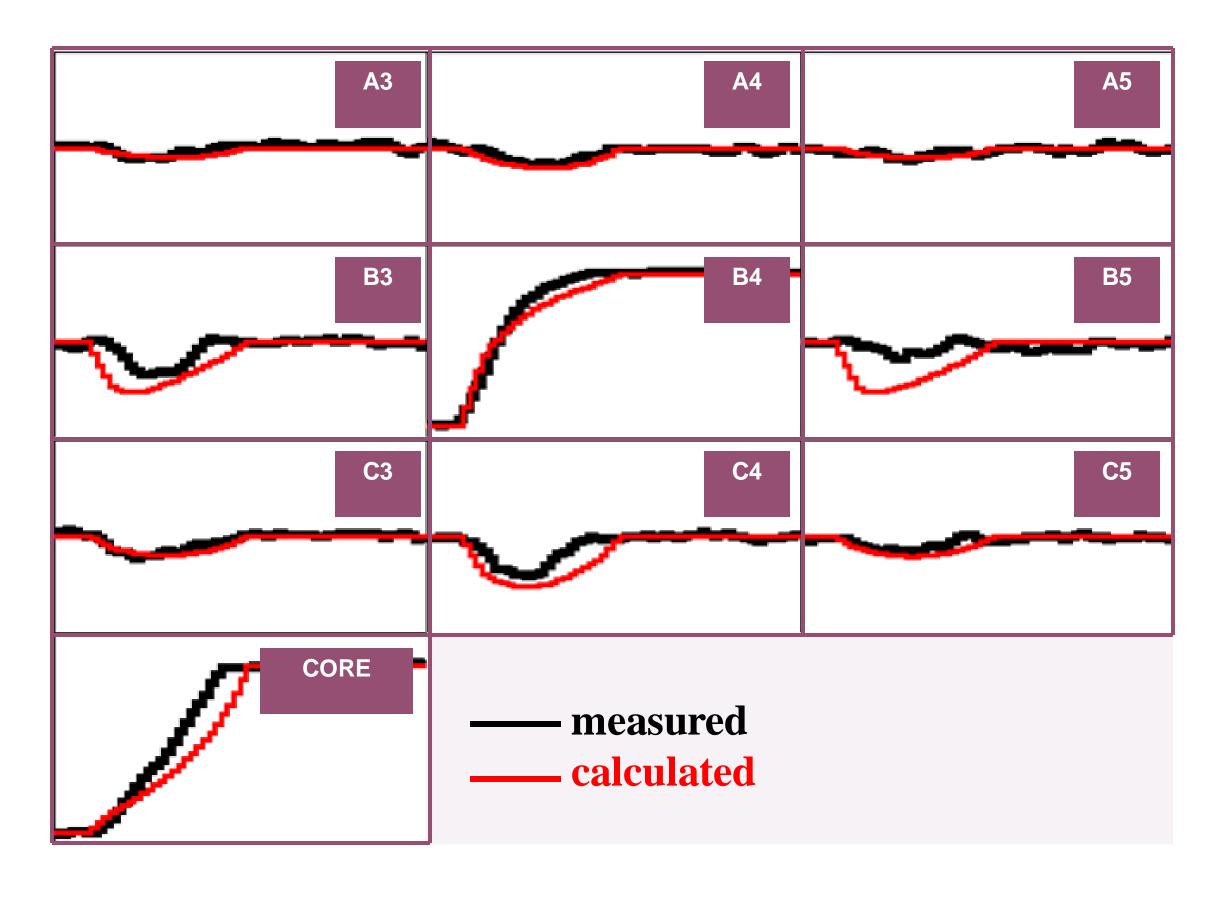
791 keV deposited in segment B4



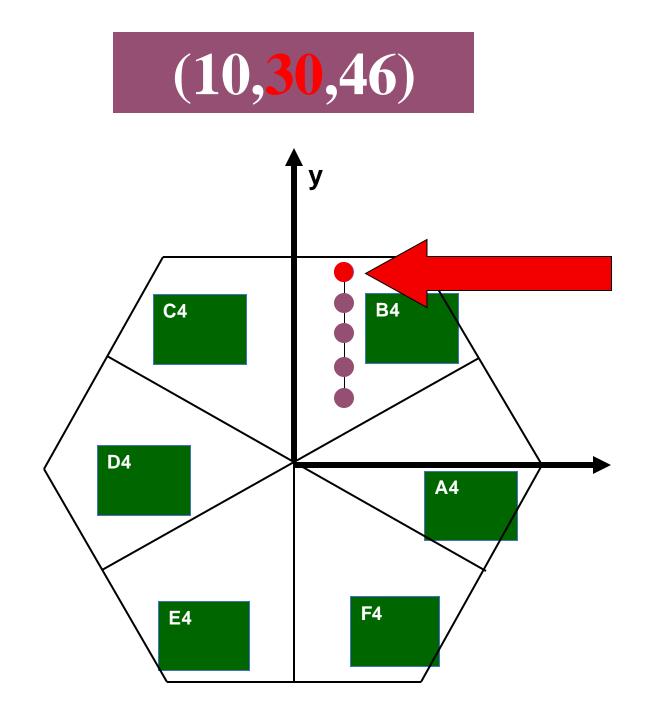


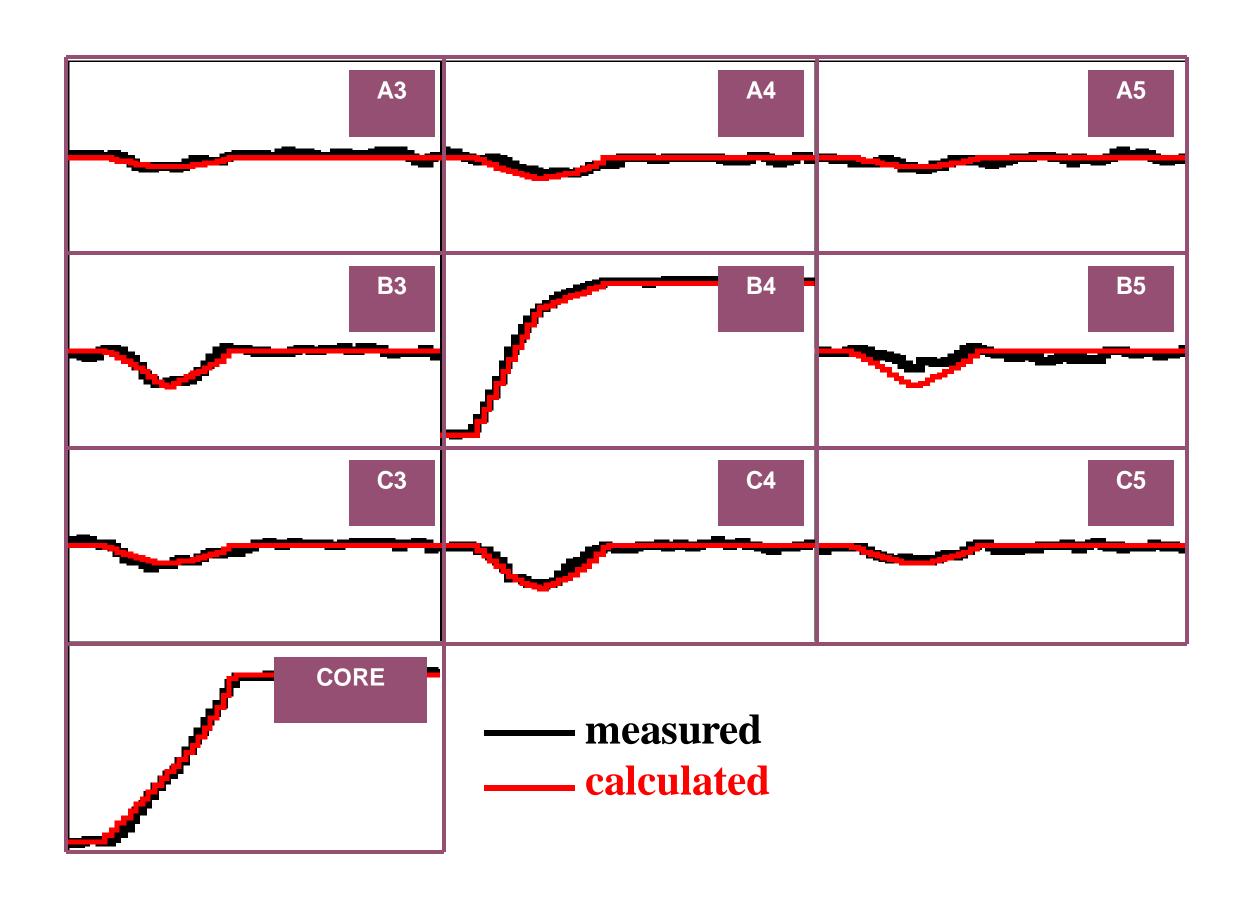
791 keV deposited in segment B4



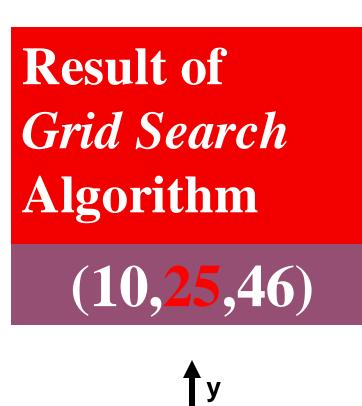


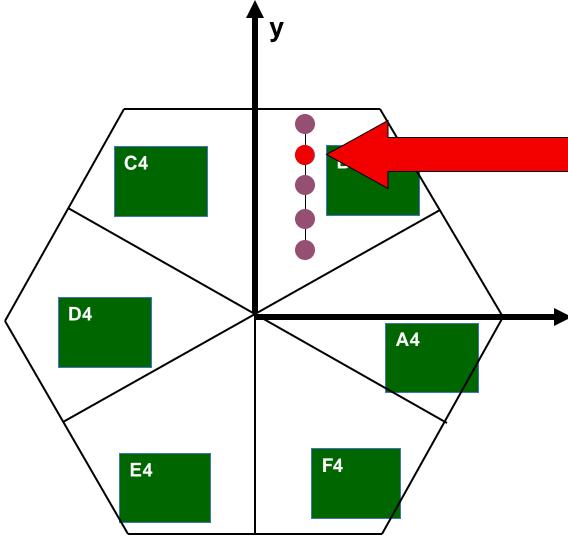
791 keV deposited in segment B4





791 keV deposited in segment B4

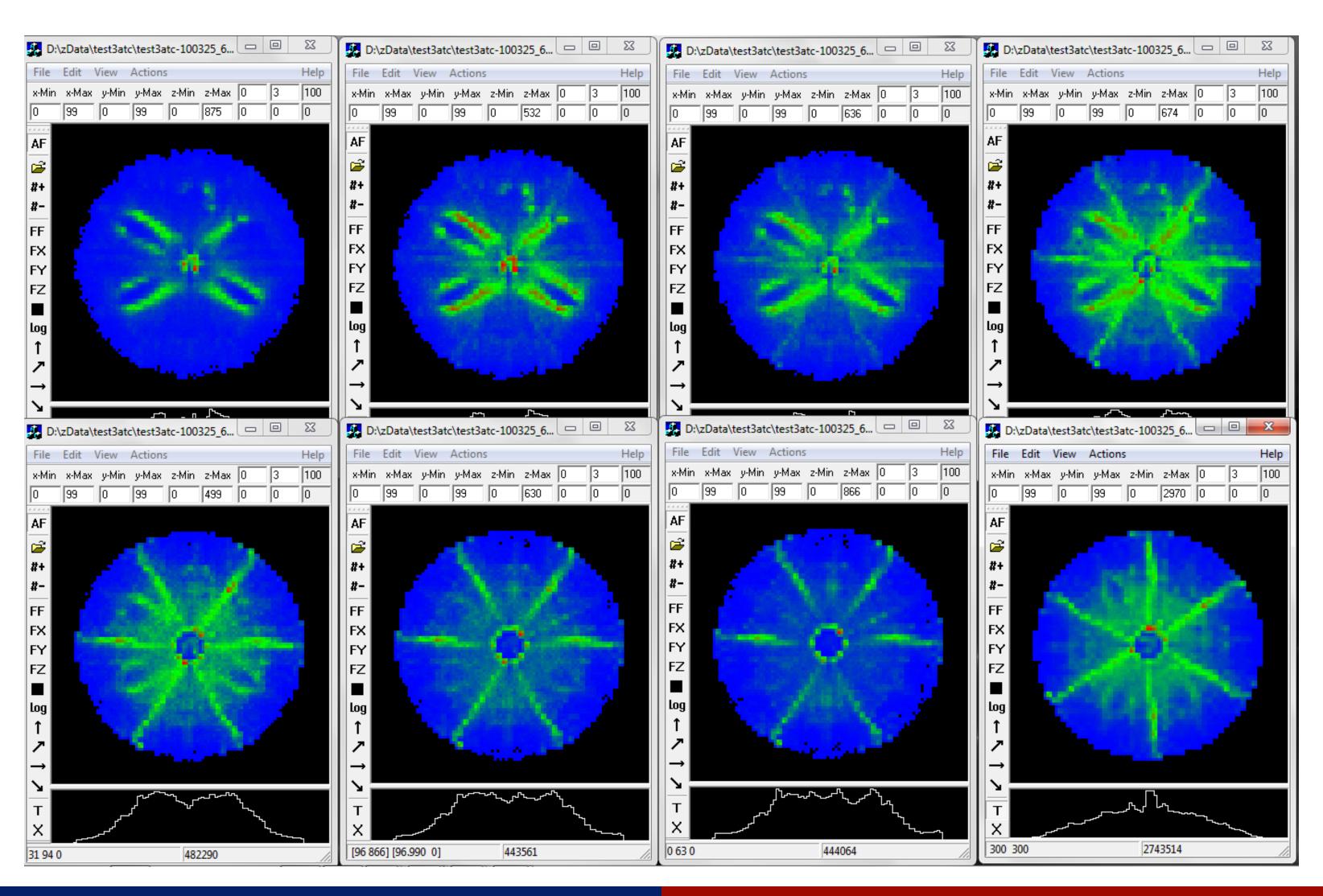




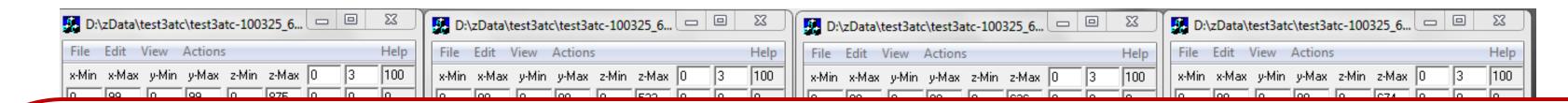
The Grid Search algorithm

- Signal decomposition assumes one interaction per segment
- The decomposition uses the transients, the net charge segment, and core pulses
- Proportional and differential cross-talk are included using the xTalk coefficients of the preprocessing
- No limit to the number of fired segments (i.e. up to 36)
- The number of used neighbours is a compile time parameter (usually 2 as Manhattan distance)
- The algorithm cycles through the segments in order of decreasing energy:
 - → the result of the decomposition is removed from the remaining signal
- Using ADL bases (Bart Bruyneel)
 - Fine grid: 2 mm step: \rightarrow ~ 48000 grid points in a crystal (700-2000 points/segments)
- Speed is:
 - → ~ 150 events/s/core for the Full Grid Search
 - → ~ 1500 events/s/core for the Adaptive Grid Search
- To speedup execution, parallelism has been implemented with
 - → blocks of ~ 300 events passed to 5 parallel threads of execution

Effect of time alignment



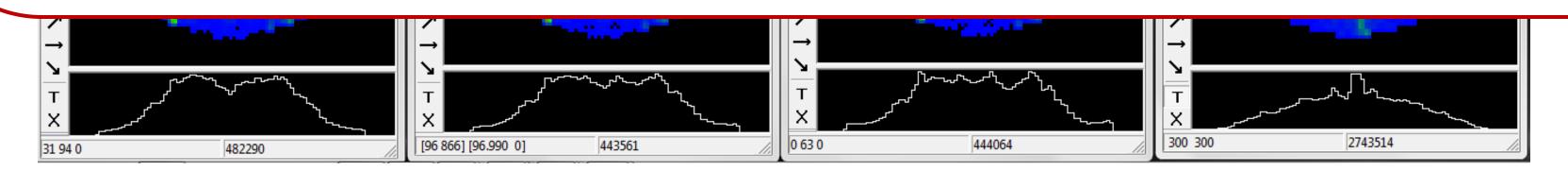
Effect of time alignment





All up to this points cannot be redone after the experiment

This is the charge of the onsite team



Post PSA Filter

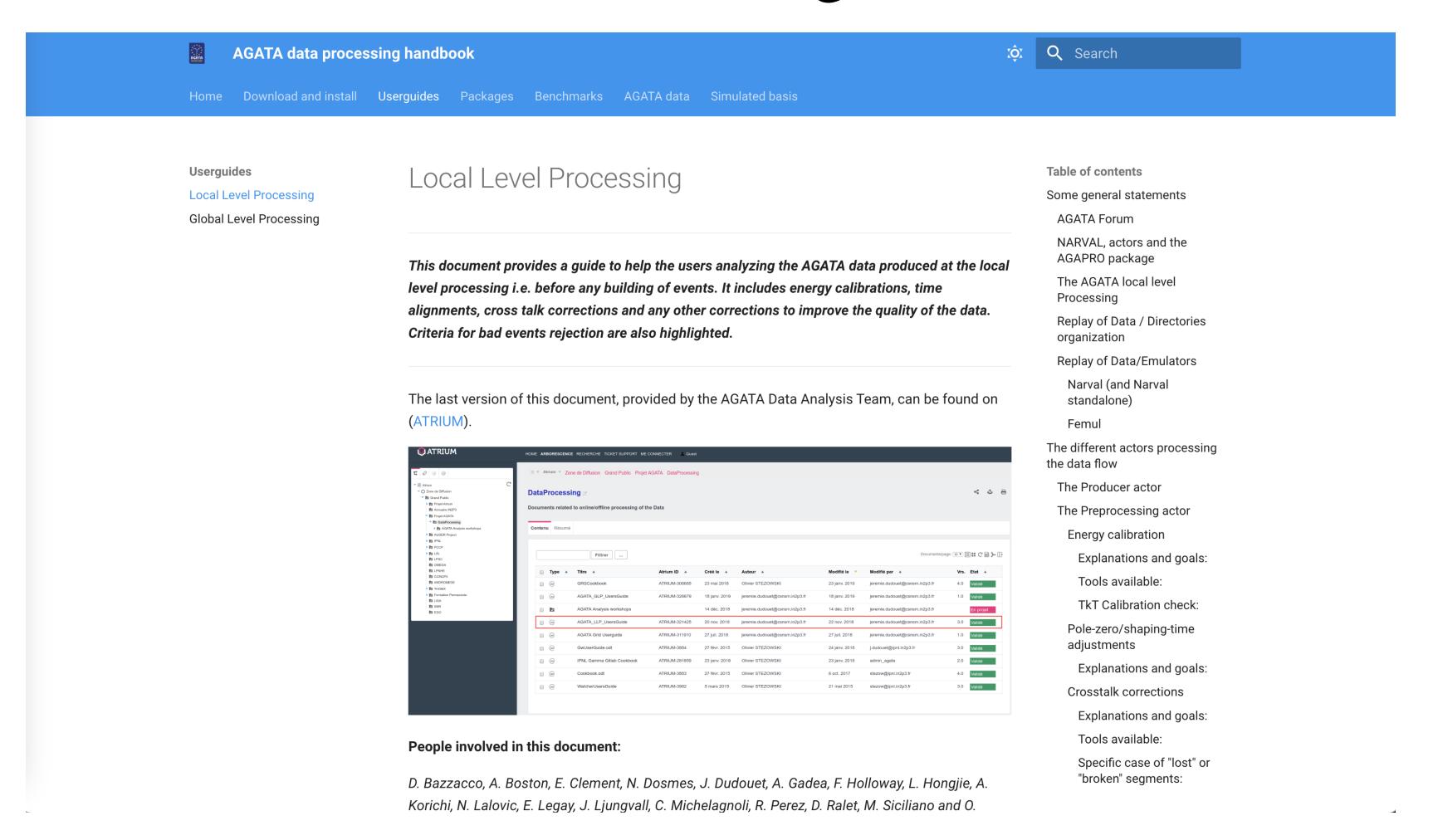
Performs:

- Final energy calibrations of cores and segments (with offset)
- Force segments to core (optional)
- Recovery of neutron damage
- Global time alignment of crystals \rightarrow important to reduce random coincidences

Global actors

- Event Builder
 - Merge all AGATA crystal in events on the basis of timestamps
- Event Merger
 - Merge AGATA built events with ancillaries on the basis of timestamps
- Tracking Filter
 - Apply the tracking algorithm using segments energies and positions
- Tree Builder
 - Dump the data on disk using ROOT tree format

AGATA user guides



https://agata.pages.in2p3.fr/handbook/



Thank you! And have fun with your data analysis