# The MEGAlib software package

written by Andreas Zoglauer

# 1 Preamble

MEGAlib stands for Medium Energy Gamma-ray Astronomy Library. It represents the complete data analysis chain from simulations to high-level data analysis like image reconstruction in the Compton scattering energy regime. It has been originally developed for the MEGA telescope.

Although the library has undergone thorough testing and although there is no part of the program, which has not already been successfully used, like all other programs, also MEGAlib is not free of errors. Since the development of MEGAlib was only intended to get the work done I need for my research, MEGAlib will probably never leave its development phase. However, especially the older parts of the software are now in use for several years and can be considered as reliable and stable. In order to minimize the number of bugs and thus to improve the performance of the library and all its programs, please report all crashes, incompatibilities and other bugs to the following email address:

zog@ssl.berkeley.edu

Moreover if you think some features are missing or if you have enhancement requests, feel free to let me know — but consider, I am astrophysicist, no software developper!

Andreas Zoglauer Berkeley, June 2006

# 2 Installation

Currently the platforms Linux, Solaris and MacOS X are supported. No other Unix versions have been tested. The programs will definitely not run under Windows.

A detailed description of the installation can be found in the file INSTALL.txt, located in \$(MEGALIB)/doc.

# 3 License

The license is located in \$(MEGALIB)/doc/License.txt. Please read it carefully since it contains important information.

# 4 Special Options

# 4.1 Main Makefile

gmake gmake clean gmake cprogram name>
gmake clean\_cprogram name>
gmake texdoc
gmake man

Compile all programs
Remove all \*.o, \*.so files
Compile this specific program and launch it
Clean all files related to the program
Tex this documentation and launch gy

Generate doxygen documentation and place it in

doc/html/index.html

# 4.2 Special scripts

 ${\it launch~ sprogram~ name} > {\it soptions} > \qquad {\it Recompile~ (if~ necessary)~ and~ launch~ the~ given~ MEGAlib}$ 

program from every directory

 ${\it debug < program \ name > < options >} \qquad {\it Recompile \ (if \ necessary) \ and \ launch \ the \ given \ MEGAlib}$ 

program from every directory in debugging mode

# 5 Program overview

This is intended as a brief overview over the different programs which are available and thus the tasks which can be performed with MEGAlib.

# 5.1 Geomega

Geomega is the detector geometry package. All geometry and detector information is represented by a \*.geo.setup file. This includes the description of the volumes and materials as well as the detector properties. The geometry can be converted to formats which the GEANT3 simulation program GMega as well as MGeant/MGGPOD can process. The Geant4 program Cosima directly loads Geomega's shared library to use the geometry package. Almost all other programs use this geometry for geometry display (eview) as well as event and image reconstruction (especially absorption probabilities and resolutions).

# 5.2 GMega

Simulation tool based on Geant3. It is capable of handling different input spectra and source geometries. For details see the documentation in the GMega directory. It produces an output file, which can be read from the event analysis programs, like Sivan or Revan.

# 5.3 Cosima

Simulation tool based on Geant4. Like GMega it is capable of handling different input spectra and source geometries. For details see the documentation in the Cosima directory. It produces an output file, which can be read from the event analysis programs, like Sivan or Revan.

# 5.4 Sivan

Sivan is intended for the analysis of simulated data and explicitly using simulation information: the underlying library is mainly used to determine responses and determine the maximum possible efficiency of an event reconstruction algorithm.

#### 5.5 Revan

The programs provides the event reconstruction, i.e. it handles the transition of the events from pure hits into interaction processes (Compton, Pair, etc.). The algorithm is split into 4 subsections: clustering (blobbing adjacent hits into one larger hit), tracking (finding showers, muons, pair events and Compton electron tracks), Compton sequence reconstruction (identifying the sequence of Compton interactions) and finally decay detection (identifying events, which might originate from decays).

# 5.6 Mimrec

Mimrecs main duty is list-mode likelihood image reconstruction. In addition it can handle other high-level data analysis, like compiling spectra, ARM-distributions, scatter-angle distribution, and more.

# 5.7 ResponseGenerator

Program to generate responses/data-space used for the Bayesian event reconstruction algorithms

#### 5.8 ResponseManipulator

Merge and view response files

#### ConvertMGGPOD 5.9

Converts fits files generated by MGeant/MGGPOD into the SIM format, which can be read by the event reconstruction programs

#### SensitivityOptimizer 5.10

Calculates the sensitivity based on a source file and several background files in a multi-dimensional data space

#### File formats 6

```
All file formats have some header information, which include the type of the file, its version, and the geometry
```

```
information: The type of the file
              <Type \in \{SIM, TRA, etc.\}>
 Type
 The version of the file
 Version
              <Version number as int>
 The geometry
 Geometry < Path to the geometry>
```

## 6.1

```
The SIM format
Start event
SE
Event ID
      <ID triggered event> <ID simulated event>
Time in seconds Linux time
TI
      <Time in sec>
Total deposited energy
     <Energy in keV>
Energy deposited in not sensitive material
      <Energy in keV>
Energy deposited in the guard ring of strip detectors
     <x position of detector center in cm>
      <y position of detector center in cm>
      <z position of detector center in cm>
      <Energy in keV>
Total energy deposit per drift chamber
     <x position of interaction in cm>
```

<y position of interaction in cm> <z position of interaction in cm>

<Energy in keV>

```
<x position of interaction in cm>
      <y position of interaction in cm>
      <z position of interaction in cm>
      <x direction>
      <v direction>
      <z direction>
      <Energy in keV>
Interaction information version 3
      <Detector ID 1=2D strip, 2=MEGA calorimeter, 3=3D strip, 4=ACS, 5=drift</p>
      chamber>
      <Type \in \{INIT, PAIR, COMP, PHOT, BREM, RAYL\}>
      <Particle number 1: Photon, 2: Positron, 3: Electron>
      <ID of this interaction>
      <ID of interaction this particle originated from>
      <x position of interaction in cm>
      <y position of interaction in cm>
      <z position of interaction in cm>
      <x direction of new particle>
      <v direction of new particle>
      <z direction of new particle>
      <Kinetic energy of new particle in keV>
Interaction information version 22
      <Type \in \{INIT, PAIR, COMP, PHOT, BREM, RAYL\}>
      <ID of this interaction>
      <ID of interaction this particle originated from>
      <Detector ID 1=2D strip, 2=MEGA calorimeter, 3=3D strip, 4=ACS, 5=drift</p>
      chamber>
      <x position of interaction in cm>
      <y position of interaction in cm>
      <z position of interaction in cm>
      <Mother particle number 1: Photon, 2: Positron, 3: Electron>
      < x direction of original particle>
      <y direction of original particle>
      <z direction of original particle>
      <x polarization of original particle>
      <y polarization of original particle>
      <z polarization of original particle>
      <Kinetic energy of original particle in keV>
      <Particle number 1: Photon, 2: Positron, 3: Electron>
      <x direction of new particle>
      <y direction of new particle>
      <z direction of new particle>
      <x polarization of new particle>
      <y polarization of new particle>
      <z polarization of new particle>
      <Interaction time in sec> (Missing in Version 21)
      <Kinetic energy of new particle in keV>
Hits
```

<Detector ID 1=D1, 2=D2, 3=D3, 4=ACS, 5=drift chamber>

HT

Electron direction information for Strip3DDirectional detectors

```
<x position of interaction in cm>
```

- <y position of interaction in cm>
- <z position of interaction in cm>
- <Energy deposit in keV>
- <Time since start of event>
- < Vector of IDs of the interactions, which contributed to this hit>

# 6.2 The GEO.SETUP format

All parameters of the MEGA detector are initialized in a setup file. Its characteristic suffix is "setup". It has an object oriented, keyword based style, for example in the beginning a detector is declared and afterwards it is initialized with several parameters.

MSiStrip Poseidon

Poseidon. Threshold 0.0 0.0 0.0 100.0 100.0 100.0

Volume WorldVolume. WorldVolume.Material Germanium WorldVolume.Shape BRIK 0.0 0.0 0.0

A complete list of all keywords can be found in chapter keywords.

In order to assure some kind of safeness the setup file is scanned four times:

- 1. Include all "Include"-files
- 2. Scan for other command keywords and object keywords
- 3. Scan for Clones of the object keywords
- 4. Assign all other parameters
- 5. Validate the input

This assures that the sequence of keywords in the file can be arbitrarily.

All dimensions of the variables are in cm, keV, g, etc.

There exist three types of keywords: Command keywords, object keywords and parameter keywords. These keywords and the whole setup file are not case sensitive.

There are several important limitations with this geometry and detector description format compared to Geant3 and Geant4:

- 1. All daughter volumes have to be completely contained in their mother volumes.
- 2. No overlapping volumes are allowed (with exception of virtual volumes)
- 3. There are only a limited number of volume types implemented so far. If you need more, let me know, but make sure, that they exist in Geant3, Geant4 and Root.
- 4. It is not possible to divide one volume in sub volumes.
- 5. All detector volumes need to be boxes (with exception of the Scintillator- and Anger Camera-type detectors)
- 6. ... and probably many more ...

# 6.2.1 Global keywords

Keyword Name <text>

Description Gives the setup a distinguishable name

Example Name MegaPrototype

Keyword Version < float>

Description Version of the current setup

Example Version 1.01

Keyword Include <file name>

Description Include the file <file name> into the setup-file. This is very useful when splitting a

large setup-file into smaller parts, e.g. defining an own file for each detector or a file containing the specific materials, etc. The file name is allowed to be a relative path

and might contain wildcards

Example Include \$(MEGALIB)/resource/geometries/materials/Materials.geo

Keyword SurroundingSphere <radius> <x-position> <y-position> <z-position>

<distance>

Description A surrounding sphere is a mandatory requirement for the simulations with GMega

or Cosima. If you do simulations in the far-field, the photons are simulated from a disk with radius <radius> pointing from a distance <distance> towards point <x-position> <y-position> <z-position>. This feigns a real far-field. Therfore, make sure you define a surrounding sphere which is as small as possible but encloses the

complete detector geometry, without intersecting any volume.

Example SurroundingSphere 25.0 0.0 0.0 15.0 25.0

Keyword // <text> or # <text>

Description Comment

Example # This is a comment

# 6.2.2 The object Material

Keyword AbsorptionFileDirectory <path>

Description Gives a default directory where to search the absorption files

Example AbsorptionFileDirectory \$(MEGALIB)/resource/geometries/materials

Keyword Material <text>

Description Defines the material <text>

Example Material CsI

 $\begin{tabular}{ll} Keyword & <& Material > . Density < float > \\ Description & The density of the volume in g/cm^3 \\ \end{tabular}$ 

Example CsI.Density 4.5

Keyword <Material>.ComponentByAtoms <A> <Z> <Number of atoms of the

component/molecule)>

Description Declares a component of a material. For example if the material is a mixture of the

two components Cs and I, they have to be declared as in the example below. The

number of atoms needs to be an integer!

Example CsI.Component 132.9 55 1 // Cs

CsI.Component 126.7 53 1 // I

Keyword <Material>.ComponentByMass <A> <Z> <Fractional mass of the component

in the material>

Description Declares a component of a material. For example if the material is a mixture of the

two components N and O, they have to be declared as in the example below. The

total fractional mass needs to be 1!

Example Air.Component 14.0 7.0 0.7 // N

Air.Component 16.0 8.0 0.3 // 0

Keyword <Material>.RadiationLength <float>

Description The radiation length of the material. This keyword is only necessary if the material

consists of only one material.

Example Silicon.RadiationLength 9.35

The directory \$(MEGALIB)/resource/geometries/materials contains a general file "Materials.geo", which includes all used materials for the MEGA geometries. The same directory contains also the absorption probability files. If you add a new material to Materials.geo, you also have to add the absorption probabilities to this directory. This is done in the following way:

- 1. Change to the GMega source directory: cd \$(GMEGA)/src
- 2. Launch GeoMega, load the geometry and create the Geant files ugeom.f and detinit.f (Geomega:  $Menu: Analysis \rightarrow Create\ Geant3\ files$ )
- 3. Switch to the main GMega directory and compile: cd \$(GMEGA); make
- 4. Switch to the directory, where you want to have your absorption files, e.g. \$(MEGALIB)/resource/geometries/materials
- 5. Run the absorptions program: absorptions
- 6. Now all absorption files are generated and you are done!

# 6.2.3 The object Volume

Please take notice of all distances except radii being half distances!

Keyword Volume <text>

Description Declares a volume with name <text>

Example Volume Wafer

Keyword <Volume>.Shape <BRIK> <half x-size> <half y-size> <half z-size>

 $\label{eq:conditional_equation} \textbf{Keyword} \qquad < \textbf{Volume} > \textbf{.Shape} \ < \textbf{SPHE} > \ < \textbf{radius min} > \ < \textbf{radius max} > \ < \textbf{theta}$ 

max> <phi min> <phi max>

Keyword <Volume>.Shape <TUBS> <radius min> <radius max> <half height> <phi min>

<phi max>

 $\label{eq:cone} \textbf{Keyword} \qquad < \textbf{Volume} >. \textbf{Shape} < \textbf{CONE} > < \textbf{half height} > < \textbf{bottom radius min} > < \textbf{bottom radius}$ 

max> <top radius min> <top radius max>

 $\label{eq:conditional} \textbf{Keyword} \qquad < \textbf{Volume} > . \textbf{Shape} < \textbf{TRD1} > < \textbf{half distance x1} > < \textbf{half distance x2} > < \textbf$ 

y> <half distance z>

Keyword <Volume>.Shape <TRD2> <half distance x1> <half distance x2> <half distance

y1> <half distance y2> <half distance z>

Keyword <Volume>.Shape <TRAP> <half distance in z> <theta> <phi> <half height

trapezium bottom> <half bottom length trapezium bottom> <half top length trapezium bottom> <alpha trapezium bottom> <half height trapezium top> <half bottom length trapezium top> <alpha trapezium to

top>

Keyword <Volume>.Shape <GTRA> <half distance in z> <theta> <phi> <twist> <half

height bottom trapezium bottom> <half bottom length trapezium bottom> <half top length trapezium bottom> <alpha trapezium bottom> <half height trapezium top> <half bottom length trapezium top> <half top length trapezium top> <alpha

trapezium top>

Keyword <Volume>.Shape <PCON> < the azimuthal angle phi at which the volume begins

(angles are counted counterclockwise)> <opening angle of the volume> <number of sections, number should be at least 2, the following three arguments are repeated

accordingly> <height> <minimum radius> <maximum radius>

Keyword <Volume>.Shape <PGON> < the azimuthal angle phi at which the volume begins

(angles are counted counterclockwise)> <opening angle of the volume> <number of sides of the cross section between the given phi limits> <number of sections, number should be at least 2, the following three arguments are repeated accordingly>

<height> <minimum radius> <maximum radius>

Description Describes the shape of the object.

BRIK: A box SPHE: A sphere TUBS: A cylinder CONE: A cone PCON: A Polycone PGON: A Polygon

TRD1, TRD2, TRAP, GTRA: trapezoids (try to avoid TRAP and GTRA!)

Detailed descriptions of the objects can be found in the Geant3, Geant4 or ROOT manual (ROOT comes with pictures!). The argument list corresponds to that of ROOT. If you want new volumes to be added, let me know, but make sure such a volume exists in Geant3, Geant4 and ROOT and provide an equation for its volume! One limitation for the current Geant4 implementation is that TRAP and GTRA are not allowed to be triangles (i.e. non of the values height, bottom and top length is allowed to be zero). So if you need triangles make sure, they contain very small

values.

Example Wafer.Shape BRIK 0.0 0.0 30.0

GeLeft.Shape TRAP 3.5 0 0 0.86602540378 3 0.0000000000000 0

0.86602540378 3 0.0000000000000 0

Keyword <Volume>.Material <Material name>

Description Material of the volume. It has to be defined somewhere else in the setup file.

Example Wafer.Material Silicon

Keyword <Volume>.Visibility <0 or 1>

Description If the value is zero, then the volume is not visible, if it is 1, the volume is visible in

the geomega viewer.

Example Wafer. Visibility 1

Keyword <Volume>.Color <int>

Description The color of the volume when display in the viewer. The numbers correspond to the

ROOT color IDs.

Example Wafer.Color 5

Keyword <Volume>.Mother <Name of volume>

Description In this mother volume the volume will be placed. Attention: the volume must be

fully contained in her mother and the mother must be defined somewhere else in the

setup file. For the world volume

Example Wafer.Mother Tracker

Keyword <Volume>.**Position** <x> <y> <z>

Description Position of the volume within its mothers coordinate system. Position is the center

of the volume. Attention the volume has to be fully contained in its mother volume!

Example Wafer.Position 0.0 0.0 30.0

Keyword <Volume>.**Rotation** <x> <y> <z>

Description Counterclockwise rotation of the volume around x-, y-, and z-axis in the mother

volume. Attention the volume has to be still fully contained in its mother volume!

Example Wafer.Rotation 0.0 0.0 90.0

Keyword <Volume>.Rotation <theta1> <phi1> <theta2> <phi2> <theta3> <phi3>

Description Rotation according to the Geant3 convention: <theta1> is the polar angle of the

> x-prim axis in the main reference system (MRS), <theta2> and <theta3> have the same meaning for the y-prim and z-prim axis. <phi1> is the azimuthal angle of the x-prim in the MRS and <phi2> and <phi3> have the same meaning for y-prim and

z-prim. Attention the volume has to be still fully contained in its mother volume!

Example Wafer.Rotation 90 90 0 0 90 0

Keyword <Volume>.Scale <double>

Description Scale (shrink or enlarge) a volume and all sub volumes. Useful if one has a large

volume tree and wants to modify it's size. This keyword can not be applied to

Copies/Clones.

Example SpaceCraftBody.Scale 0.5

#### 6.2.4**Detector objects**

There exist 7 different detector types: A 2D strip detector like the MEGA Silicon wafers (2DStrip), a 3D strip detector (3DStrip) like the NCT Germanium detectors, directional 3D strip detectors, where some information of the electron direction is retained, a Drift Chamber detector including capabilities for light sensing - needed for liquid Xe and gas microwell detectors, a calorimeter like in MEGA and a one-volumetype detector, which can be used as MEGA ACS as well as SPI Germanium detector, or any other thumb detector, which can only measure energy information, and finally a 3D voxel detector.

#### The detector keywords 6.2.4.1

Keyword Strip2D <detector name>

Description Declares a 2d strip detector with name <detector name>. The strips are always

oriented in x and y direction. If you want another orientation, simple rotate the

detector volume.

Example Strip2D bachus

Keyword Strip3D <detector name>

Description Declares a 3D strip detector with name < detector name >. It inherits all capabilities

from Strip2D. In addition it has a depth resolution, which is always the z direction.

Example Strip3D nct1

Keyword Strip3DDirectional <detector name>

Description Declares a 3D strip detector with name <detector name>. It inherits all capabilities

from Strip3D. In addition it has a directional resolution: It can detect all directions of electrons originating from Compton interaction. Due to limitations in the simulation, it is currently not possible to detect the direction of electrons simply passing through the silicon layer. If you want to use this detector, you need full "IA" information from the simulation. Thus if you use mggpod, make sure to use INIT2 and ACT2 options.

Example Strip3DDirectional SiWafer

Keyword DriftChamber <detector name> Description Declares a drift chamber detector with name < detector name >. It inherits all capa-

bilities from Strip3D. In addition, specific information like the (optical) light speed in this material ("LightSpeed"), the light sensitive detector side ("LightDetectorPosition") and its energy resolution ("LightEnergyResolution") as well as the electron

drift parameters ("DriftConstant", "EnergyPerElectron") can be set.

Example DriftChamber Chamber

Keyword Calorimeter < detector name >

Description Declares a calorimeter with name <detector name>

Example Calorimeter Fortuna

Keyword Scintillator < detector name >

Description Declares a large, one channel and non position sensitive scintillation detector with

name <detector name>. The special feature of this volume is, that it does not need to be box-like and that it can consists of multiple volumes. A detector of this type can also be used to model the SPI-Germanium detectors. Hits in this detector are

always centered.

Example Scintillator SPI12

Keyword Voxel3D < detector name >

Description Declares a box-shaped volume consisting of voxels in all three dimensions. This de-

tector has no known real-world counterpart, as it has no passive material (electronics, connectors, etc.) between the voxels. Its primary function is currently simulation

diagnostics. Hits in this detector are always centered.

Example Scintillator SPI12

6.2.4.2 Common keywords

Keyword < Detector>.SensitiveVolume < name of volume>

Description Volume, in which positions and energies of interactions are measured. Typical exam-

ples are one CsI-crystal or one single Si-wafer of the MEGA prototype.

Example MEGACal.SensitiveVolume CsICrystal

Keyword < Detector>. DetectorVolume < name of volume>

Description A larger volume which contains several evenly spaced sensitive volume. The position

of the sensitive volumes are specified in the "Structural" parameters. They must be identical with the positions given in the volume description! Typical examples are the MEGA calorimeter, which consists of 120 CsI crystal or one layer of the MEGA tracker, which consists of 9 Si-wafers. The sensitive volume is either the same as the detector volume or entirely and **unrotated** contained in the detector volume!

Example MEGACal.DetectorVolume CsIDetector

Keyword <Detector>.StructuralPitch <x> <y> <z>

Description Spacing between the sensitive volumes (distance between the end of last sensitive

volume to the start of the next sensitive volume). This is not the pitch between the individual strips of a strip detector! If you have a tracker then also add the distance

between the layers as z-component!

Example MEGACal.StructuralPitch 0.07 0.07 0.0

Keyword < Detector>.StructuralOffset <x> <y> <z>

Description Distance between the edge of the detector volume to the beginning of the first sensitive

volume. Calculated from negative to positive axis!

Example MEGACal.StructuralOffset 0.235 0.235 0.185

Keyword <Detector>.NoiseThreshold <double>

Description All hits in one voxel of the detector which are below this energy (in keV) are assumes

not to be measured.

Example MEGACal.NoiseThreshold 50

Keyword <Detector>.TriggerThreshold <double>

Description A hit need to deposit at least this energy (in keV) to raise a trigger signal.

Explanation The difference between noise and trigger threshold: Each strip has a certain amount

of electronics noise. So when one reads it out, one gets a signal in ADC counts, which only reflects the noise of the electronics. So normally one only uses hits which are well above the noise, e.g. 5 sigma above it. Trigger threshold is something different. A hit needs to produce a certain voltage (i.e. has to deposit a certain energy) to initiate

the read-out of the detector.

Example MEGACal.TriggerThreshold 100

Keyword <Detector>.EnergyResolutionAt <energy> <measurements error>

Description Energy resolution at the given energy in keV (one sigma)

Example MEGACal.EnergyResolution 662 60

Keyword < Detector > . Failure Rate < double >

Description Random failure rate – values are between 0 (no failures) and 1 (complete loss of

detector)

Example MEGACal.FailureRate 0.1

# 6.2.4.3 Keywords specific to calorimeters

Keyword <Detector>.DepthResolution <energy in keV> <noise in cm (one sigma)>

Description Give the depth resolution at the given energy in cm (one sigma). If this keyword is

not given, then no depth resolution is assumed, i.e. it behaves like a Strip2D detector.

Example MEGACal.DepthResolution 500 1.2

# 6.2.4.4 Keywords specific to strip detectors (Strip2D, Strip3D, Voxel3D, DriftChamber) .

Keyword  $\langle \text{Detector} \rangle$ . **Offset**  $\langle x \rangle \langle y \rangle \langle z \text{ (Voxel3D only)} \rangle$ 

Description Distance between the edge of the sensitive detector to the beginning of the first strip

Example SMegaSD.Offset 0.142 0.142

Keyword  $\langle \text{Detector} \rangle$ . **StripNumber**  $\langle x \rangle \langle y \rangle \langle z \text{ (Voxel3D only)} \rangle$ 

Description Number of strips

Example SMegaSD.StripNumber 128 128

Keyword <Detector>.GuardringTriggerThreshold <double>

Description A hit in the gurad ring needs to deposit at least this energy (in keV) to raise a trigger

(veto) signal.

Example SMegaSD.GuardringTriggerThreshold 100

Keyword < Detector>.GuardringEnergyResolutionAt < energy> < measurements error>

Description Energy resolution at the given energy in keV (one sigma) for the guard ring

Example SMegaSD.EnergyResolution 100 10

For 3D strip detectors (with and without electron direction resolution) and DriftChambers there is another special keyword:

Keyword <Detector>.DepthResolution <Energy in keV> <Resolution in cm (one sigma)>
Description Set the one sigma depth resolution. If this keyword is not given, then no depth

resolution is assumed.

Example GeStrip.DepthResolution 200 0.2

Keyword <Detector>.DepthResolutionThreshold <Energy in keV>

Description Below this threshold no depth resolution can be measured, i.e. the center (z-axis) of

the detector is used as z-position.

Example GeStrip.DepthResolutionThreshold 25

For 3D strip detectors with directional resolution exists another special keyword:

Keyword < Detector>. Directional Resolution < Energy in keV> < Resolution in degree (one

sigma)>

Description Set the one sigma directional resolution.

Example SiStrip.DirectionalResolution 200 30

DriftChambers have additional keywords

Keyword < Detector>.LightSpeed < light speed in cm/s>

Description Light speed of the scintillation light in the drift chamber

Example Chamber.LightSpeed 18E+9

Keyword <Detector>.LightDetectorPosition <0: none; 1: +x; -1:-x; 2: +y; -2 -y; 3: +z;

-3:-z>

Description Represents the side of the detector which is light sensitive (i.e. equipped with PMTs

or diodes). If this value is zero, then no light detector is assumed.

Example Chamber.LightDetectorPosition 3

Keyword < Detector > . DriftConstant < constant in cm >

Description One sigma of the opening cone of the drift in the E-field  $(c \times \sqrt{Distance\ to\ wires})$ .

If this is zero, a simple and fast projection is used.

Example Chamber.DriftConstant 0.01

Keyword < Detector > . EnergyPerElectron < Energy in keV >

Description Energy of one drifting electron Example Chamber.DriftConstant 0.01

Keyword < Detector>.LightEnergyResolutionAt < Energy> < Resolution in keV (one

sigma)>

Description One sigma energy resolution of the detected light.

Example Chamber.DriftConstant 0.01

# 6.2.4.5 Keywords specific to ACS-like detectors

The special feature of this detector type is that it may contain several sensitive volumes of different shapes.

Keyword <Detector>.**HitPosition** <Volume name 1> <Volume name 2> <x> <y> <z> Description A hit in volume <Volume name 1> is moved by x, y, z into <Volume name 2>. All

sensitive volumes of the detector have to be covered!

Example GeDet.HitPosition GeCentral GeCentral 0 0 0

# 6.2.5 Triggers

The current implementation of triggers is not completely finished and not obvious throughout the package: It has only been tested for MEGA type triggers (D1+D2) and the simulation does not necessarily take into account all options: While GMega uses part of this information (it only translates the info to "TriggerByChannel" and "DetectorType", which is the more open trigger and GMega ignores vetoes and trigger thresholds), Cosima has in its setup file an own trigger condition. The approach is the following: First define if this trigger is a veto or not. Then define if the triggering happens by channel or by detector. "TriggerByChannel" means that the channels are counted for the trigger, "TriggerByDetector" means that the detectors are counted (irrelevant of how many channels in this detector have trigger - if at least one is above the trigger threshold). Then define the triggering detectors (by type or name) and the number of hits they must accumulate. Pay attention, you can only have one type of detector keyword per trigger, i.e. one trigger is allowed to contain only "DetectorType" keywords or only "Detector" keywords or only "GuardringDetector" keywords!

Keyword Trigger <text>

Description Declares a trigger condition with name <text>

Example Trigger D1D2

Keyword <Trigger>.Veto <true/false>

Description Determines if this is a real trigger (acceptance) or a veto trigger (rejection). If this

option is not given then veto is false!

Example ACSVeto.Veto true

Keyword <Trigger>.TriggerByChannel <true/false>

Description Determines that the hits are accumulated channel wise. This is the default.

Example D1D2.TriggerByChannel true

Keyword <Trigger>.TriggerByDetector <true/false>

Description Determines that the hits are accumulated detector wise. TriggerByChannel is the

default

Example D1D2.TriggerByChannel true

Keyword <Trigger>.DetectorType <Detector type name> <Number of hits>

Description <Number of hits> are necessary in detector type <Detector type name> to raise a

trigger. If this keyword occurs multiple times all conditions have to be fulfilled. The <Detector type name> follows the MEGAlib convention: Strip2D is a 2D strip detector, Calorimeter is a MEGA calorimeter, Strip3D is a 3D strip detector, Scintillator is a scintillator/ACS type detector and DriftChamber is of course the

drift chamber.

Example D1D2.DetectorType Strip2D 1

D1D2.DetectorType Calorimeter 1

Keyword <Trigger>.Detector <Detector name> <Number of hits>

Description <Number of hits> are necessary in <Detector name> to raise a trigger. If this

keyword occurs multiple times all conditions have to be fulfilled.

Keyword <Trigger>.GuardringDetectorType <Detector type name> <Number of hits>

Description <Number of hits> are necessary in the guard ring of detector type <Detector type

name> to raise a trigger. If this keyword occurs multiple times all conditions have to be fulfilled. The <Detector type name> follows the MEGAlib convention: Strip2D is a 2D strip detector, Calorimeter is a MEGA calorimeter, Strip3D is a 3D strip detector, Scintillator is a scintillator/ACS type detector and DriftChamber is of

course the drift chamber.

Example D3.GuardringDetectorType Strip3D 1

Keyword <Trigger>.GuardringDetector <Detector name> <Number of hits>

Description <Number of hits> are necessary in the guard ring <Detector name> to raise a trigger.

If this keyword occurs multiple times all conditions have to be fulfilled.

Example D3.GuardringDetector MyStrip3D 1

# Here are more examples:

The MEGA prototype has an electron tracker and a calorimeter. Thus, a reasonable trigger condition would require at least two layers of the tracker and one calorimeter triggering. In addition any events with hits in the veto dome should be rejected. Thus we define two trigger conditions:

Trigger Main
Main.Veto false
Main.TriggerByDetector true
Main.Detector MyD1 2
Main.Detector MyD2 1

Trigger AntiCoincidence
AntiCoincidence.Veto true
AntiCoincidence.TriggerByDetector true
AntiCoincidence.Detector MyAnticoidence 1

A thick Germanium detector might require at least three hits for Triple Compton coincidence. Since the detector is thick, those hits do not need to be in different detectors but only different channels need to trigger. In addition, events which deposit energy in the guard rings of the detector are going to be rejected:

Trigger Main
Main.Veto false
Main.TriggerByChannel true
Main.DetectorType Strip3D 3

Trigger Guardring
Guardring.Veto true
Guardring.TriggerByDetector true
Guardring.GuardringDetectorType Strip3D 1

The triggering is performed during reading the events from the \*.sim file into revan or sivan. No triggers are tested for the \*.evta files, since those events represent detector data. A trigger is raised, when the deposited energy in the given channel is above the trigger threshold of the detector or guard ring.

# 6.2.6 Additional features

# 6.2.6.1 Templates and the keyword Copy .

In a usual detector geometry some volumes will appear several times. To avoid any copy and paste, the keyword Copy has been introduced, which copies the characteristics of one volume to another. Normally the base volume is defined as template, i.e. it does not have a mother and it is not positioned.

Volume GeWafer

GeWafer.Material Germanium

 ${\tt GeWafer.Visibility}\ 1$ 

GeWafer.Color 6

GeWafer.Shape BRIK 4.0 4.0 1.0

// Arrange the Ge-detectors

GeWafer.Copy GeE1N001

GeE1N001.Position 0.0 0.0 3.75

GeE1N001.Mother WorldVolume

GeWafer.Copy GeE1N002

 ${\tt GeE1N002.Position~0.0~0.0~1.25}$ 

GeE1N002.Mother WorldVolume

Up to now, the keyword Copy is only implemented for volumes!

Keyword <Volume>.Copy <existing volume>

Description Create the new volume < existing volume > and copy all characteristics of < Volume >

to the new volume

Example SiLayer.Copy bachus

# 6.2.6.2 The keyword Virtual .

One major restriction of the geomega format is that no overlapping volumes are allowed. But sometimes it is useful to have a larger volume, in which several other volumes are grouped, to use as template. In the case this larger volume has no real meaning other than being a container for other volumes, it can be declared as virtual. Those virtual volumes are allowed to overlap with other volumes – as long as their content does not overlap with any other volume! During the creation of the geometry, virtual volumes are removed from the volume tree!

Keyword <Detector>.Virtual <br/>bool><br/>Description Declare a volume as virtual

Example TrackerContainer.Virtual true

# 6.2.6.3 The keyword Constant

A very experimental feature is that one can define constants in the setup file:

Keyword < Constant> < string: Replace this> < string: with this>

Description Replace some text with another text

Restrictions It is not checked if the string is a token, it is even replaced if it is part of another

string. Don't use any keywords!

Example Constant Size 2.0

Example Wafer.Shape BRIK Size Size 0.025

#### 6.2.6.4 The maths environment.

Another very experimental feature is that one can do very basic mathematical calculations in the setup file. The signs { and } start and end the maths environment. Everything within those brackets is considered as one token:

```
Constant Size 2.0
Wafer.Shape BRIK {1.2*Size} {1.2*Size} {0.5*(log(Size)-1)}
```

The maths environment relys on the ROOT interpreter. So whatever manipulation ROOT knows can be done in the maths environment.

# 6.2.7 Closing remarks

Before you start coding a geometry, take a look at the examples in the directory \$(MEGAlib)/resources/geometries. Try to write your code as close as possible to the examples, in order to avoid any trouble.

It is very time-consuming to protect against each possible input error in an open file format like this. So do not expect to get a warning/error message for each time you make a mistake – sometimes the program will warn you with an error message, sometimes it will simply accept the error and produce false results and sometimes it will simply crash. Whenever you come across such a problem, let me know and I will implement a protection.

Something which is very difficult to implement is a check for overlapping volumes – so this is not done at the moment. Therfore it is the users responsibility to make sure no volumes overlap.

In addition, since most of the input parameters need to be positive and almost no sanity checks for positivity are implemented, it is the users responsibility to ensure positive values where reasonable.

# 6.2.8 Practices for writing a good geometry

- Use meaningful descriptions, e.g. WorldVolume instead of VAC0
- Don't make your geometry flat! The more daughters a volume has, the more volumes have to be searched when a particle moves from one volume to the next! It i significantly better to have a steep geometry: The world volumes contains daughters, which contain daughter volumes, which contain daughter volumes, etc.
- Use multiple files representing different objects: If you have distinguishable objects such as individual detectors with all their surround electronics and mounting, describe them in an individual file! Simply "Include" this file in the file where its mother volume is described. This makes your geometry much more clearly laid out.

# 6.3 The TRA format

# 6.3.1 Common key words

```
Start event
SE
```

Event type

ET 
$$\langle \text{Type} \in \{PA, CO\} \rangle$$

Event ID (same as in SIM)

Time in seconds Linux time

TI <Time in sec>

Time walk between D1 and D2 trigger in ns - only for real MEGA data TW  $\,$   $\,$   $\,$   $\,$   $\,$   $\,$  Time in ns  $\,$ 

x- and z-axis of orientation of detector (spherical coordinates)

RX 
$$< x > < y > < z >$$
 (Default: 0 1 0)

RZ 
$$< x > < y > < z >$$
 (Default: -1 0 0)

Orientation of instrument in galactic coordinates

GX <Longitude in degree> <Latitude in degree>

GZ <Longitude in degree> <Latitude in degree>

# 6.3.2 Compton events

```
Energies in keV:
      <Energy of scattered gamma-ray>
      <Energy error of scattered gamma-ray>
      <Energy of recoil electron>
      <Energy error of recoil electron>
Positions and directions of the first interaction
     <x position of first interaction in cm>
      <y position of first interaction in cm>
      <z position of first interaction in cm>
      <x position error of first interaction in cm>
      <y position error of first interaction in cm>
      <z position error of first interaction in cm>
      <x position of second interaction in cm>
      <y position of second interaction in cm>
      <z position of second interaction in cm>
      <x position error of second interaction in cm>
      <y position error of second interaction in cm>
      <z position error of second interaction in cm>
      <Electron direction x>
      <Electron direction y>
      <Electron direction z>
      <Error of electron direction x>
      <Error of electron direction y>
      <Error of electron direction z>
Length of Compton sequence in "detected interactions"
      <Length (1, 2, 3, etc.)>
First and second Compton quality factor (aka. "Compton test statistics")
      <Probability best sequence> <Probability second best sequence>
Length of first track
     <Length (1, 2, 3, etc.)>
Deposited energy of first track in its first layer
     <Energy in keV>
Shortest distance in the sequence (not first distance)
LA
      <Distance in cm>
Time-of-flight between first and second interaction
      <Time-of-flight in s> <Uncertainty of time-of-flight in s>
COMPTEL PSD value
PD <PSD a.u.> <Uncertainty PSD a.u.>
```

# 6.3.3 Pair events

Position of the gamma conversion

PC < x in cm > < y in cm > < z in cm >

Energy and initial direction of the first generated particle

PE <Energy in keV> <Energy error in keV> <Direction x> <Direction y> <Direction z>

Energy and initial direction of the second generated particle

PP <Energy in keV> <Energy error in keV> <Direction x> <Direction y> <Direction z>

Energy deposit in the initial layer

PI <Energy in keV>

# 7 ConvertMGGPOD

# 7.1 Command line options

Mandatory:

-f <filename> Load this fits file (\*.fits) (this command can appear several times!)

-g <filename> Load this geometry file

Optional:

-o <filename> Name of the output file. Per default \*.sim is appended to the base name of the fits

file.

-m Sets the maximum number of steps, which the fits events may have per detector type

before the discretization (default: 5000)

-s <int> Version number of the generated \*.sim file. Default is 3, for Bayesian event recon-

struction you need 22.

-i Ignore hits which consist of only one hit

-use-nohits-events Writes output also for those events, which do not have a hit

-ignore-veto To reduce the file size, events which produced a veto are per default thrown away.

Use this flag ignore all vetos. (if this flag is NOT set, the veto is applied at this step, using the energy threshold for the anticoincidence detector(s) given in the geometry file. NO noising is applied to the veto — so the result is a step-function veto threshold at the nominal threshold energy! Later, revan will apply noising and another veto

decision to the remaining events!)

-no-discretization Do not do the discretization of hits into voxels

-no-gui Avoids the progress bar screen

-d Dump the content of the fits file. Special option, which ignores everything but -f &

-g!

# 8 GeoMega - Geometry for Mega

# 8.1 Command line options

Geomega is a GUI based application. Thus, you do not need to give command line options. In fact only the most frequently used options are implemented as special tasks.

Optional:

-g or -geometry <filename> Load this geometry file (\*.geo.setup)

-d Activates debug mode

-s or -startvolume < name> Use this volume as world volume. If this volume exists as several

copies, use one of the Copy names!

Special tasks:

-create-mggpod <filename suffix> Create mggpod files with this file name suffix

Create mggpod files with the default file names (setup.geo, media.med, materials.mat)

# 9 Revan - Real event analyzer

# 9.1 Command line options

Revan is a GUI based application. Thus, you do not need to give command line options. In fact only the most frequently used options are implemented as special tasks.

Optional:

-f <filename> Load this sim or evta file (\*.sim, \*.evta)
-g <filename> Load this geometry file (\*.geo.setup)

-d Activates debug mode

-no-gui Avoids the progress bar screen

Special tasks:

-a <filename> Load this configuration file and start the reconstruction of the events. If -f and

-g are given, then those override the values of the configuration file.

# 9.2 Graphical user interface

Menu File:

Open a simulation (\*.sim) or events (\*.evta) file

Load Geometry
Load the geometry, which belongs to the simulation or events file.
Load Configuration
Load a configuration file, which contains all GUI information.
Save Configuration file, which contains all GUI information.

Exit

List of previously used revan files

Menu Reconstruction:

Selection of algorithm

Start Event Reconstruc- Start

tion

nstruc- Starts the event reconstruction

Selects different reconstruction algorithms – the following four menus will have

different contents for different algorithms

Clustering options No clustering:

No options

Clustering by distance:

Enter the maximum distance between two interactions in cm up to which to

hits are combined to a cluster

Use center of the cluster (instead of each sub hit individually) as reference point for distance calculations: In default mode, if one has already a cluster to which an additional hit should be added, then all sub-hits are checked if the new hits is within the distance. If the option is checked, then new hits have to be within a minimum distance radius around the center of gravity/energy of the cluster

to be added to the cluster.

Clustering of adjacent voxels:

Select the numbers of neighbors (4 closest neighbors or 8 clostest neighbors) of the adjacent voxels. The clusterization can be aborted if the z-distance is larger than a certain sigma of the position resolution. Ignore this if the sigma is negative.

Tracking options No tracking:

No options

General options for all tracking algorithms:

Check if the algorithm should check for pair events, MIPs or Compton electron tracks.

Maximum layer jump for Comptons offers to set the maximum allowed distance of two hits (in layer distances). One means the hits have to be in consecutive layers, two means that there is at maximum one layer between the hits.

Maximum number of different track sequences to keep This means that not only the best, but the two, three, four, five, best track sequences are transferred to CSR. For MEGA this did not turn out to improve the performance (but worsens it), but is intended for thick Silicon trackers for handling two layer interactions. Reject purely ambiguous sequences Do not try to analyze tracks which have not any straight beginning, but play ping-pong between the layers, or which curl. Those tracks are more likely to be wrongly reconstructed, but at low energies there are a lot of them...

# Bayesian electron tracking

Give the  $\it file \ containing \ the \ tracking \ data$  . This has to be the file ending with \*.goodbad.rsp

# No Compton tracking

No options

# Compton sequence reconstruction without energy recovery

Use two site events without track means to use events which consist of nothing but two hits

Only use events which start in 2D strip detectors rejects events which start in anything else but a 2D strip detector

Reject events which are limited to one detector type rejects events which do not interact at least in two different detector types like 2D strip detector and 3D strip detector or 2D strip detector and calorimeter

Assume undecided events start in 2D strip detectors (tracker) assumes for all 2 site events, whose direction of motion can not be determined, that they start in a silicon strip detector

Always assume events starts in the 2D strip detectors (tracker) always forced 2 site events to start in the tracker,

Allowed range of Compton quality factor rejects events whose final quality factor is not within the limits

Maximum number of interactions: Since the complexity of the algorithm goes with the factorial of the number of interactions, we have to cut it off here.

All Comptons have to originate from objects in this file checks if the Compton cone intersects volumes in the given geometry file. Only those events are kept.

# Compton sequence reconstruction with energy recovery tbd.

# Bayesian Compton tracking

Give the *file containing the BCT data*, i.e. the files containing the Bayesian matrices. You only need to give only file, the others are assumed to reside on the same directory. This has to be the file ending with \*.goodbad.rsp

Maximum number of interactions: Since the complexity of the algorithm goes with the factorial of the number of interactions, we have to cut it off here.

# Compton sequence reconstruction with time of flight

tbd.

No decay detection

No options

Decay detection for Comptons after Compton tracking

File containing decay lines wants the file, which contains the individual energies of the decays, for which the algorithm should search.

Sequencing options

Decay option

General event selections Select events by their total deposited energy, their minimum lever arm between

any two Compton interactions or event ID (negative numbers means no ID

limit)

Menu Applications:

Initial energy distribution Shows the energy distribution before the event reconstruction

Energy distribution be- Compares the energy distribution before and after the event reconstruction

fore/after

Energy per cluster per de- For each detector type show a histogram of how many energy per cluster is

tector type deposited

Spacial hit distribution Shows a 3D hit distribution Number of clusters Dumps a cluster distribution

# 10 Mimrec - Mega image reconstruction

# 10.1 Command line options

Optional:

-f <filename> Load this sim or evta file (\*.sim, \*.evta)
-g <filename> Load this geometry file (\*.geo.setup)
-c <filename> Load a configuration file (\*.cfg)

-d Activates debug mode

Special tasks:

-i Create an image.
-s Create a spectrum
-a Create an ARM

-e Dump the event selections

Additional options for the

special tasks

-n or -no-gui Do not use a graphical user interface

-k or -keep-alive Do not quit after executing a batch run, if we do have a gui

# 10.2 Graphical user interface

Menu File:

Open a simulation (\*.sim) or events (\*.evta) file

Load Geometry
Load the geometry, which belongs to the simulation or events file.
Load Configuration
Load a configuration file, which contains all GUI information.
Save Configuration
Save a configuration file, which contains all GUI information.

Exit

List of previously used revan files

Menu Backprojection:

Coordinate system Choose between the three different coordinate systems: Spherical, Cartesian

2D and Cartesian 3D

Image dimension Choose the range of the different axes and the number of bins (Do not modify

the image axis vector!)

Backprojection algorithm Choose between the backprojection to points and areas. The shape of the area

is controlled by later GUIs

Response selection Only Universal adaptable Compton response is currently working

Response parameters Give the length (longitudinal) of the Compton cone (for tracks only) and the

width (transversal) of the Compton cone as well as the width of the pair response - all values are 1 sigma. If you want to use absorptions probabilities be prepared to wait for the result... The fit parameter file is currently not working.

Not working

Event selections Choose from a wide range of different event selections. For somebody with

a basic knowledge of Compton scattering/Compton telescopes most of them should be obvious. The only sophisticated one is the earth horizon cut: You have the choice between: (1) No cut (2) A cut which rejects all events which intersect the horizon - this cut does not differentiate between tracked and not tracked events. Tracked events are assumed to be a full circle. (3) A probability based cut. It is calculated how likely it is that the event came from earth. If this probability exceeds the given value (GUI: maximum earth probability), then the event is rejected. For not tracked events this probability is determined by the portion of the cone which came from below the horizon. For tracked events this is a little bit more complicated and needs a response file, which has to be given via the GUI: The response file is basically a 2D-distribution, Energy versus SPD, and allows to determine the portion of the arc, which is below the horizon. If no response file is given in probability mode, we default to untracked mode! The location of the horizon is given by the last GUI element: "Angle between direction of earth center and horizon". In the GUI, the earth is currently fixed to be at the nadir of the detector (although everything else can already handle any earth position - theoretically). If you enter 0 deg in the GUI then you basically have no earth horizon, if you enter 90 deg then the horizon is your lower hemisphere.

Memory management You normally want only use 1 byte for the response matrix

Menu Likelihood:

Sensitivity

Algorithm OS-EM is broken, so use Classic EM. As stop criteria you can use the number

of iterations (recommended) or a minimum allowed likelihood change.

Penalty Broken

Menu Reconstruction:

Start image reconstruc- Nomen est omen

tion

Stop image reconstruction Nomen est omen

Menu Response: If somebody doesn't understand the menu names, he is not worth working

with mimrec! To all those function here the event selections set in Menu:  $Backprojection \rightarrow Event selection$  apply. Some points (angular resolutions, energy spectra) open an additional window where the position of the original source can be set and a window (radius) around this position. If that window

is negative, it is ignored.

# 11 SensitivityOptimizer

Calculates the point source sensitivity based on a source file and several background files in a multidimensional data space. The result (point of best sensitivity in the data space) dumped to a log file.

# 11.1 Command line options

General options:

-n <name> name (suffix) for the output files

-s <file> <sim photons> <sim area> source file, including the number of simulated photons and their start area -b <file> <time> background file name and measurement time -c < file >Mimrec Configuration File: This has to be an file, with no event restrictions in Compton Sequence Length, Earth horizon cut, Compton scatter angle, Compton quality factor, ARM and Energy! -w < min > < max >use a larger energy window for collection of background events (in keV) -csl <min low edge> <min high edge> Compton sequence length: the minimum varies between its <max> low and high edge, the maximum is fixed to max. "2 2 2" means there is one bin representing two-site events, "2 2 7" means there is one bin representing hits with a length between 2 and 7, "2 7 10" we have six bins, representing the lengths 2 to 10, 3 to 10, 4 to 10, 5 to 10, 6 to 10 and 7 -ehc < min > < max > < steps >Earth horizon cut in deg (0..min..max). Null means no cut -phi < min > < max > < steps >Compton scatter angle in deg (0..min..max) -cqf < min > < max > < steps >Compton quality factor (0..min..max) -pos < theta > < phi >Position of the source -arm < min > max > steps >ARM in deg (Radius: 0..min..max)  $-\mathrm{spd} < \min > < \max > < \mathrm{steps} >$ SPD in deg (Radius: 0..min..max) - not necessary for not tracked events! -egy < mean > < min > < max > < steps >Energy in keV (Radius: 0..min..max) Print this help -h

# 11.2 Example

SensitivityOptimizer

- -s MySims/Source847bl/Source847bl.tra 2000000 41547.56
- -b MySims/AlbedoPhotons/AlbedoPhotons.tra 2759.52
- -b MySims/CosmicPhotons/CosmicPhotons.small.tra 6441.8
- -b MySims/CosmicProtons/CosmicProtons.prompt.tra 376.752
- -b MySims/CosmicProtons/CosmicProtons.activation.tra 12000.0
- -b MySims/TrappedProtons/TrappedProtons.activation.tra 16000.0
- -c Open.cfg
- $-\operatorname{csl} 3\ 3\ 7 \operatorname{ehc}\ 0\ 100\ 6 \operatorname{phi}\ 80\ 140\ 4 \operatorname{cqf}\ 0.4\ 1.0\ 7 \operatorname{pos}\ 0\ 0 \operatorname{arm}\ 1.0\ 2.0\ 6 \operatorname{spd}\ 10.0\ 30.0\ 5 \operatorname{egy}\ 847\ 16.0\ 16.0\ 1$
- -n 847blSens

# 12 How-to guides

# 12.1 Use Case 1

This example shows the data analysis path from simulation to imaging for two  $^{88}Y$  sources located on top of the iglu of the MEGA prototype.

# 12.1.1 Generation of a geometry file for GMega

• Run geomega:

- Load the geometry file: Menu: File → Open and select the geometry Prototype.geo.setup from the directory <path to MEGAlib>/resource/geometries/simplifiedprototype. Alternatively use the command line switch "geomega -f <file name>" to load the geometry.
- ullet View the geometry:  $Menu: Analysis \to View geometry$ . If you don't see the geometry, check the command line for errors and correct them!
- Generate the Geant3 geometry information with Menu: Analysis → Create Geant3 files. This generates the files ugeom.f and detinit.f. Here ugeom.f contains the geometry information and detinit.f the detector information.
- When the files are generated, copy them into the src directory of GMega: user@earth:~/MEGAsoft/MEGAlib> cp ugeom.f <path to GMega>/src user@earth:~/MEGAsoft/MEGAlib> cp detinit.f <path to GMega>/src
- Switch to the GMega directory and recompile: user@earth:~/MEGAsoft/GMega> gmake

# 12.1.2 Generation of a geometry file for MGeant/MGGPOD

- Switch to the directory where you need the MGeant/MGGPOD files
- Run geomega: user@earth:~/MEGAsoft/MEGAlib> geomega
- Load the geometry file: Menu: File → Open and select the geometry Prototype.geo.setup from the directory <path to MEGAlib>/resource/geometries/simplifiedprototype. Alternatively use the command line switch "geomega -f <file name>" to load the geometry.
- ullet View the geometry:  $Menu: Analysis \to View geometry$ . If you don't see the geometry, check the command line for errors and correct them!
- Generate the MGGPOD/MGeant geometry information with *Menu: Analysis* → *Create MGeant/MGGPOD files*. This generates the files media.med, materials.mat, and setup.geo.
- Now you are ready to run MGeant or MGGPOD

Cosima does not need any special generated geometry file: You only announce the name of the geometry in the \*.source file.

## 12.1.3 Generation of a simulation file

## 12.1.3.1 Generation of a simulation file with GMega

• First make sure that the file Restart.dat does not exist. This file contains restart information in case the last run has been interrupted:

```
user@earth:~/MEGAsoft/GMega> rm -f Restart.dat
```

- Then start GMega: user@earth:~/MEGAsoft/GMega> GMega sgeo/UseCase1.src
- ... and wait until the simulation has finished.

# 12.1.3.2 Generation of a simulation file with MGeant/MGGPOD

- Make sure you compile MGeant/MGGPOD with the ACT2/INIT2 extension (check Imakefile)
- The result of MGeant/MGGPOD is a fits file. This has to be converted into the MEGAlib format via the tool ConvertMGGPOD:
  - user@earth:~/MEGAsoft/GMega> ConvertMGGPOD -f UseCase1.fits -g \$MEGALIB/resource/geometries/simplif
- ... now you have a sim file which can be used with MEGAlib.

## 12.1.3.3 Generation of a simulation file with Cosima

- After you have successfully compiled Cosima, simply launch it with the appropriate source file: user@earth:~/MEGAsoft/Cosima> Cosima source/UseCase1.source
- ... and wait until the simulation has finished.

# 12.1.4 Analyzing the data

# 12.1.4.1 Analyzing the data with Sivan

• Start sivan:

user@earth:~/MEGAsoft/MEGAlib> sivan

- Load the simulation file:  $Menu: File \rightarrow Open$ . Switch to the directory GMega or Cosima and select the simulation file UseCase1.sim or UseCase1.inc1.id1.sim.
- The second step is to load the geometry belonging to the simulation file: Menu: File → Load geometry.
   In the directory <path to GMega>/resource/geometries/simplifiedprototype select the geometry Prototype.geo.setup.

The last step is not necessary, if you have generated your simulation with Cosima, because then the path to the file is included in in the sim file.

In case the geometry is wrong, the analysis will stop with a command similar to: Cannot noise hit ....

• Start the analysis with  $Menu: Analysis \rightarrow Analyze events$  and create \*.tra file This creates a \*.tra file which contains the interaction information.

## 12.1.4.2 Analyzing the data with Revan

• Start revan:

user@earth:~/MEGAsoft/MEGAlib> revan

- The second step is to load the geometry belonging to the simulation file: Menu: File → Load geometry. Switch to the directory <path to MEGAlib>/resource/geometries/simplifiedprototype and select the geometry Prototype.geo.setup.
- Choose all options from *Menu: Reconstruction* → *Selection of algorithm* and **below** which are reasonable. Pay attention that if you use the Bayesian algorithms you need to have generated Response files in advance. Since this is a difficult process, stay with the figure-of-merit algorithms right now.
- Start the analysis with  $Menu: Reconstruction \rightarrow Start event reconstruction$ . This creates a \*.tra file which contains the interaction information.

# 12.1.5 Point source imaging in the near-field

• Start mimrec:

user@earth:~/MEGAsoft/MEGAlib> mimrec

- Load the \*.tra file:  $Menu: File \rightarrow Open$ . Switch to the directory GMega or Cosima and select the simulation file UseCase1.tra or UseCase1.inc1.id1.tra.
- Set the coordinate system to Cartesian 2D in Menu: Backprojection  $\rightarrow$  Coordinate system.
- Set the correct zoom via Menu: Backprojection  $\rightarrow$  Image dimensions. Set the following coordinates:  $x_{min} = -5$ ,  $x_{max} = +5$ ,  $x_{bins} = 50$ ,  $x_{min} = -5$ ,  $x_{max} = 40.9$ ,  $x_{max} = 41.1$ ,  $x_{bins} = 1$ .

- The only reasonable back projection algorithm in Menu: Backprojection  $\rightarrow$  Backprojection algorithm is Areas
- ullet In Menu: Backprojection o Response selection use universal adaptable Compton response
- For the response fit parameters in Menu: Backprojection  $\rightarrow$  Response parameters use Compton longitudinal = 60, Compton transversal = 1.5, Pair = 5.
- In *Menu:* Backprojection → Event selections make sure all event types are allowed except photo effect events (tracked and not tracked Comptons as well as Pair events). The good start point for the energy selection is 0/1200 keV. Make sure that the second energy range is set to on 0/0, that the time walk starts at negative values (e.g. -100), and that the energy range of the scattered gamma-ray starts at negative energies.
- In *Menu:* Backprojection → Memory management make sure that the response accuracy is set to one byte. 4 bytes are needed for elaborate schemes such as coded masks, where the response varies on small scales.
- If you want to look at the spectrum click *Menu: Response* → *Energy spectrum* . Ignore the position coordinates and set a negative radius. (If the radius is positive, then only those events are selected, whose Compton cone is within the radius distance from the given position.)
- If you want to get the image, then in Menu: Likelihood  $\rightarrow$  Algorithm set the number of iterations to 10 and afterwards simply press the "Play" button.

# 12.2 Calculating the effective area of a telescope

The effective area of a telescope is the corresponding area of an ideal telescope which would detect all photons passing through its surface area (and measure their full energy). For Compton telescopes, the effective area is normally significantly smaller than the surface area.

The formula for calculating the effective area is:

$$A_{eff} = A_{start} \frac{N_{Photons\ within\ your\ energy\ and\ ARM\ selection}}{N_{Simulated\ events}} \tag{1}$$

The area from which the photons have been started can be obtained from the simulations. If you have done the simulations with Cosima or GMega, then look into your \*.geo.setup. The start area is a disk whose radius is defined by the first parameter of the "SurroundingSphere". If you have done the simulations with mggpod, then you have to look into your fits file and locate the "BMPAR19" header information, which represents the area you look for. The number of simulated events is the input to your simulations.

To get the number of photons within your energy and ARM selection, follow these steps with mimrec:

- Do all your event selection in the mimrec event selector (Tracks yes or no, Length of Compton sequence, Compton quality factor, Range of Compton scatter angles, Earth horizon distance, etc.), with exception of the energy resolution. For the energy range, you take a window which is significantly larger than the peak width but not to large you want to have the peak width extend over at least 20 bins!
- Look at the energy spectra (the numbers for the GUI before the selection are: 0, 180, -1, 100). Do a fit: Right-click on the top of one of the green bins. You get a menu which is named "TH1D::EnergySpectrum" Select "Fitpanel". Select "gaus". Select with the bottom grey bar the fit window. Click fit. The screen output now tells you the sigma of your Gauss peak. Play a little around to find the best fit.
- Set a new energy window in the event selector (e.g.  $\pm 1.4 \times 1$ -sigma energy resolution)
- Look at the ARM (the numbers for the GUI before the selection are: 0, 180, 5, 100 for on axis sources). The function is automatically fitted by 2 Lorentzians and one (asymmetric) Gaussian. If the fit is not ok, play with your bin size and ARM window ("Radius" parameter in the GUI). Make sure the FWHM of the peak is spread over at least 20 bins! The FWHM determined in the fit is written to the screen.

- Now do the fit again: As radius in the screen, which appears after clicking "ARM of scattered gammaray" set the FWHM of the ARM fit. Generate the ARM and look at the screen output: "Compton and pair events in histogram" are your events in your ARM and energy window.
- Do the math!

# 12.3 Generate a response for Bayesian event reconstruction

The Bayesian Compton and Electron Tracking (BCT & BET) algorithms work reliably and significantly out-perform all previous types of Compton reconstruction algorithms: More photons are reconstructed into the photo peak and background can be better suppressed.

However there is a price to pay - a significant increase in complexity: Firstly one has to generate a large data-space, based on on hundreds of millions of simulated events and secondly the event reconstruction takes almost 10 times longer, due to the large amount of absorption probabilities, which have to be calculated.

# 12.3.1 Generation of the data space

First, one has to generate the data spaces, based on which a quality factor for each possible event combination is calculated via Bayesian statistics. This quality factor corresponds to the probability that the events is good, based on the given measurement parameters.

The data-space has 8 dimensions, which is split into 2 six-dimensional and 1 five dimensional space, to take into account independence of some variables and to simplify its generation process. This process splits into three parts:

- 1. Simulation of a significant amount of photons
- 2. Filling the data spaces
- 3. Potentially merging data spaces of different runs

12.3.1.1 Simulation The simulation should resemble the real measurements as close as possible. However, the response generator can currently only handle photons reliably. For a low-earth orbit satellites, a simulation where 10% of the photons come from the upper hemisphere and 90% from the lower hemisphere should do.

Please consider to make large enough simulations to fill the data-spaces with decent statistics. When the \*.compton.good.rsp response file (BCT) or the \*.central.good.rsp (BET) has an average statistics of 100 counts per bin (details in 12.3.1.3) than you have reached the minimum requirements. Depending on you geometry this might mean to simulate 500 million events and more. Also take into account that simulating the data is the fastest part, filling the data-space and therefore calculating all absorption probabilities takes 20+ times longer!

Remarks for MGGPOD users - for Cosima (Geant4) everything works per default:

- It is necessary to make the simulation with the INIT2 and ACT2 keyword
- Since you are probably running dozens of simulations in parallel, make sure to always use a different set of random numbers!
- Set the electron cut off in the input dat file to  $10^{-7}$ .
- You have to convert the fits file into the sim file with the special option "-s 21" of ConvertMGGPOD. This retrieves significantly more information from the fits file, so that the sim-file will definitely get larger than the fits-file!

**12.3.1.2** Filling the data space The filling of the data-space is done with the program ResponseGenerator. Its mandatory options are:

- -g: geometry file name
- -f: sim file name
- -r: output response file name
- -m: modes: t = track or c = Compton
- -c: revan configuration file

It is very important to specify the options "-m c" (or "-m t" for a tracking response) and the revan configuration file. The revan configuration file contains all options, with which the events are reconstructed. All relevant options can be set via the revan GUI. Simple press the menu option "Save configuration" after you have configured revan

The output of ResponseGenerator are several \*.rsp files with represent several aspects of the response.

12.3.1.3 Merging response files You probably have done dozens of simulations in parallel, generated hundreds of GB of simulation data and now want to merge all sets of \*.rsp files into one set. This is done via the program ResponseManipulator:

user@earth:~/MEGAsoft/MEGAlib> ResponseManipulator -f New.blabla.rsp -a A.rsp -a B.rsp This has to be done for all different response files.

If the call "ResponseManipulator -s BaseName.compton.good.rsp" tells you that you have a statistics of at least 100 cts/bin, then you are done!

# 12.3.2 Event reconstruction

Now load the revan configuration file  $Menu: File \to Load\ configuration$ , switch to the reconstruction  $Menu: Reconstruction \to Selection\ of\ algorithm\ and\ load\ the\ *.rsp\ files\ Menu:\ Reconstruction \to Sequencing\ options$ . Here, you only need to give the \*.goodbad.rsp file. All other files are expected to have the same base name and to reside in the same directory as the main file.

During the event reconstruction the most likely event sequence is determined as well as an overall quality factor of the event. The quality factor represents the probability that the event is good (correctly reconstructed and completely absorbed) based on the given measurements.

This quality factor (CQF) is the main event selection criteria for mimrec. A CQF of 0.1 means that the event is good based on the measurements with a probability of 90%.

# 12.3.3 Additional warnings

- Whenever you change your detector, you need to redo the whole analysis!
- Always do a test with a small file first, before you start to generate 100+ GB of data!

# 12.4 Generate a response for the earth horizon cut

Use the program ResponseGenerator on a sufficiently large source data set. The simulation has to be done with full simulation information (ACT2 and INIT2 in case you used mgeant/mggpod).

The call is:

 ${\tt user@earth:} {\sim}/{\tt MEGAsoft/MEGAlib>} \ {\tt ResponseGenerator-g~Bla.geo.setup-f~Bla.sim-r~Bla-m~e-c} \\ {\tt Revan.cfg-b~Mimrec.cfg-s~10000}$ 

The mimrec configuration file should be reasonably similar to what is used for the sensitivity calculations - of course with exception of the earth horizon cut.

When a call to ResponseManipulator, tells you that the average bin content is well above 100, then you have enough statistics:

 $user@earth: \sim / MEGAsoft/MEGAlib> \\ ResponseManipulator -s \ Bla.compton.ehc.rsp$ 

You can view the content of the Response Matrix via:

user@earth:~/MEGAsoft/MEGAlib> ResponseManipulator -v Bla.compton.ehc.rsp x y 0 0 0 0