

# Status of Tracking for Agata

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1 Online Tracking

2 Status of OFT

3 To do....

# Online Tracking

- OFT disabled for experiments with event rate  $> 1.5\text{-}2$  kHz

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→ 4x faster

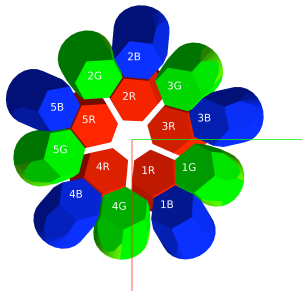
# Online Tracking

- OFT disabled for experiments with event rate  $> 1.5\text{-}2$  kHz
- mgt implemented into Narval environment (3x faster than OFT)
- optimization of OFT
  - 4x faster
  - tracking no longer bottleneck of the DAQ



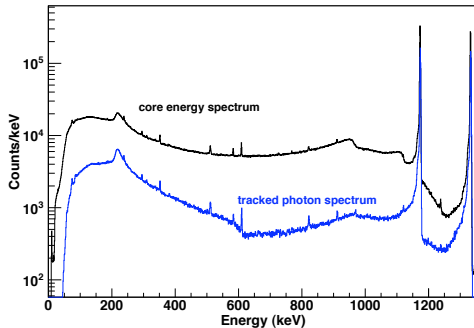
## Source run

- tracking with source data taken in April 2010:
  - 3 triple clusters : ATC1, ATC2, ATC4 (NB: non adjacent detector !)
  - $^{60}\text{Co}$  @  $z=27.2$  cm (nominal position :  $z=0$ )
  - PSA interaction points (actor revision # 879) written to Agata-type file



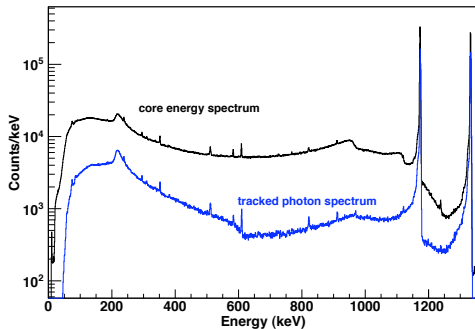


# Issues



- $P/T \sim 35\%$  and  $\epsilon_{tracking} \sim 53\%$  (with standard tracking parameters)

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- single-hit threshold

## Why such a performance ?

- use AGATA simulation code to try to find the origin(s) of the problems

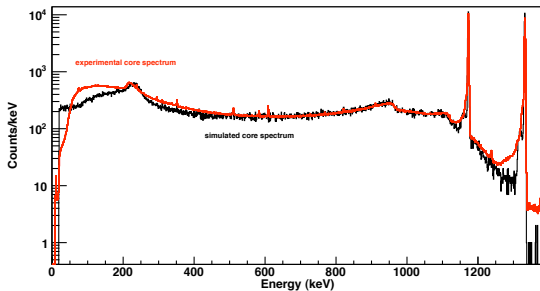
configuration of the simulations:

- ATC1,2,4 with passivated areas, capsules and cryostats
- 2 mm Al chamber
- PRISMA ancillary (pushed back by 8 cm wrt nominal position @  $Z=40$  cm)
- packing per segment at the energy-weighted barycenter
- standard energy resolution for segments
- 20-keV threshold on segment energy
- emission multiplicity = 2



# Why such a performance ?

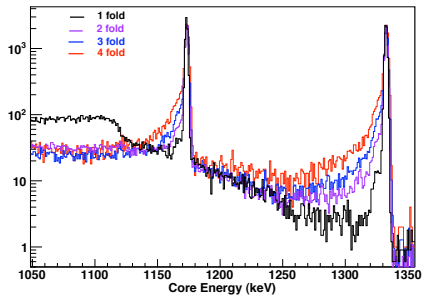
core-energy spectra normalised to peak+tail



- worse experimental input spectral quality (P/T=15% vs 19%)

# Tails

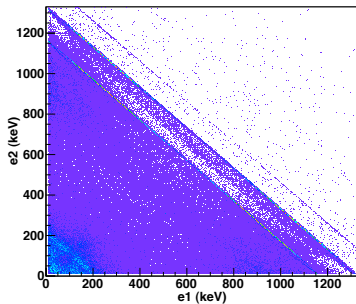
core-energy spectra normalised to peak



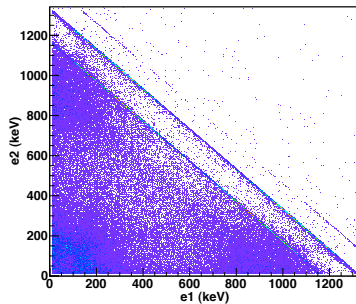
- tails appear for events with more than 1 segment hit in a detector

# Tails

fold 2 (slice1)



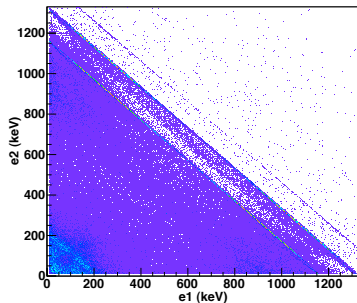
fold 2 (slice2)



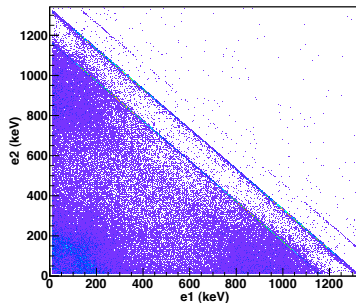
- Bow effect

# Tails

fold 2 (slice1)



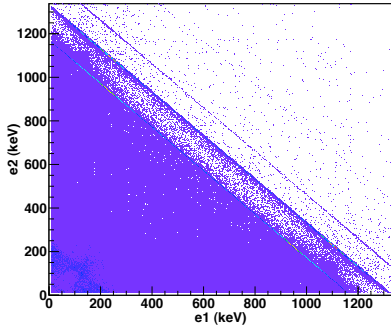
fold 2 (slice2)



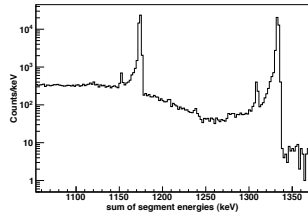
- Bow effect
- How does this affect interaction positions ?

# Tails

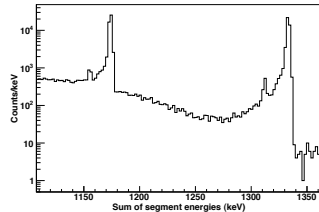
fold 2 (coinc. between slices 2-3)



fold 2 slices 2-3



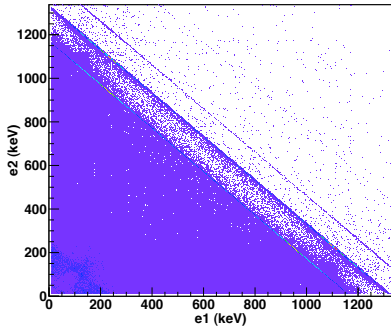
fold 2 slices 1-2





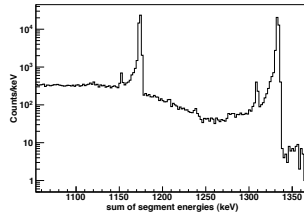
# Tails

fold 2 (coinc. between slices 2-3)

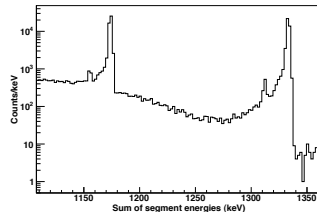


- problem arises with coincident segs A1,A2,A3 (individual spectra OK)

fold 2 slices 2-3

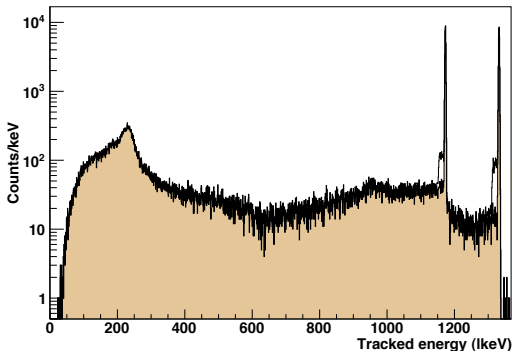


fold 2 slices 1-2



# Core-energy correction

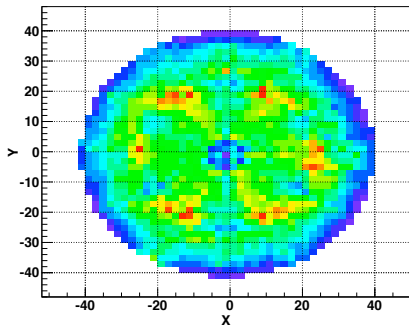
difference between core energy and  $\Sigma e_{segments}$  is redistributed among all interaction points



- peak is recovered (+5%) and P/T : 45% → 47%

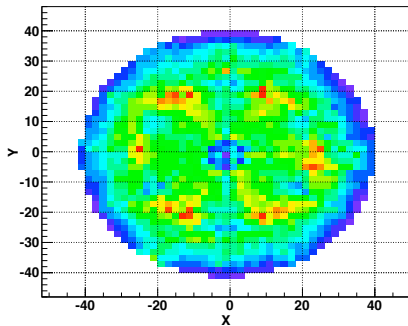
# Distribution of interaction points

april 2010

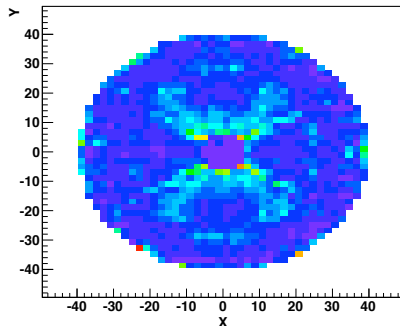


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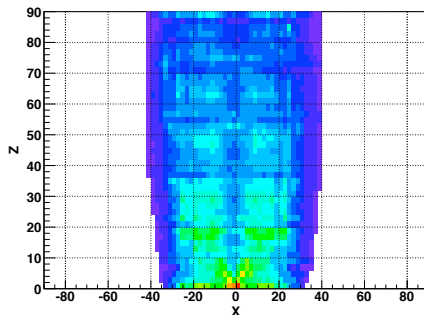


august 2009



- smoother (X,Y) distribution & 6-fold symmetry

# Distribution of interaction points



- But - clustering of interaction points & quantization in depth

## To do (help is welcome!)

- improve quality (reality) of simulations to understand tracking response
- modify single-interaction treatment to account for high-energy single interactions and depth quantization
- compare mgt/OFT
- track with core-energy-corrected data (replay recently performed)