



Histogram
tools: NEST
and HistoFit

Oleksandr
Volynets

NEST

Motivation
Implementation
Results

HistoFit

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Implementation
Dummy data
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Histogram tools: NEST and HistoFit

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Outline

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- Introduction
- NEST(New Energy Spectrum Toolkit):
 - Motivation
 - Implementation
 - Results
- HistoFit
 - Motivation
 - Implementation, method
 - Creating dummy data for check
 - Results of decomposing spectras



Introduction

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- MaGe/Geant4 is a powerful tool to get the expected energy spectrum of the experiment.
- However, all we get is a set of files with spectras from each holder, cable, detector etc.
- But the Data only contains one single part which is naturally a sum of all contributions.
- We have to do offline histogram analysis to get all those files together to see how good we understand our detector.
- In this presentation two histogram tools for spectra analysis will be shown



NEST(New Energy Spectrum Toolkit)

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- The detector has many parts, we simulate (dominant) radioactive sources in separate parts;
- In order to plot MC spectrum we have to add all background contributions with its' weights according to activities/masses/time;
- To deal with all these histograms a tool with simple control of masses, activity etc. of all sources is needed;



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The tool called NEST(New Energy Spectrum Toolkit) was created by Jens Schubert. It has the following features:

- Draw separately *Total energy in all crystals/Total energy in every crystal/Energy in the individual segment*;
- Draw separately *All/Single-crystal/Single-segment events*;
- All parameters of the sources are read from a single text file; all the parameters of the geometry are read from a single XML-file;
- Flexible control of output (counts in energy window, expected counts of every single background contribution etc.);
- Tag the contributions and get numbers (e.g. background index) for only needed tags;
- ...



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Example list of parameters:

```
# +-----+
# | +-----+ |
# | | LIFE TIMES | |
# | +-----+ |
# +-----+
#HALFlifetime__Agm108          418 years  "the HALFlifetime of Agm108"
lifetime__Agm108              1.9032785e10 s  "the lifetime of Agm108"
#HALFlifetime__Agm110          249.9 days  "the HALFlifetime of Agm110"
lifetime__Agm110              3.11497e7 s   "the lifetime of Agm110"
# +-----+
# | CONTAMINATION LEVELS  -- in Phase-1-Crystals |
# +-----+
contamination_Th232__Xtal_ANGx          5.5e-8      Bq/kg  "The contamination of
contamination_Th232_MeasurementDate__Xtal_ANGx/date 2008-01-01 none  "The contamination of
```



Implementation

Example geometry description (PhaselIdeal.xml):

```
<ListOfElBaCos>
  <Replacer>
    <ReplaceExpressionName> <RM>ProcBaseName</RM><RM>Nnu</RM> </ReplaceExpressionName>
    <ReplaceExpressionValue> <RM>DBD_2nu2beta</RM><RM>2 </RM> </ReplaceExpressionValue>
    <ReplaceExpressionValue> <RM>DBD_0nu2beta</RM><RM>0 </RM> </ReplaceExpressionValue>
  </Replacer>

  <!-- loop over all phase-1 crystals -->
  <ListOfElBaCos switch="on">

    <Replacer>
      <ReplaceExpressionName> <RM>XtalName</RM><RM>GeXXFrac</RM><RM>PhysVolIndex </RM>
      <ReplaceExpressionValue switch="on"> <RM>ANG1 </RM><RM>ANG1 </RM><RM>6 </RM>
      <ReplaceExpressionValue switch="on"> <RM>ANG2 </RM><RM>ANG2 </RM><RM>12 </RM>
      <ReplaceExpressionValue switch="on"> <RM>ANG3 </RM><RM>ANG3 </RM><RM>13 </RM>
      <ReplaceExpressionValue switch="on"> <RM>ANG4 </RM><RM>ANG4 </RM><RM>7 </RM>
      <ReplaceExpressionValue switch="on"> <RM>ANG5 </RM><RM>ANG5 </RM><RM>11 </RM>
      <ReplaceExpressionValue switch="on"> <RM>GTF1 </RM><RM>GTFx </RM><RM>5 </RM>
      <ReplaceExpressionValue switch="on"> <RM>GTF2 </RM><RM>GTFx </RM><RM>3 </RM>
      <ReplaceExpressionValue switch="on"> <RM>GTF3 </RM><RM>GTFx </RM><RM>2 </RM>
      <ReplaceExpressionValue switch="on"> <RM>GTF4 </RM><RM>GTFx </RM><RM>1 </RM>
      <ReplaceExpressionValue switch="on"> <RM>GTF5 </RM><RM>GTFx </RM><RM>4 </RM>
      <ReplaceExpressionValue switch="on"> <RM>GTF6 </RM><RM>GTFx </RM><RM>0 </RM>
      <ReplaceExpressionValue switch="on"> <RM>RG1 </RM><RM>RG1 </RM><RM>9 </RM>
      <ReplaceExpressionValue switch="on"> <RM>RG2 </RM><RM>RG2 </RM><RM>10 </RM>
      <ReplaceExpressionValue switch="on"> <RM>RG3 </RM><RM>RG3 </RM><RM>8 </RM>
    </Replacer>
  </ListOfElBaCos>
  ...

```




Results

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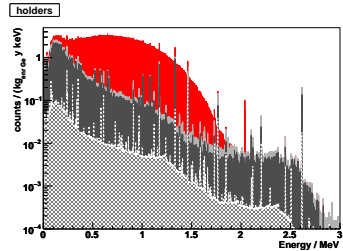
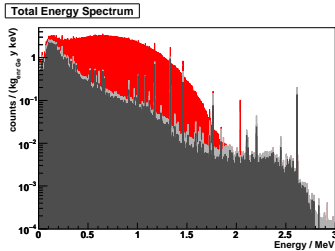
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Let's imagine the situation:

- Monte Carlo energy spectras calculated



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Let's imagine the situation:

- Monte Carlo energy spectras calculated
- Data energy spectrum obtained



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Let's imagine the situation:

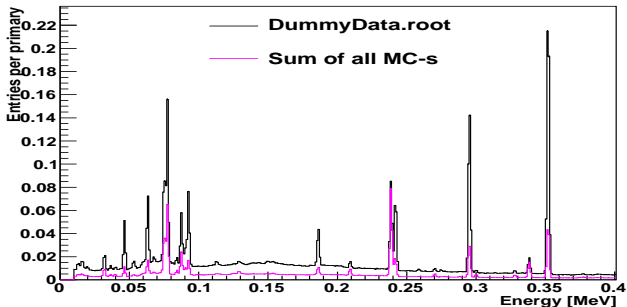
- Monte Carlo energy spectras calculated
- Data energy spectrum obtained
- Monte Carlo describes the shape, but not the numbers



Motivation

Let's imagine the situation:

- Monte Carlo energy spectras calculated
- Data energy spectrum obtained
- Monte Carlo describes the shape, but not the numbers



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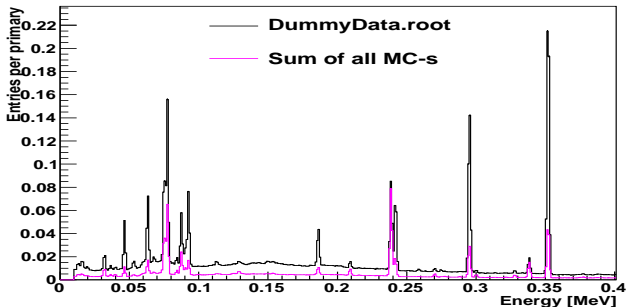
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Let's imagine the situation:

- Monte Carlo energy spectras calculated
- Data energy spectrum obtained
- Monte Carlo describes the shape, but not the numbers
- Background or Signal contribution estimated wrong?





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Tool developed for:

- Fitting several main MC contributions to data spectrum
- Getting correct estimates of the contribution



Motivation, implementation

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Tool developed for:

- Fitting several main MC contributions to data spectrum
- Getting correct estimates of the contribution

Method:

$$\chi^2 = \sum_{i, \text{bins}} \frac{(D_i - \alpha_0 MC_i^0 - \alpha_1 MC_i^1 - \dots)}{(\sigma_{D_i}^2 + \alpha_0^2 \sigma_{MC_i^0}^2 + \alpha_1^2 \sigma_{MC_i^1}^2 + \dots)}$$

Use MINUIT to minimize and get α_i parameters



Creating dummy data spectrum

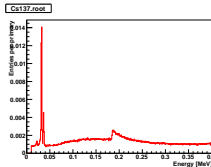
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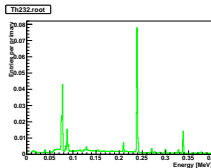
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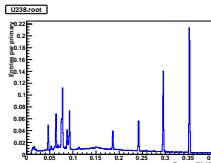
To check whether it works a dummy data was created:



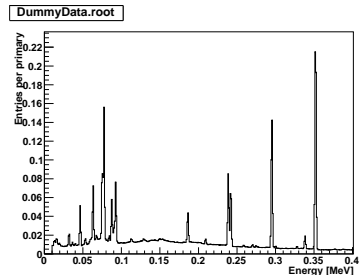
$\times 2$



$\times 1 \Rightarrow$



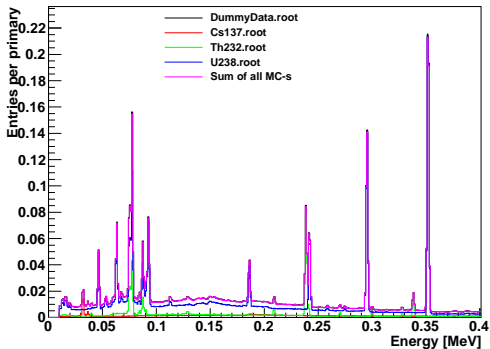
$\times 5$





Self-check: Decomposing spectrum

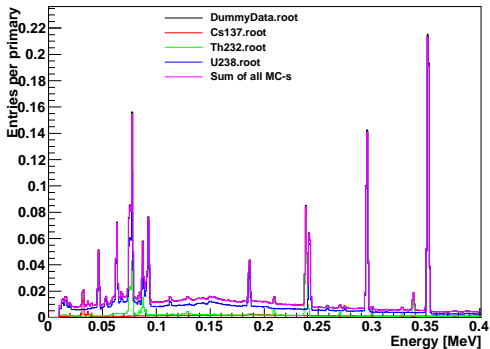
Using the same 3 histograms to fit to “Dummy Data”:





Self-check: Decomposing spectrum

Using the same 3 histograms to fit to “Dummy Data”:



Parameter 0 = 2 ± 0.00679609

Parameter 1 = 0.999997 ± 0.00316915

$\chi^2 \approx 10^{-5}$

Parameter 2 = 4.99999 ± 0.00722358





Self-check 2: Decomposing wrong spectrum

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Same procedure, but now Dummy Data contains deposits from 3 sources: ^{137}Cs , ^{232}Th and ^{238}U , but set of MC-spectras contains only 2 of them. And it converges but with wrong results (using given spectras):



Self-check 2: Decomposing wrong spectrum

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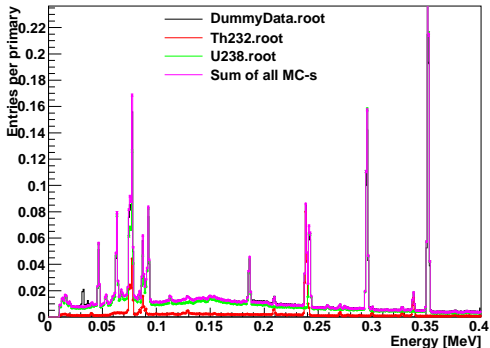
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Parameter 0 = *not used*

Parameter 1 = 1.029 ± 0.003

Parameter 2 = 5.638 ± 0.008

$$\chi^2 = 150000$$



Self-check 2: Decomposing wrong spectrum

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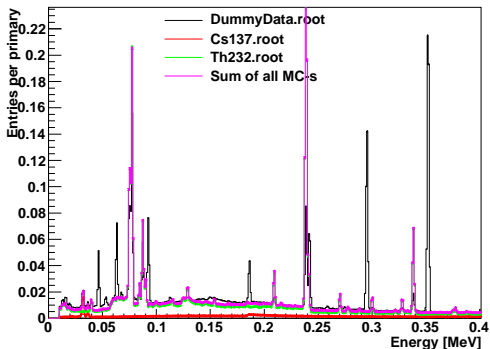
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Parameter 0 = 2.266 ± 0.009

Parameter 1 = 4.794 ± 0.008

Parameter 2 = *not used*

$$\chi^2 = 530000$$



Conclusions and discussion

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Next steps:

- Wait for the data at GERDA, or
- Find some data with measuring several sources simultaneously

Is the following needed:

- Fitting MC-s within different regions
(e.g. $Sp1=0..1.5$ MeV; $Sp2=1..2.3$ MeV) - complicated but possible
- Else?