SHIELD-HIT12A - User’s Guide

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  [http://www.inr.ru/shield](http://www.inr.ru/shield)
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## Contents

1 Introduction .................................................. 1
  1.1 SHIELD-HIT12A ........................................... 1
    1.1.1 Particle transportation ............................ 2
    1.1.2 Nuclear physics .................................... 2
    1.1.3 Atomic physics .................................... 3
    1.1.4 Scoring of physical quantities .................... 3
    1.1.5 Other features .................................... 4

2 Quick start guide ............................................ 5
  2.1 Linux: Installation from binary distributions .......... 5
  2.2 Windows ................................................ 5
  2.3 Converter script :convertmc : .......................... 5
  2.4 Setup a SHIELD-HIT12A run ............................ 6
  2.5 Example ................................................ 6
  2.6 Analyzing output ....................................... 9

3 Input files .................................................. 11
  3.1 Overview ................................................. 11
  3.2 mat.dat - Target medium ............................... 12
    3.2.1 Loading external stopping power files ............ 12
    3.2.2 Voxelized structures ................................ 13
    3.2.3 List of descriptors for mat.dat .................. 14
    3.2.4 Examples - Material initialization ............... 15
  3.3 beam.dat - Projectile, statistics, seeds ............. 17
    3.3.1 List of descriptors for beam.dat ................ 18
    3.3.2 USECBEAM - External source file ................. 21
    3.3.3 USEPARLEV - External parameter file ............. 22
    3.3.4 USEBMOD - Beam modulator ....................... 23
    3.3.5 Examples - Beam initialization .................. 26
  3.4 geo.dat - Geometry .................................... 27
    3.4.1 Title .............................................. 27
    3.4.2 Bodies ............................................ 27
    3.4.3 Voxelized Bodies ................................... 28
    3.4.4 Zones ............................................ 29
    3.4.5 Media ............................................ 30
    3.4.6 Examples - Geometry ................................ 30
  3.5 detect.dat - Scoring ................................... 34
    3.5.1 GEOMAP - Mapping the geometry .................. 34
    3.5.2 ZONE - Scoring by zone ........................... 35
<table>
<thead>
<tr>
<th>Table</th>
<th>Title</th>
<th>Page</th>
</tr>
</thead>
<tbody>
<tr>
<td>A.1</td>
<td>Units</td>
<td>58</td>
</tr>
<tr>
<td>A.2</td>
<td>Particles</td>
<td>59</td>
</tr>
<tr>
<td>A.3</td>
<td>Nuclear target identifiers</td>
<td>60</td>
</tr>
<tr>
<td>A.4</td>
<td>ICRU materials</td>
<td>62</td>
</tr>
<tr>
<td>A.5</td>
<td>Input and output files</td>
<td>69</td>
</tr>
<tr>
<td>A.6</td>
<td>PARLEV parameters</td>
<td>72</td>
</tr>
</tbody>
</table>
Chapter 1

Introduction

The Monte Carlo particle transport code SHIELD-HIT\(^1\) is designed to precisely simulate therapeutic beams of protons and ions in biological tissue relevant for ion beam cancer therapy. SHIELD-HIT (Heavy Ion Therapy) evolved from the common SHIELD code that models interactions of hadrons and atomic nuclei in complex extended targets in the energy range up to 1 TeV/nucleon.

SHIELD and SHIELD-HIT employ the same models to simulate nuclear interactions, which were developed at JINR, Dubna\(^2\) and INR RAS (Moscow)\(^3\). The models are grouped together in the MSDM-generator (Multi Stage Dynamical Model) which allows to simulate a whole nuclear reaction in the exclusive approach. Contrary to the inclusive approach, every generated secondary particle is tracked and processed until it stops, decays, leaves the simulation universe or interacts destructively. Parameters of all secondaries including residual nuclei are retained, thereby fulfilling all conservation laws. Neutron transport below 14.5 MeV is simulated using the 28 group neutron data system ABBN\(^1\) both in SHIELD and SHIELD-HIT.

SHIELD calculates ionization losses of charged hadrons and nuclear fragments according to the Bethe-Bloch equation. The heavy ion version SHIELD-HI contains also ATIMA stopping powers\(^2\). The "medical" version SHIELD-HIT includes various models and data sets to compute mean ionization loss, fluctuation of ionization losses and multiple Coulomb scattering. In addition, external stopping power tables (including MSTAR and ICRU tables) can be provided using the libdEdx\(^3\) stopping power library.

A more elaborate history of the various releases of SHIELD-HIT is given in section B.2.

1.1 SHIELD-HIT12A

SHIELD-HIT12A was forked from SHIELD-HIT08 and its source code includes massive changes. The physics engine was tuned to new experimental nucleus-nucleus interaction data, many performance improvements were made and generally the entire code base was restructured, now eliminating the need for users to recompile code for typical usage applications. Several bug fixes from the original SHIELD-HIT branch are forwarded and included in SHIELD-HIT12A. Further development of the Physics engine and bug fixes from the original SHIELD-HIT branch enter continually in SHIELD-HIT12A and vice verse.

Characteristic features of SHIELD-HIT12A can be summarized in the following way:

---

\(^{1}\)Official homepage: http://www.inr.ru/shield

\(^{2}\)http://www.jinr.ru/

\(^{3}\)http://www.inr.troitsk.ru/
1.1.1 Particle transportation

- Transport of neutrons, pions, kaons, atomic nuclei, and a number of anti particles in energy range up to 1 TeV/nucleon for SHIELD-HIT. SHIELD-HIT12A is limited to 2 GeV/A. Lower energy cutoff is $E_{\text{cut}} = 25$ keV/nucleon except for atomic nuclei with $Z > 20$ in SHIELD-HIT12A where $E_{\text{cut}} = 1$ MeV/nucleon.

- Available initial beam configurations: Gaussian, flat square or flat circular in any direction. Beam divergence can be Gaussian or flat beam; focused or defocused beams can be specified.

- Geometric configuration of the target as an arbitrary combination of geometric bodies bounded by the second order surfaces (Combinatorial Geometry compatible) [4, 5].

- Arbitrary chemical and isotopic composition of materials in target zones can be defined based on the table of available isotopes shown in table A.3.

- Full memorization of the extra-nuclear cascade tree during simulation without any loss of physics information.

- Formation of neutrons ($E_n < 14.5$ MeV) as well as electrons/positrons and $\gamma$-rays, which were created during simulation of extra-nuclear cascade and $\pi^0$ decay. However, only neutrons are transported.

- The possibility to switch on/off various physics processes (energy straggling, multiple scattering, nuclear interactions) on user request.

1.1.2 Nuclear physics

- Simulation of inelastic hadron-nucleus and nucleus-nucleus interaction in exclusive approach using a Multi Stage Dynamical Model (MSDM-generator) [6]. The total and inelastic cross sections of the hadron-nucleus and nucleus-nucleus interaction are calculated according to [7–9] with modifications by [10–12]. These cross sections are used for sampling of the nuclear interaction path length as well as for choice of the interaction type (inelastic/elastic). The MSDM-generator describes all stages of the nuclear reaction in the exclusive approach. Current versions of known Russian nuclear models are interfaced in the MSDM-generator:
  - Fast, cascade stage of the nuclear reaction
    * Intranuclear cascades is handled by the Dubna Cascade Model (DCM) [13]
    * Independent quark-gluon string model (QGSM) [14][16]
    * The coalescence model [13]
  - Precompound emission of nucleons and lightest nuclei [17]
  - Equilibrium de-excitation of the residual nucleus
    * Fermi break-up of light nuclei [18]
    * Evaporation/Fission competition [18][19]
    * Multifragmentation of highly excited nuclei (SMM) [20]

- Neutron transport ($E_n < 14.5$ MeV) on the basis of the 28-group neutron data ABBN [1]. A phase space file of these neutrons can be exported for use in other programs such as MCNP or MCNPX [21].
1.1. SHIELD-HIT12A

- Two- and three-particle modes of decay of pions and kaons.
- Neutron transport below 14.5 MeV is simulated by the original neutron transport code LOENT (Low Energy Neutron Transport) [22] using the 28 group neutron data system ABBN [1]. The LOENT code may be used both separately and as a part of the SHIELD code, since SHIELD and LOENT use the same geometry parser (Combinatorial Geometry) [4, 5].

The LOENT code uses the following tables from the neutron data system ABBN:

- st - total cross section
- sf - fission cross section (n,f)
- n - mean number of fission neutrons
- sc - capture cross section (n,c)
- sin - inelastic scattering cross section (n,n’), including the reaction (n,2n)
- se - elastic scattering cross section (n,n)
- m - mean cosine of the angle of the elastic scattering
- sin(g,g+k) - matrix of inter group transitions at the inelastic scattering.

The LOENT code gets neutrons from an external neutron source and follows them, one by one, until the end of the neutron trajectory. Multiplication of neutrons is possible via the reactions (n,2n) and (n,f). Each neutron has its statistical weight attached and a cumulative timer, which accumulates the time from the beginning of the neutron transport history. After transition of the neutron to the thermal group, its energy does not change in further collisions.

1.1.3 Atomic physics

- Ionization losses of charged hadrons and nuclear fragments according to the Bethe-Bloch equation and the Lindhard Scharff equation at low energies.
- Multiple Coulomb scattering simulated with Moliere’s or a Gaussian model, similar to that implemented in Geant [23].
- Fluctuations of the ionization energy loss (energy straggling) simulated by a fast implementation (article in preparation) of Vavilov’s model [23, 24] or Gaussian distribution.
- Possibility to load external mass stopping power tables to override internal ones. A script interfacing to libdEdx [3] is available.

1.1.4 Scoring of physical quantities

A generic scoring system was developed—that is specified by the user—which provides:

- Detectors for a wide range of physical quantities, which can be made sensitive to specific particles or particle groups.
- Arbitrary scoring in Cartesian or cylindrical scoring grids.
- Alternatively, detectors can be assigned to zones that constitute the target geometry.
- Possibility to generate spectra and depth-dose files in TRiP98 format (.spc and .ddd).
● Routines to merge results from multiple runs from parallel processing system.

In addition to the generic scoring system, the legacy scoring system is retained, providing:

● Scoring of the production rates of radioisotopes in the entire system.

● Track Length Estimation (TLE) of differential (differential in energy) fluences and energy fluence of secondary particles and nuclear fragments averaged over each geometric zone of the target.

● Scoring of contributions to the energy deposition from various types and from different generations of particles and nuclear fragments separately.

However, the post-processing scripts available for this scoring system cannot merge the ASCII output from parallel runs.

1.1.5 Other features

● Reading and processing of beam source and ripple filter files (see sections 3.3.2 and 3.3.4, respectively).

● Support for parallelization (see chapter 6).
Chapter 2

Quick start guide

2.1 Linux: Installation from binary distributions

SHIELD-HIT12A should come in a tarball of the form: shield_hit12a_vXXX.tar.gz, where XXX refers to the release version. Untar the file:

$ tar xvfz shield_hit12a_vXXX.tar.gz
$ cd shield_hit12a_vXXX
$ sudo make install

which will copy the binaries to /usr/local/bin. This should work on Ubuntu type systems. If you wish to install to other directories, you may have to edit the Makefile accordingly. On non-sudo type systems (such as Debian or RedHat based Linux systems) one can become root with su instead of sudo and the last line then reads:

$ su
# make install

2.2 Windows

SHIELD-HIT12A was tested on Windows 7 and Windows XP systems, but may in principle also run on other Windows systems. SHIELD-HIT12A is provided as a zip file for either 32 or 64 bit systems. Once you unzipped it, you will find a ready-to-run shieldhit.exe file.

2.3 Converter script convertmc

The output from SHIELD-HIT12A is in a binary format, which is not human readable. In order to convert the output to something sensible, such as ASCII data, or perhaps a plot, you must use the convertmc script, which is provided by the pymchelper package you can find on [https://github.com/DataMedSci/pymchelper](https://github.com/DataMedSci/pymchelper). On Linux systems, make sure you have the python package manager pip installed (Ubuntu and debian: apt-get install pip Fedora/RedHat: dnf install pip or yum install pip). Then simply run

$ pip install pymchelper

to install it as a user. For system wide installation you can also run the same command as root user. If you have already installed pymchelper you may want to upgrade it:
2.4 Setup a SHIELD-HIT12A run

You can run one of the examples found in base directory, simply by entering:

```bash
$ shieldhit examples\simple
```

or change into the directory and run `shieldhit` without arguments.

Windows users can open a shell (Run - “Enter a command” - cmd) and change to the directory of where the `shieldhit.exe` file is found. To run the example, simply type:

```bash
$ shieldhit examples\simple
```

Or simply drop the `shieldhit.exe` file into the folder you want to run, and double click on `shieldhit.exe`.

2.5 Example

Here is a quick description of the most important files in the example `examples/simple` supplied with the distribution:

**geo.dat**

The file `geo.dat` describes the simulated geometry. Here we want to simulate a target which is a cylinder where the center of its base is located in origin (0,0,0) cm. The height of the cylinder is described by a vector along the Z axis, and is here set to 30 cm. Finally the cylinder radius is set to 10 cm. The cylinder is surrounded by a universe consisting of vacuum which does not interact with the particle. Finally that vacuum is surrounded by a black hole medium. Any particle hitting this medium will be disintegrated, and not tracked any further.

All this information is formatted in “cards”. Cards are lines which are no longer than 80 characters long—a legacy from last centuries punch card systems. Only ASCII characters are allowed; as a white character only space is allowed while tabs are not.

Each card has a number of arguments, which are data being passed to the code. The position and lengths of the arguments varies, thus one should make sure that the exact positions are used.

Lines beginning with * are ignored, and can be used for comments. So, in the case of the simulation described earlier, our file looks like this:

```plaintext
*---><---><------><------------------------------------------------>
 0 0 C12 200 MeV/A, H2O 30 cm cylinder, r=10, 1 zone
*---><---><------><------><-----><------><------><------><------><------>
RCC 1 0.0 0.0 0.0 0.0 0.0 30.0
  10.0
RCC 2 0.0 0.0 -5.0 0.0 0.0 35.0
  15.0
RCC 3 0.0 0.0 -10.0 0.0 0.0 40.0
  20.0
END
*.<---><--->..<--->OR<--->OR<--->OR<--->OR<--->OR<--->OR<--->OR<--->
```
The first line, is a comment line, which is ignored. It is inserted here, to better see where the fields are which are interpreted by SHIELD-HIT12A. The following line specifies a title of the geometry setup. The title is preceded by two integers which are described later in section 3.4. After another comment line, the next two lines describe the geometry of the aforementioned target cylinder. The \texttt{RCC} card selects a body (here a Right Circular Cylinder). The body is assigned to a number 1. The \(x,y,z\) coordinates of the base follow, along with a vector which spans to the top of the cylinder. Since the cylinder is following the \(Z\) axis, all entries are zero, except for the height of the cylinder, i.e. the \(z\)-component of the vector. The following card is a continuation card of the \texttt{RCC} card, and holds only one value, namely the radius of the cylinder. All values are in \(\text{cm}\). A list of default units in SHIELD-HIT12A is provided in section A.2. Afterwards two more cylinders follow, each expanded by an additional 10 cm in all three directions.

The geometry section, which only contains one body here, must be terminated by the \texttt{END} card.

The lines

\begin{verbatim}
  *..<-><--->..<--->OR<--->OR<--->OR<--->OR<--->OR<--->OR<--->OR<--->OR<--->OR<--->OR<--->
  001 +1
  002 +2 -1
  003 +3 -2
\end{verbatim}

assigns the bodies we just specified to zones. For the first body (our target cylinder) the body and the zone are identical. The surrounding vacuum is represented by the second cylinder, but subtracted the inner cylinder, since this is already described by zone 001. Similarly zone 003 consists of the largest body (number 3), subtracted the body number 2. There is no reason to subtract body number 1 here, as it is already entirely contained in body number 2. The point is that a zone can be any Boolean composition of previously defined bodies, which is described in detail later in section 3.4. Any point inside the black hole must be assigned to a zone, and only one zone.

The section describing the zones must also be terminated with the \texttt{END} card.

Finally, two lines follow: the first specifies a list of zones, and the second assigns a material number to that zone. In this example, each line consists only of three numbers since we have three zones. In the second line we tell which materials each zone consists of. The numbers 0 and 1000 are predefined, any other number will be specified in the \texttt{mat.dat} file. 0 is black hole, 1000 is vacuum and the inner most cylinder is made of material with the id\# 1.

\textit{mat.dat}

The file \texttt{mat.dat} specifies the medium of the zone we just specified in \texttt{geo.dat}.

We define medium number 1 to be water. The most simple way to do this is by writing:

\begin{verbatim}
  MEDIUM 1
  ICRU 276
  END
\end{verbatim}
The first line tells SHIELD-HIT12A that we now are going to assign MEDIUM number one. The second line says that we want to use a default ICRU material with material number 276. The numbers of predefined materials (as given by ICRU) are listed in table A.4 in section A.5 with 276 being water. Each MEDIUM section must be terminated with an END line for SHIELD-HIT12A to initialize the material. The specification materials, other than those listed in table A.4 will be explained in section 3.2.

beam.dat

For beam.dat we should initially only worry about a few lines. Fairly at the top, JPART0 specifies the number of the primary particle according to table A.2.

<table>
<thead>
<tr>
<th>JPART0</th>
<th>25</th>
<th>! Incident particle type</th>
</tr>
</thead>
<tbody>
<tr>
<td>HIPROJ</td>
<td>12.0</td>
<td>6.0</td>
</tr>
</tbody>
</table>

Here we request a beam of heavy ions (JPART0 = 25), which is specified further with the HIPROJ card. By stating $A = 12$ and $Z = 6$ we select carbon-12 ions to be our primary particle.

The energy is specified with the TMAX0 card, here set to 391.0 MeV/nucleon.

| TMAX0 | 391.0 | 0.0 | ! Incident energy; (MeV/nucl) |

The statistics, i.e. the amount of primary particles to be simulated is specified in the NSTAT line:

| NSTAT | 2000 | 1000 | ! NSTAT, Step of saving |

With no further input we get a default pencil beam with $\sigma_x = 0.0$ $\sigma_y = 0.0$ cm which starts at origin $(0,0,0)$.

detect.dat

Finally, we can specify detectors to score a variety of quantities. The file detect.dat is optional, but very useful. In this example, we want to score a depth dose-curve along the water target.

We can setup a cylinder, following the Z axis, located inside the water target. It does not need to match any geometry specified in geo.dat. Note, this file requires fixed format.

| CYL | 0.0 | 0.0 | 0.0 | 10.0 | 7.0 | 30.0 | 1 | 1 | 300 | -1 | ENERGY | ex_cyl |

The first card specifies cylindrical scoring, staring from point $(0,0,0)$, and with the radius 10 cm, covering $2\pi$ (by specifying $7\frac{1}{3}$) of the circumference of the cylinder, and with a height of 30 cm along the Z axis. Currently, the CYL scoring cannot be rotated, and is always placed parallel to the X-axis. In the next line the first three arguments say that we want 1 bin for the radius, 1 bin for the angle spanning the cylinder and 300 bins along the height. The result will then be a one-dimensional file, where the scored quantity is listed in steps of 30.0 cm / 300 bins (i.e. 1 mm). The next three arguments tell that we want the ENERGY to be scored of the particle $-1$ which means the energy deposited by all particles crossing the bins. ex_cyl is the name of the output file.

---

1 Any number larger than $2\pi$ means we do not subdivide the cylinder along its radial lines, i.e. we do not cut the cylinder in pieces of pie.

2 I.e. scoring along the entire circumference and not into pie pieces.
2.6. Analyzing output

Alternatively we could also specify a cartesian scoring mesh. The two next lines specifies a mesh with one bin from -5 to 5 cm along the X and Y axis, and then 300 bins from 0 cm to 30 cm along Z. The result is saved in the file ex_zmsh.

```
MSH -5.0 -5.0 0.0 5.0 5.0 30.0
  1   1   300  -1 ENERGY ex_zmsh
```

We can here also score along another axis if we want. The next example scores 100 bins from -5 cm to 5 cm along the Y axis.

```
MSH -1.0 -5.0 10.0 1.0 5.0 12.0
  1  100   1  -1 ENERGY ex_ymsh
```

or if we want to see a 100x300 pixel 2-D map of the energy deposited along the beam path in the target:

```
MSH -5.0 -5.0 0.0 5.0 5.0 30.0
  1  100  300  -1 ENERGY ex_yzmsh
```

2.6 Analyzing output

After SHIELD-HIT12A terminated the run you find several new files in your directory. Here the results from the four detectors, defined in detect.dat, are stored in the files ex_zmsg.bdo, ex_ymsh.bdo, ex_yzmsh.bdo and ex_cyl.bdo. These files are in binary form and the script convertmc can be used to convert them to ASCII format. Stepping into the directory and applying,

```
$ convertmc txt ex_zmsh.bdo
```

will produce a human-readable output file ex_zmsh.txt, which can be plotted. You can also directly produce a plot in .png format

```
$ convertmc image ex_zmsh.bdo
```

convertmc is also capable generating plots directly, please see the documentation of pymchelper at

https://pymchelper.readthedocs.io/

Uncertainties are briefly covered in section 3.5.11.
Chapter 3

Input files

3.1 Overview

The SHIELD-HIT12A transport code works with at least three input files.

*mat.dat* – chemical composition of materials in target zones.

*beam.dat* – several parameters, explained below (like seed, projectile, statistics, etc.).

*geo.dat* – geometry of the target using combinational geometry (CG), similar to FLUKA.

Optionally, the user can include an additional file:

*detect.dat* – for simple scoring of physical quantities in independent geometries.

These four files are described in more detail in the sections to follow.

As extension of the file *beam.dat* up to three optional ASCII formatted files (with user-specified names) can be supplied that specify: a ripple filter, a beam source file, and parameters for nuclear interaction models. Examples for these three files are shown adjacent to the description of file *beam.dat* in Section 3.3. If chosen by the user, SHIELD-HIT12A may also look for external ASCII files containing stopping power data as described in Section 3.2.

All input files must be grouped in the same folder. Its location can be specified by providing a command line argument to the SHIELD-HIT12A executable. All output files are placed into the same directory. The SHIELD-HIT12A executable itself can be run from anywhere if the exact path to the directory with the input files is given. If the executable is run from the directory containing the input files the specification of the path can be omitted.
3.2 **mat.dat - Target medium**

The `mat.dat` file defines the materials that are used in the simulation. It consists of cards (lines starting with a descriptor followed by possible arguments) and comment lines. Any line starting with a * or a < or a blank line are regarded as a comment, which is ignored and may appear at any place. The ordering of the cards has to obey certain rules. Some cards can be omitted, and SHIELD-HIT12A will assign the default values given in the descriptor list below. It is recommended to specify all parameters explicitly since default values may change in future releases of SHIELD-HIT12A. While the ordering of the arguments must be as specified for the respective card, they can be positioned in any column (free format).

Each parseable line starts with a descriptor, which specifies what parameters are to be described. After this descriptor, none, one or two arguments follow, depending on the type of the descriptor. The descriptor and each argument must be separated with at least one space or tab. The rest of the line is not parsed. It can be used for comments if it is separated from the last argument with at least a space or tab. Only the first 128 characters are read; any characters beyond will be ignored.

Each material specification starts with the MEDIUM card. A medium can consist of a single element or of a compound. A compound again can be a chemical compound, or simply a mix of elements and/or isotopes, e.g.:

- Pure medium - element with one fixed Z (e.g. O, Fe, Cu, etc.).
- Chemical composition, e.g. water H$_2$O.
- Isotope mixture, if isotopes of the same Z have different neutron properties at low energies, e.g. mixture of $^{235}$U and $^{238}$U.
- a chemical compound containing isotopes, e.g. $^6$LiF.

The maximal total number of different media NUMMED is limited to 100, excluding internal and outer vacuum which are predefined as medium numbers 1000 and 0, respectively. Furthermore, if CT-images are imported using the VOX card, the available medium slots are reduced depending on the Hounsfield unit to medium segmentation algorithm applied. This may be up to 40 media, leaving 60 media for the user.

The user can select one of the predefined media listed in the ICRU material table in section A.5. Alternatively, the user can define a MEDIUM by specifying the chemical elements or isotopes, which can be found in this medium. The constituent elements are limited to a total number of 13. A chemical element is described by the variable NUCLID. Usually, NUCLID=Z selects the element from the periodic table corresponding to the atomic number Z (1 < Z < 100). Further isotopes, which differ in the number of neutrons, are predefined with values of NUCLID > 100 and are listed in table A.3. If an isotope does not appear in the list, no low energy neutron cross section data are included in SHIELD-HIT12A. In this case SHIELD-HIT12A will not start the simulation, if the neutron transport cutoff energy (NEUTRLCUT in beam.dat see section 3.3) is set below 14.5 MeV.

Note that the corresponding neutron cross sections are either for natural isotope mixtures or for a specific isotope.

### 3.2.1 Loading external stopping power files

The user can provide external files containing mass stopping power data by specifying the LOADEDX card with an appropriate value from the material list in table A.5. Each file contains data for the lightest 18 ions from $^1$H$^+$ (proton) to $^{40}$Ar$^{18+}$ interacting with one specific target material.
These external files are expected to be ASCII formatted, space or tab separated columns, and should follow the ICRU 73 [25] energy grid (53 energy nodes from 25 keV/u up to 1 GeV/u). SHIELD-HIT12A expects the grid to contain the first 18 ions, i.e. from protons ($Z=1$) to Argon ions ($Z=18$). The mass stopping power data must be in units of MeV cm$^2$/g. Comment lines starting with * are allowed anywhere in the file, but no line (either comment or data) may be longer than 512 characters. External stopping power files may contain data different from ICRU, e.g. ATIMA [2] data, but the energy grid must be compatible to ICRU 73. The libdEdx [3,26] computer library can serve as a source for stopping power data. A script shield.dEdx, described in section 5.2.4, can generate these tables for most ICRU materials using libdEdx.

Examples of external stopping power files are: Water.txt, Air.txt, A-150.txt, Kapton.txt, which are supplied along with the SHIELD-HIT12A distribution. In table A.5 the naming convention of the external stopping power files is listed.

### 3.2.2 Voxelized structures

A CT cube loaded with the VOX card in geo.dat must be assigned to a medium number, which is specified as a special voxel type medium in mat.dat. This is done using the VOXMED card in mat.dat. Specifying this card, will prepare a list 24 materials, based on a paper by Schneider et al. [27], which is valid for Hounsfield units between -1000 to +1600. Additionally, a density scale is created, and assigned to each voxel, depending on the Hounsfield units.

$$\rho(HU) = \begin{cases} 
1.03091 + 1.0297 \cdot 0.001HU & : HU < -98 \\
1.0018 + 0.893 \cdot 0.001HU & : -98 \leq HU < 14 \\
1.03 & : 14 \leq HU < 23 \\
1.0003 + 1.169 \cdot 0.001HU & : 23 \leq HU < 100 \\
1.017 + 0.592 \cdot 0.001HU & : 100 \leq HU 
\end{cases}$$

with $\rho$ in units of [g/cm$^3$]. Materials with densities below 0.5 will automatically have the STATE set to 1 (gaseous), and 2 (condensed) otherwise.

User-specified Hounsfield conversion tables may be implemented upon request to the developers.
3.2.3 List of descriptors for *mat.dat*

The recognized descriptors are listed alphabetically, and not by relevance.

**AMASS**: 1 argument. Optional card which can override the atomic mass \( A \) of an element when specified after a **NUCLID** card.

**END**: No argument. Mandatory card that terminates the description of a medium that was started by the **MEDIUM** card.

**ICRU**: 1 argument. Optional card which will select a material from the ICRU list in section A.5. This card may not be used together with the **NUCLID** card for a **MEDIUM**, as it may lead to unpredictable behaviour.

**IVALUE**: 1 argument. Optional card that can override the default mean excitation energy \((I\text{-value})\) of the element that was specified with the preceding **NUCLID** card. Units are eV.

**LOADDEDX**: 1 optional argument. Optional card which will trigger SHIELD-HIT12A to look for an external stopping power file following the naming scheme stated in the table in section A.5. The argument must be the ICRU number of the material to be loaded. However, if used after the **ICRU** card, the ICRU number is already known, and the argument can be omitted. The external stopping power files should contain mass stopping power data for the medium for ions ranging from \(^1\text{H}^+\) (proton) to \(^{40}\text{Ar}^{19+}\). See section 3.2.1 for a description of the format.

**MEDIUM**: 1 argument which specifies the medium number. The numbering must be sequential, starting at 1 for the first medium which is specified.

**NUCLID**: 2 arguments. This card adds a single element to the medium. First argument is the **NUCLID** number specifying the element or isotope from table A.3. You can also use a \( Z \) value that is missing in this list but there are no neutron cross section data attached to it. Second argument is the relative stoichiometric fraction, which will automatically be normalized by SHIELD-HIT12A. Up to 13 elements can be defined per **MEDIUM**. SHIELD-HIT12A will refuse to start, if there are no neutron cross sections for the requested element, and the lower neutron energy cut is set below 14.5 MeV.

**RHO**: 1 argument which specifies the density of the medium in g/cm\(^3\). This card is mandatory in conjunction with the **NUCLID** card. It should be stated after the **MEDIUM** card and before the first **NUCLID** card. In the case that the material is setup using the **ICRU** card, **RHO** will override the default value.

**STATE**: 1 arguments which specifies the state of the medium. Appropriate mean excitation energies \((I\text{-values})\) are selected for each element in a compound, following the ICRU49 guidelines [28]. This card is mandatory in conjunction with the use of a **NUCLID** card. It should be stated after the **MEDIUM** card and before the first **NUCLID** card.

SHIELD-HIT12A discriminates between a chemical compound and just a mix of elements. If **STATE 0**, then SHIELD-HIT12A uses \( I \)-values for elements. If **STATE 1** or **STATE 2** SHIELD-HIT12A uses \( I \)-values for compounds in gas or liquid/condensed phase, respectively. In the case that the material is setup using the **ICRU** card, **STATE** will override the default state of the ICRU material.

**VOXMED**: Specifies that this medium will be a voxel structure, prepares a material database consisting of 24 materials following Schneider et al. [27]. Density will be scaled to Hounsfield units, as described in section 3.2.2.
3.2.4 Examples - Material initialization

Example 1: Specifying five materials

An example of \textit{mat.dat} with five materials is given below

1. Water, ICRU default setup. External stopping powers will be loaded from \textit{Water.txt}.

2. Air, ICRU default setup.

3. PMMA (also known as Perspex or Lucite), is defined. It is a compound in solid state. In this example we assume slightly different parameters, i.e. we override the default density with a density of 1.20 g/cm$^3$. Also the default I-value of hydrogen is changed to 21.9 eV. (This example is merely intended for illustration, and is not motivated to mimic a real situation.)

4. Alanine, ICRU default setup, but overriding the density.

5. Lithium-6 Fluoride (LiF), which is composed of Lithium-6 isotopes which has a very high thermal neutron cross section. External stopping power tables are loaded from \textit{LiF.txt}.

* ----------------- WATER -----------------
MEDIUM 1
ICRU 276
LOADDEDX
END

* ---------------- AIR -------------------
MEDIUM 2
ICRU 104
END

* ---------------- PMMA ------------------
MEDIUM 3
STATE 2
RHO 1.20
NUCLID 1 8
IVALUE 21.9 ! Override I-value of hydrogen
NUCLID 6 5
NUCLID 8 2
END

* --------------- NPL ALANINE ------------
MEDIUM 4
ICRU 105
RHO 1.23 ! Override density, since NPL alanine is less dense
END

* ------------- Li6-Fluoride -------------
MEDIUM 5
STATE 2
RHO 2.635
NUCLID 105 1
NUCLID 9 1
LOADDEDX 185 ! Better stopping powers from external lib.
END
Example 2: Voxelized CT structure

Another example, where a voxelized structure is assigned to medium number 2:

```
* ------------------ WATER ------------------
MEDIUM 1
ICRU 276
END
* --------- VOXEL STRUCTURE---------
MEDIUM 2
VOXMED
END
```
3.3  *beam.dat* - Projectile, statistics, seeds...

The *beam.dat* file characterizes the particle beam transport by defining: the primary beam properties; the required statistics and random seed; details of the physics engine; and additional options.

The format is free, meaning that the arguments can be given in any columns, but they must be ordered as in the description of the respective card. After the last argument, any text string for comments are allowed. Cards can be arbitrarily ordered. Any card (with exception of HIPROJ, see below) can be omitted, and SHIELD-HIT12A will then use default values. The default values are given in the descriptor list below, but may change unnoticed in future releases of SHIELD-HIT12A. Therefore it is recommended to avoid using default parameters, if they are important to the simulation.

In *beam.dat* any line starting with a * or a < is regarded as a comment and is ignored. Blank lines will be ignored too.

Each parsable line starts with a descriptor that specifies what parameters are to be described. After this descriptor, one to three arguments follow, depending on the type of the descriptor. The descriptor and each argument must be separated with at least one space or tab. The rest of the line is not parsed, and can be used for comments. However it is important that the comments are separated too from the last argument with a space or tab. Only the first 128 characters are read, any characters beyond will be ignored.
3.3.1 List of descriptors for beam.dat

The recognized descriptors are listed alphabetically, and not by relevance.

APCORR : Optional card. 1 argument that switches the antiparticle physics correction on and off. This will affect the annihilation physics of antiprotons, $\bar{p}$ and $\bar{K}^-$. The argument can be either 0 or 1. 0 for using the default settings – calculating the capture probability based on the Fermi-Teller $Z$-law [29], and using the default antiproton cross sections. 1 for scaling the antiproton cross section by a factor of 1.08 and using a more physical correct theory for the capture probability. See section B.3.

If this card is not specified, APCORR is set to zero.

BEAMDIR : Optional card. 2 arguments that specify the beam direction: the polar and azimuth angles $\theta$ and $\phi$, both in degrees, as explained in figure 3.1.

Initially, the beam divergence and shape is setup assuming transport along Z axis. No matter where the card is specified, the actual rotation of the beam into the requested beam direction will always take place as a last step, before transportation begins. If this card is not specified, then beam transport along the Z axis is assumed.

If this card is not specified, $\theta$ and $\phi$ are set to zero. That is, by default the beam points in Z direction.

BEAMDIV : Optional card. 3 arguments that specify the divergence and focal point of the beam. First two arguments define the divergence in Gaussian sigmas in X and Y direction assuming beam transport in Z direction. (A subsequent rotation of the beam can be specified with the BEAMDIR card.) The beam divergence must be in mrad. Negative values will generate a flat distribution instead. The distributions are weighted along the polar angle, so requesting a flat angle distribution will also result in a flat beam spot.

The third argument sets the distance from the beam source position to the focal point $k$ in cm at which the undisturbed beam (i.e. transport in vacuum) has the width and shape specified by the BEAMSIGMA card. A positive value $k > 0$ describes a defocused beam with the focal point upstream the beam at a distance $k$, while a negative value $k < 0$ results in a focused beam with the focal point downstream, relative to the source position of the beam. If $k = 0$ the beam has it focus coinciding the beam source position. The beam model follows equation (25) in reference [30], which gives a beam with a slant emittance ellipse. The same reference describes the Gaussian case only, however $k$ affects any beam shape in SHIELD-HIT12A.

In case of a beam divergence larger than $2\pi$, i.e. $> 6283.2$ mrad, then an isotropic distribution is assumed. Here, random sources are simulated located on a sphere with a radius given by the third argument. Each initialized particle will have a random direction. The second argument has no effect. This is useful for simulating the environment found in space.
If this card is not specified, all values are set by default to zero.

**BEAMPOS**: Optional card. 3 arguments that specify the start position of the beam in \((x,y,z)\) coordinates, respectively. All units are in cm.

By default the beam starts at the origin \((0,0,0)\).

**BEAMSIGMA**: Optional card. 2 arguments that specify the lateral extension of a Gaussian-shaped beam by the sigmas in cm along X and Y, respectively. If both values are negative, then a flat square distribution is generated, where \(\sigma_x\) and \(\sigma_y\) represent half the width of the sides. If \(\sigma_x\) is larger than 0 and \(\sigma_y\) is smaller than 0, then a flat circular distribution is generated with radius \(|\sigma_y|\). The \(\sigma_x\) value is unused here, but it must be larger than 0.0. If the focus is specified in the optional BEAMDIV card then the beam size and shape are obtained at the focal point.

Default values are zero sigma in X and Y direction.

**BMODMC**: Optional card. 1 argument that specifies the mode of the beam modulator simulation: 0 for modulus beam modulator version retaining the spatial period information of the beam modulator (such as in ripple filters); 1 for Monte Carlo sampling of beam modulator material thickness—useful when generating SPC files (see section 3.5.7) and simulating range modulator wheels.

Default value is 0.

**BMODTRANS**: Optional card. 1 argument that specifies the interpretation of data in the input files loaded with the USEBMOD card: 0 lists of material thicknesses (default), 1 lists of vacuum thicknesses.

Default value is 0.

**DELTAE**: Optional card. 1 argument that specifies the relative mean energy loss per transportation step, that is a finite fraction smaller than unity. This value times the particle energy at a given step determines the absolute energy loss.

Default is 0.05. (Meaning 5%).

**DEMIN**: Optional card. 1 argument that specifies the minimum allowed energy loss per transportation step in MeV/n.

Some models are only valid if the absolute magnitude of energy loss (as calculated by DELTAE) is larger than a minimal value, such as Moliere multiple scattering.

Default value is 0.025 MeV/n.

**EMTRANS**: Future switch for photon/electron/positron transportation. Not implemented, default is 0 (off).

**EXTSPEC**: Specifies if external source file to be read with USECBEAM is an array of monoenergetic beams (EXTSPEC 0) or a binned spectrum (EXTSPEC 1). If a binned spectrum, then the energies are assigned to the lower energy of that bin. The last bin marks the endpoint of the energy spectrum, and its weighting value is thus ignored.

This card is optional, and default is 0 (no spectrum). The card has only effect if USECBEAM is called after EXTSPEC.
HIPROJ: If JPART0=25 then the of this card 2 arguments specifies the number of nucleons $A$ and the charge $Z$ of the beam particles. $Z$ has to be larger than 2, since nucleons with $Z=1$ and $Z=2$ have their own particle ID JPART0, as listed in table A.2.

If this card is omitted, then default values are $Z=6$ and $A=12$.

JPART0: Optional card. 1 argument that specifies the primary particle ID, see table A.2. If JPART0=25, then the HIPROJ card can be called as well. If HIPROJ is not called, then carbon-12 is assumed by default.

If JPART0 is not specified, then the default particle 2 (protons) are assumed.

MAKELN: Optional card. 1 argument that invokes phase-space output of all secondary neutrons with energies below 14.5 MeV which are created within the target: 1 output is written to for028, see section 4.1, 0 no output of secondary neutrons.

Default value is 0.

MSCAT: Optional card. 1 argument that switches between the available types of multiple scattering: 1 for Gaussian- and 2 for Moliere-type multiple scattering.

Moliere multiple scattering (2) is activated by default.

NEUTRFAST: Optional card. Toggles fast neutron transport for energies above 14.5 MeV. If set to 0, no neutrons above 14.5 MeV are transported.

Default is 1, that is fast neutron transport is activated.

NEUTRLCUT: Optional card. Lower energy cut-off value for neutron transport in MeV. Must be lower than 14.5 MeV but larger or equal 0.0 MeV. (This was previously called OLN in older versions of SHIELD-HIT.)

Default value is 0.0. In this case, neutrons are transported until they are absorbed or exit the simulation universe.

NSTAT: Optional card. 2 arguments that specify the requested total number of primary particles to be simulated and the number of transported primaries after which an intermediate save to the scorers will be invoked, respectively.

Default values are 10000 primary particles simulated, and intermediate saving after 5000 transported primary particles.

Setting the first argument of to -1 will cause an infinite amount of particles to be transported (or until integer overflow). This is useful in combination with the –time commandline option.

Setting the second argument to 0 or a negative number will disable saving the results, until either –time or the requested total number of primary particles have been simulated.

NUCRE: Optional card. 1 argument that switches nuclear reactions on or off: 1 with nuclear reactions; 0 all nuclear reactions are turned off. (This card was called INUCRE in earlier versions of SHIELD-HIT.)

Default value is 1, i.e. nuclear reactions are activated.

RNDSEED: Optional card. 1 argument that specifies the random number seed. Using the same seed for an identical simulation should yield the same results.

Default value is 89736501.
STRAGG : Optional card. 1 argument that switches between the available models for energy straggling: 1 for Gaussian- and 2 for Vavilov-type energy straggling; 0 for no straggling. Vavilov straggling (2) is activated by default.

TMAXO : Optional card. 2 arguments that specify the initial energy of the primary particle in MeV/nucleon and the energy spread in MeV/nucleon, expressed as one standard deviation of a Gaussian distribution (1σ). If a negative value is given, then it is treated as momentum [MeV/c].

Default values are 250 MeV/nucleon and zero energy spread.

USEBMOD : Optional card. 2 arguments that specify the zone number (integer) and an external beam modulator file. See section 3.3.4. No file name is specified by default, and by default an invalid zone number is specified.

USECBEAM : Optional card. 1 argument that specifies the filename of an optional external beam source file. See section 3.3.2. No file name is specified by default

USEPARLEV : Optional card. 1 argument that specifies the file name of an external file. The file may contain model parameter values that override the default PARLEV model parameters. See section 3.3.3.

No file name is specified by default.

3.3.2 USECBEAM - External source file

If a file name is specified, e.g. sobp.dat, a projectile will be sampled from the file. Beam settings specified in the input file beam.dat such as BEAMSIGMA and TMAXO will be overridden. The weighting is typically taken as the particle number for the individual beam components (or spots). The total particle number is passed on in the header of the .bdo in case a VOXSCORE card is specified in detect.dat.

Example of a typical beam file, which was generated from a hadron therapy carbon ion control file:

```
*ENERGY (GEV) X (CM) Y (CM) FWHM (CM) WEIGHT
0.270550 -1.00 1.00 0.48 9.3700e+06
0.270550 -0.90 1.00 0.48 9.3700e+06
0.270550 -0.80 1.00 0.48 9.3700e+06
0.270550 -0.70 1.00 0.48 9.3700e+06
0.270550 -0.60 1.00 0.48 9.3700e+06
0.270550 -0.50 1.00 0.48 9.3700e+06
0.270550 -0.40 1.00 0.48 9.3700e+06
0.270550 -0.30 1.00 0.48 9.3700e+06
0.270550 -0.20 1.00 0.48 9.3700e+06
0.270550 -0.10 1.00 0.48 9.3700e+06
(...)
```

Notice that the energy is in GeV to maintain compatibility with other Monte Carlo codes. Lines starting with * are ignored and can be used for comments.

If 6 columns are specified, the following format is assumed:

1. Note that [MeV/nucleon] is different from [MeV/amu].
The FWHM parameter is the Gaussian FWHM (and not sigma), contrary to the BEAM-SIGMA card. However, if FWHM is negative (both x and y in the 2-d case), then a box with similar side length is generated. If only one of the two values are negative in the 2-d case, then a flat circular distribution is assumed, with FWHMy as the circular radius. FWHMx is currently ignored, but may be used for ellipsoid beamlets in future.

If 7 columns are specified, the following format is assumed:

<table>
<thead>
<tr>
<th>*ENERGY(GEV)</th>
<th>SigmaT0(GEV)</th>
<th>X(CM)</th>
<th>Y(CM)</th>
<th>FWHMx(cm)</th>
<th>FWHMy(cm)</th>
<th>WEIGHT</th>
</tr>
</thead>
<tbody>
<tr>
<td>0.270550</td>
<td>1.1e-03</td>
<td>-1.00</td>
<td>1.00</td>
<td>0.48</td>
<td>0.44</td>
<td>9.3700e+06</td>
</tr>
<tr>
<td>0.270550</td>
<td>1.1e-03</td>
<td>-0.90</td>
<td>1.00</td>
<td>0.48</td>
<td>0.44</td>
<td>9.3700e+06</td>
</tr>
<tr>
<td>0.270550</td>
<td>1.1e-03</td>
<td>-0.80</td>
<td>1.00</td>
<td>0.48</td>
<td>0.44</td>
<td>9.3700e+06</td>
</tr>
<tr>
<td>0.270550</td>
<td>1.1e-03</td>
<td>-0.70</td>
<td>1.00</td>
<td>0.48</td>
<td>0.44</td>
<td>9.3700e+06</td>
</tr>
<tr>
<td>0.270550</td>
<td>1.1e-03</td>
<td>-0.60</td>
<td>1.00</td>
<td>0.48</td>
<td>0.44</td>
<td>9.3700e+06</td>
</tr>
<tr>
<td>0.270550</td>
<td>1.1e-03</td>
<td>-0.50</td>
<td>1.00</td>
<td>0.48</td>
<td>0.44</td>
<td>9.3700e+06</td>
</tr>
<tr>
<td>0.270550</td>
<td>1.1e-03</td>
<td>-0.40</td>
<td>1.00</td>
<td>0.48</td>
<td>0.44</td>
<td>9.3700e+06</td>
</tr>
<tr>
<td>0.270550</td>
<td>1.1e-03</td>
<td>-0.30</td>
<td>1.00</td>
<td>0.48</td>
<td>0.44</td>
<td>9.3700e+06</td>
</tr>
<tr>
<td>0.270550</td>
<td>1.1e-03</td>
<td>-0.20</td>
<td>1.00</td>
<td>0.48</td>
<td>0.44</td>
<td>9.3700e+06</td>
</tr>
<tr>
<td>0.270550</td>
<td>1.1e-03</td>
<td>-0.10</td>
<td>1.00</td>
<td>0.48</td>
<td>0.44</td>
<td>9.3700e+06</td>
</tr>
</tbody>
</table>

where SigmaT0 marks the energy spread of the primary energy assuming a Gaussian distribution. The energy spread is 1σ standard deviation in GeV/nucleon. This will override the second argument specified with the TMAX0 card in beam.dat.

### 3.3.3 USEPARLEV - External parameter file

The predefined values of the PARLEV parameters can be redefined by those given in the optional PARLEV input file thereby overwriting the carefully selected and benchmarked default values. Further information on PARLEV parameters can be found in Appendix A.8 and the default values are listed in Table A.6.

It is recommended to use the default values. They have been tested against available data. Most likely you want to use the default values to remove the USEPARLEV card in the beam.dat file entirely.

---

*Thereby compatibility to FLUKA can be retained.*
Example of typical PARLEV input file with two columns. The first column specifies the ID of a PARLEV parameter while the second specifies its new value (either space or tab separated).

<table>
<thead>
<tr>
<th>ID</th>
<th>Value</th>
</tr>
</thead>
<tbody>
<tr>
<td>33</td>
<td>1.0</td>
</tr>
<tr>
<td>34</td>
<td>1.0</td>
</tr>
<tr>
<td>39</td>
<td>1.20</td>
</tr>
<tr>
<td>40</td>
<td>1.20</td>
</tr>
</tbody>
</table>

Lines starting with * are ignored and can be used for comments.

### 3.3.4 USEBMOD - Beam modulator

A beam modulator is a special geometry, which transports a particle hitting it by a certain amount in the Z-direction, following a user-specified transportation table. The amount of transportation can be sampled randomly from that table, or be linked to a certain position along X and/or Y axis.

The beam modulator geometry is useful for simulating complex geometries such as:

- Ripple filters [31] (RiFi).
- Ridge filters
- Range modulator wheels

A beam modulator can be inserted into the beam at an arbitrary position. This works by specifying a zone in `geo.dat`, and attaching a user generated, external modulator file to that particular zone using the `USEBMOD` command in `beam.dat`.

The beam modulator file is structured as a series of bins, holding a certain thickness. The first line contains the start position of the first bin, and the subsequent lines contain the end position of the bin and its respective thickness.

Shown below are the first lines of a typical user generated (1D) displacement map:

<table>
<thead>
<tr>
<th>Y coordinate</th>
<th>Z displacement</th>
</tr>
</thead>
<tbody>
<tr>
<td>-0.797654000000</td>
<td>0.299843308865</td>
</tr>
<tr>
<td>-0.797583286279</td>
<td>0.296589497459</td>
</tr>
<tr>
<td>-0.79076849238</td>
<td>0.292032749859</td>
</tr>
<tr>
<td>-0.790878035008</td>
<td>0.28747741389</td>
</tr>
<tr>
<td>-0.784271597967</td>
<td>0.28292066629</td>
</tr>
<tr>
<td>-0.771129305477</td>
<td>0.277060982496</td>
</tr>
<tr>
<td>-0.771072840203</td>
<td>0.274457933371</td>
</tr>
<tr>
<td>-0.76466403162</td>
<td>0.269901185771</td>
</tr>
<tr>
<td>-0.751366459627</td>
<td>0.26599378882</td>
</tr>
<tr>
<td>-0.744760022586</td>
<td>0.26143704122</td>
</tr>
<tr>
<td>-0.738167701863</td>
<td>0.257531055901</td>
</tr>
</tbody>
</table>

(\ldots many more data points\ldots)

SHIELD-HIT12A assumes by default that the beam modulator file lists the thickness of the ripple filter as a function of position. If a displacement in vacuum is stated then the `BMODTRANS` card must be set in `beam.dat`. 
Line breaks and lines starting with * are ignored and can be used for comments.

The beam modulator can either be one or two dimensional. If the beam modulator is one dimensional, the periodic structure changes along the Y axis only. The structure is repeated over the entire beam modulator on the y-position of the incoming particle. Alternatively, the structure can be sampled by a random number using the BMODMC card in beam.dat. This way the spatial structure is lost, but pencil beams hitting the same spot on the modulator will still be modulated over the entire structure, where the Y-bins are used as weights instead. This is useful for generating TRiP98 depth dose kernels, which require a full modulated pencil beam.

The data must be either space or tab delimited and the maximum number of steps is set to 200. The size of the beam modulator is specified in the geo.dat file. In the example shown below, the beam modulator is represented by the zone number 3. Successful application of the ripple filter will be stated at run time when the first particle hits the ripple filter surface.

**Important:** The thicknesses or displacements of the external file should not exceed the zone dimensions along the z-axis, as this may lead to undefined behaviour of the code.

Example: geo.dat with ripple filter (body and zone number 3).

```
0 0 C-12 on water with RiFi 21.10.2011
RPP 1 -100.0 100.0 -100.0 100.0 0.0 100.0
RPP 2 -150.0 150.0 -150.0 150.0 -1.300 -1.0
RPP 3 -8.0 8.0 -8.0 8.0 -1.300 -1.0
RPP 4 -200.0 200.0 -200.0 200.0 -200.0 200.0
END
TAR 1 +1
AIR 2 +2 -1 -3
RIF 3 +3
OUT 4 +4 -2
END
1 2 3 4
1 2 3 0
```

The beam.dat links the zone number 3 to a certain beam modulator file, by adding the line

```
USERIFI 3 rifi.dat
```

Additional information about the implementation of the beam modulator as a ripple filter can be found in [32] and [10].

Two-dimensional beam modulators can be specified as well. SHIELD-HIT12A detects this automatically if the specified file contains three columns of data instead of two. The supplied file should be a mesh of data, describing a full period of the ripple filter. Either the X coordinate can increment fastest followed by the Y position, or vice versa, yet the data must be ordered in increasing order along X and Y. The third column gives the thickness in Z direction. Again, X and Y denote the end-positions of the bins, except for the first line which defined the starting positions of the first bin. Line breaks and lines starting with * are ignored. The maximum number of allowed lines is hard coded to be 200x200 steps (i.e. 40,000 data points). The table below shows an excerpt of a 2D ripple filter file example:

```
* 2-D ripple filter example.
* X coordinate [cm]  Y coordinate [cm]  Z displacement [cm]
0.0000  0.0000
```


0.0000  0.0030  0.0000
0.0000  0.0060  0.0000
0.0000  0.0090  0.0000
0.0000  0.0120  0.0000
0.0000  0.0150  0.0058
0.0000  0.0180  0.0139
(...many lines skipped here...)
0.0060  0.1350  0.0278
0.0060  0.1380  0.0175
0.0060  0.1410  0.0082
0.0060  0.1440  0.0000
0.0060  0.1470  0.0000
0.0060  0.1500  0.0000

0.0090  0.0000  0.0000
0.0090  0.0030  0.0000
(...many lines skipped here...)
0.1500  0.1470  0.0000
0.1500  0.1500  0.0000
3.3.5 Examples - Beam initialization

Example 1: A SOBP carbon ion beam with ripple filter

Example of typical beam.dat file:

```plaintext
* Input file beam.dat for the SHIELD-HIT Transport Code
RNDSEED 89736501 ! Random seed
JPART0 25 ! Incident particle type
HIPROJ 12.0 6.0 ! Carbon ions
TMAX0 400.0 0.0 ! Incident energy; (MeV/nucl)
BEAMSIGMA 2.0 2.0 ! SigmaX, SigmaY at focus point
BEAMPOS 0.0 0.0 -1.50 ! Beam XYZ start pos
NSTAT 20000 5000 ! NSTAT, Step of saving
NEUTRLCUT 0.0 ! Cutoff for neutron transport
MAKELN 0 ! 1 - Make neutron phase space file
DELTAE 0.020 ! Delta E (relative share ~0.1)
DEMIN 0.030 ! Minimum Energy step 0.025 (MeV/n)
STRAGG 2 ! Straggling: 0-Off 1-Gauss, 2-Vavilov
MSCAT 2 ! Mult. scatt 0-Off 1-Gauss, 2-Moliere
NUCRE 1 ! Nucl.Reac. switcher: 0-Off, 1-On
USEBMOD 3 rifi.dat ! Zone# and file name for beam modulator
USECBEAM sobp.dat ! Filename of beam sourcefile
```
3.4 geo.dat - Geometry

The SHIELD-HIT code uses known Combinatorial Geometry (CG), similar to the Monte Carlo particle transport codes MORSE, FLUKA.

The geometry is defined by a number of so called bodies which are primitive objects such as boxes and cylinders, which by Boolean logic are combined to zones. For instance a square slab with a round hole can be constructed using two primitives, i.e. a box subtracted with a cylinder. Similarly a hollow box can be described by a larger box subtracted with a smaller box inside it.

The zones must then be assigned to a material identifier, which is an integer defined by the user in the material definition file mat.dat, see section 3.2. Every point in the simulated universe must be assigned to a zone, else errors may occur. So, if a hollow box is constructed, one must remember also to assign the void inside the box to a material code (e.g. internal vacuum 1000). One should also understand that a single body cannot directly be assigned to a material, but must be assigned to a zone first, even if the zone just consists of a single body.

The definition of the bodies, zones and the material assignments are set in the geo.dat file, in exactly that order.

Anywhere within geo.dat lines starting with * are skipped.

3.4.1 Title

The first line of geo.dat contains two integers JDBG1, JDBG2 and a string of characters describing the title of the geometry. The format of the title card is:

- 5 columns (1-5): integer JDBG1, see below
- 5 columns (6-10): integer JDBG2, see below
- 10 columns (11-20): unused
- 60 columns (21-80): any title of the geometry

The first integer JDBG1 selects whether the file for017 containing the geometry debugging information should be kept (0) or deleted (1) after the geometry parser was initialized. The second integer JDBG2 describes the lower cutoff value of transportation step size in powers of 10, i.e. $10^{-|JDBG2|}$. If set to zero, the minimum step size is set to $10^{-4}$ cm. If any of the numbers are omitted, they are interpreted as (0). The fields for both numbers are five columns wide.

3.4.2 Bodies

A list of bodies describing the geometry follows the title card explained in the section before. A body is a primitive object with a given geometrical configuration. The available primitive objects are listed section A.6. A “top level” body (which is mandatory in the MC code FLUKA) can optionally be added which contains all other bodies, which will be the simulation universe, but this is not strictly necessary in SHIELD-HIT12A. There are no requirements to how this body must be named or where it is placed in the body list, but for convenience it can be either the first or last body. The list of bodies must be terminated with the END card. The format of a body card is:

- 2 columns (1-2): unused, e.g. two empty spaces
- 3 columns (3-5): name of the body (see sec. A.6)
- 1 column (6): unused
Chapter 3. Input files

- 4 columns (7-10): an integer describing the number of the body
- 6 x 10 columns (11-70): 6 arguments describing the body as mentioned in [A.6]

A continuation card may be necessary, depending on what body is described. A continuation card must be blank in the first 10 columns (i.e. no name and no body number). The body section must be terminated with an END card.

3.4.3 Voxelized Body

The V0X body is a special body, which will load a CT-cube formatted as the VOXELPLAN format used by TRiP98. A CT-cube in the VOXELPLAN format consists of two files:

1. a header file with the .hed suffix, containing human readable information on the properties of the CT-cube, such as number of slices, number of bins along the X and Y axis, voxel dimensions etc.
2. a CT-cube file, containing the voxel Hounsfield units in binary floating point format.

The V0X body is treated as if it would be a normal BOX, however the dimensions are automatically extracted from the header file. However, SHIELD-HIT12A still needs to know where to place the CT-cube, and whether it should be rotated. SHIELD-HIT12A always assumes that the point (0,0,0) in the simulated universe coordinate system will be the isocenter. Thus, the isocenter position in SHIELD-HIT12A should be marked inside the CT-cube. This is conveyed to SHIELD-HIT12A using the first five arguments of the V0X card:

1. X - simulated universe isocenter x-position inside CT cube coordinate system [cm]
2. Y - simulated universe isocenter y-position inside CT cube coordinate system [cm]
3. Z - simulated universe isocenter z-position inside CT cube coordinate system [cm]
4. Couch angle [degrees]
5. Gantry angle [degrees]
6. Optional planned target dose as specified in TRiP98 [Gy]

The last optional argument needs more clarification: when the V0XSORE card is used in detect.dat, then TRiP98 formatted data cubes may be created during postprocessing. These TRiP98 formatted cubes store the dose value units relative to the planned PTV dose. SHIELD-HIT12A calculates however the dose per simulated primary particle. So, in order to convert to the relative TRiP dose, SHIELD-HIT12A thus needs to know 1) The planned target dose (given by the last argument in the V0X card), and 2) the total particle budget. The total particle budget are derived from the external beam specifications loaded with the USECBEAM card, assuming these were crafted with the actual particle numbers deposited in each beam spot. Note, that specifying this option does not change any scoring or transport behaviour of SHIELD-HIT12A. Any raw output .bdo will still contain the values in their respective units, the difference is just, that the information on the target dose is shipped along in the header of the output files, providing useful information to any external postprocessing routines.

Arguments for second card:

1. The zone description follows afterwards. In order to construct different zones out of the bodies defined at the beginning of `geo.dat`, Boolean logic is used. Examples of how to construct zones out of two bodies are shown in figure 3.2.

A line in the zone description consists of:

- 2 columns (1-2): unused, e.g., two empty spaces
- 3 columns (3-5): name of the zone
- 5 columns (6-10): an optional integer describing the number of the zone

![Figure 3.2: Two bodies (marked as 1 and 2) describing 5 different zones.](image)
• 9 x (2+5) columns (11-72): where each set of 9 columns are composed of two components. The two first columns hold the Boolean operator OR or are left blank if needed. The following 5 columns contain a signed (+,−) integer body number. It is essential that the alignment is strictly kept.

If more than 9 bodies are used for the zone description additional continuation cards can be added directly afterwards. These continuation lines must not have a zone name, i.e. there must be no entry at columns 3-5. The zone section must also be terminated with END card.

Example:

*..<-><--->..<--->OR<--->OR<--->OR<--->OR<--->OR<--->OR<--->OR<--->OR<-----*
001  +1  -2  -3  -4  -5
002       +2
003       +3
004       +4
005       +5
006       +6  -1
END

It is important that any point (except for zone boundaries) where a particle may go, is described by exactly one zone, no more, no less. SHIELD-HIT12A may produce unexpected or random behavior if particles cross areas where two zones overlap, or a zone which is not defined. To some extent SHIELD-HIT12A will warn the user in such cases, but SHIELD-HIT12A may not catch all cases, therefore a careful assignment of zones is essential.

3.4.5 Media

Finally, what follows is the assignment of media to the various zones. External vacuum behaves like a black hole, any particle hitting it will be disintegrated and not transported further. The “top level” zone must be assigned to external vacuum (id = 0). Inside this zone you can place a zone with internal (i.e. real physical) vacuum (id = 1000) or any other material. In internal vacuum regular particle transport takes place, as opposed to the external vacuum. Even if not strictly mandatory, the external vacuum should encompass the entire system to prevent infinite loops where stray particles are transported forever, and to avoid unintended behavior of the code. The media assignment is done in two equally formatted lists. The first list is a list of zones:

• 14x5 columns (1-70): list of integer zone numbers

The second list are the material numbers:

• 14x5 columns (1-70): list of integer media identifiers

If one line is not enough, then it is simply continued on the following line, until all zones are described. All zones must be assigned to medium, thus, both lists must be equally long.

3.4.6 Examples - Geometry

Example 1: Water tank surrounded by air

A simple example of the input geometric file geo.dat is given below.
Example 2: Proton therapy beam line

A more complex example is a model of the beam line at the cyclotron facility in Clatterbridge, UK. It features two zones with external vacuum, where the second one is used to collimate the beam:

```
0 0 \hspace{1cm} \text{COO} \hspace{1cm} \text{beam line Clatterbridge} \hspace{1cm} 13.05.2011
RCC 1 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} -25.6 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 1.00
\hspace{1cm} 1.0
RCC 2 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} -25.6 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 1.00
\hspace{1cm} 0.3
RCC 3 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} -22.6 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 0.003
\hspace{1cm} 1.0
RCC 4 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} -0.66 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 0.660
\hspace{1cm} 0.286
RCC 5 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 0.002
\hspace{1cm} 1.0
RCC 6 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 5.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 0.005
\hspace{1cm} 1.0
RCC 7 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} -30.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 35.005
\hspace{1cm} 1.0
RCC 8 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 27.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 0.001
\hspace{1cm} 2.0
RCC 9 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 55.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 1.00
\hspace{1cm} 2.0
RCC 10 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 55.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 1.00
\hspace{1cm} 2.0
RCC 11 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 115.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 0.002
\hspace{1cm} 2.0
RCC 12 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 115.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 0.002
\hspace{1cm} 2.0
RCC 13 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 175.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 7.80
\hspace{1cm} 2.0
RCC 14 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 175.0 \hspace{1cm} 0.0 \hspace{1cm} 0.0 \hspace{1cm} 7.80
\hspace{1cm} 1.70
```
RCC 15 0.0 0.0 182.80 0.0 0.0 0.80
20.0
RCC 16 0.0 0.0 182.80 0.0 0.0 0.80
1.5
RCC 17 0.0 0.0 5.005 0.0 0.0 178.995
25.0
RCC 18 0.0 0.0 184.0 0.0 0.0 30.0
20.0
RCC 19 0.0 0.0 -100.0 0.0 0.0 500.0
100.0
RCC 20 0.0 0.0 -200.0 0.0 0.0 1000.0
200.0
RCC 21 0.0 0.0 -26.0 0.0 0.0 1.00
15.0
RCC 22 0.0 0.0 -26.0 0.0 0.0 1.00
0.3
END

TAR  1  +18
BR1  2  +1  -2
BR2  3  +9  -10
BR3  4  +13 -14
BR4  5  +15 -16
HL1  6  +2
HL2  7  +10
HL3  8  +14
HL4  9  +16
SF1  10 +3
SF2  11 +5
STP  12 +4
KWD  13 +6

VAC  14 +7  -1  -3  -4  -5  -6  -21
WHL  15 +8
ICM  16 +11
ICA  17 +12
CAR  18 +17 -8  -9  -11 -12 -13 -15
OAR  19 +19 -7  -17 -18
OUT  20 +20 -19
IBH  21 +21 -22
IHL  22 +22
END

1  2  3  4  5  6  7  8  9  10  11  12  13  14
15  16  17  18  19  20  21  22
1  2  2  2  2  2  2  3  3  9  9  2  5  1000
8  6  4  3  3  0  0 1000
Example 3: Voxelized CT cube

This example demonstrates how a TRiP formatted CT cube is loaded. The couch angle is set to -90 degrees, and the iso-center (which is the (0,0,0) point of the SHIELD-HIT12A coordinate system) is located at (15.5, 15.5, 8.7) cm in the CT cube coordinate system.

```plaintext
*---><---><--------><------------------------------------------------|

 0 0 Voxelplan test

*---><---><--------><--------><--------><--------><--------><-------->

* sphere - black body
SPH 1 0.0 0.0 0.0 10000.0 0.0 0.0

*---><---><--------><--------><--------><--------><--------><-------->

* sphere - vacuum
SPH 2 0.0 0.0 0.0 1000.0 0.0 0.0

*---><---><--------><--------><--------><--------><--------><-------->

* box/vox under consideration, water or CT
VOX 3 15.5 15.5 8.6 -90.0 0.0

../../res/TST001/tst001000

*---><---><--------><--------><--------><--------><--------><-------->

END

001  +1  -2
002  +2  -3
003  +3

END

1  2  3

* 0 - black hole, 1000 vacuum, 1 - water
0 1000 1
```
3.5 \textit{detect.dat} - Scoring

Different geometrical scoring types, called \textit{estimators}, are available for auxiliary scoring: \textit{GEOMAP}, \textit{ZONE}, \textit{DZONE}, \textit{CYL}, \textit{MSh}, \textit{PLANE DCYL}, \textit{DMSH}, \textit{DMSHZ} and \textit{DPLANE}. Each estimator can be invoked with a scoring card and possible following cards in the file \textit{detect.dat}. Most estimators need additional specification of a \textit{detector}, which controls what kind of quantity is scored. The available detectors are described afterwards in section 3.5.10.

Each field is exactly 10 chars long, except for the output filename, which may be 4096 characters long. The output of each estimator is written, in binary form, to a \textit{.bdo} file, which is specified by the user. The \textit{.bdo} suffix is automatically attached, if missing. The recommended post-processing scripts \texttt{convertmc} is open-source and is currently not included in the \texttt{SHIELD-HIT12A} distribution. It must be installed separately from the \texttt{pymchelper} package.

\url{https://github.com/DataMedSci/pymchelper}

3.5.1 \textit{GEOMAP} - Mapping the geometry

\textit{GEOMAP}: Mapping the geometry. This card is useful for debugging the geometry. Using the arguments below, a 1, 2 or 3-D mesh can be created, where the points will hold either the zone number, medium number specified in \textit{mat.dat} or the density of the medium in [g/cm$^3$]. For voxelized structures, the actual density in the individual voxels will be scored.

The card is only invoked during initialization, and has therefore no affect on run-time performance of \texttt{SHIELD-HIT12A}. The resulting \textit{.bdo} files are independent of any particles transported, i.e. when the user is debugging the geometry, the particle number can be set to zero (either by using the \texttt{-n} command line option or the \texttt{NSTAT} card in \textit{beam.dat}, in order to quickly generate the output files.

In case of parallelization using the \texttt{-N} flag, the \textit{GEOMAP} card will be ignored and \textit{.bdo} files are not generated, as it makes no sense to clutter the directory with multiple data cubes with identical data in.

The arguments for the \textit{GEOMAP} files are:

1. $X_{\text{min}}$: lowest X position.
2. $Y_{\text{min}}$: lowest Y position.
3. $Z_{\text{min}}$: lowest Z position.
4. $X_{\text{max}}$: highest X position.
5. $Y_{\text{max}}$: highest Y position.
6. $Z_{\text{max}}$: highest Z position.

Arguments for the second card:

1. $X_{\text{bin}}$: number of bins in X direction.
2. $Y_{\text{bin}}$: number of bins in Y direction.
3. $Z_{\text{bin}}$: number of bins in Z direction.
4. unused
5. \texttt{ZONE/MEDIUM/RHO}: Quantity to be mapped. This can be either the number of the zone (\texttt{ZONE}), or the number of the medium (\texttt{MEDIUM}), or the density of the material (\texttt{RHO})
6. \texttt{outputfile}: output file name.
3.5.2 ZONE - Scoring by zone

ZONE: Scoring by zones $w_i$. Zones are defined from bodies in geo.dat. A single or a range of zones can be specified. Arguments are:

1. $w_i$: first zone of a range of zones to be scored.
2. $w_j$: optional last zone of a range of zones. If this is not specified, then only a single zone $w_i$ will be scored. Else every zone within the interval $[w_i, w_j]$ will be scored.
3. unused.
4. JPART: particle type. See list in section A.3. Setting this value to -1 scores all particles. If heavy ions are scored (JPART=25) then a continuation card is required, where the first argument is an integer representing the particle charge $Z$, and the second argument is an integer representing the particle mass $A$.
5. DETECTOR: quantity that should be scored. Important: Zone scoring works straightforwardly only with the detectors ENERGY, CROSSFLU, DLET, COUNTER and PET as described in section 3.5.10. All other detectors can be applied as well, but require knowledge of the zone volume, which is not known by SHIELD-HIT12A (unlike the MSH and CYL scorer). Therefore the user must divide the scoring result with the corresponding zone volume (in cm$^3$) for all other detectors than those listed above. This may be changed in a future release (see ticket #177).
6. outputfile: output file name.

Analogue to CYL and MSH, a continuation card is only required if JPART=25 or if DETECTOR is specified as LETFLU (and only LETFLU). Arguments for the continuation card are:

1. $Z$: an integer specifying the charge of the particle to be scored.
2. $A$: an integer specifying the number of nucleons of the particle to be scored. Setting this value to -1 scores all isotopes.
3. MEDIUM: an integer specifying the medium for LET weighted scoring. If left unspecified, the default is the medium material that the bin is made of. Any other integer from the medium list specified in mat.dat is valid here. In that case, scoring will apply the stopping powers of the medium specified here. Transport will irrespective of the scoring always be done in the as medium specified in geo.dat. This is useful for calculating stopping power ratios needed by various cavity theories in radiation dosimetry.

4. unused.
5. unused.
6. unused.
3.5.3 CYL - Cylindrical scoring

**CYL**: Cylindrical scoring. Cylindrical coordinates are used. Currently, scoring can only be done along the Z axis. This card requires a second succeeding card. Arguments are:

1. \( R_{\text{min}} \): inner radius of scoring cylinder. If \( R_{\text{min}} > 0 \) then the scoring volume is a cylinder shell.
2. \( \theta_{\text{min}} \): minimum value of angular segment.
3. \( Z_{\text{min}} \): start position of cylindrical scoring in Z direction.
4. \( R_{\text{max}} \): outer radius of scoring cylinder.
5. \( \theta_{\text{max}} \): maximum angle of angular segment. If this value is set to number larger than \( 2\pi \) then \( \theta_{\text{max}} \) will be fixed at \( 2\pi \), covering the entire disc.
6. \( Z_{\text{max}} \): end position of cylindrical scoring in Z direction.

Arguments for the second card:

1. \( R_{\text{bin}} \): number of bins in radial direction.
2. \( \theta_{\text{bin}} \): number of angular bin segments.
3. \( Z_{\text{bin}} \): number of bins in Z direction.
4. **JPART**: particle type. See list in section A.3 Setting this value to -1 scores all particles. If heavy ions are scored (JPART=25) then a third continuation card is required, where the first argument is an integer representing the particle charge \( Z \), and the second argument is an integer representing the particle mass \( A \).
5. **DETECTOR**: quantity that should be scored. A list of available detectors is given in section 3.5.10
6. **outputfile**: output file name.

Analogue to ZONE and MSH a continuation card is only required if **JPART=25** or if **DETECTOR** is specified as LETFLU (and only LETFLU). Arguments for the third card are:

1. **Z**: an integer specifying the charge of the particle to be scored.
2. **A**: an integer specifying the number of nucleons of the particle to be scored. Setting this value to -1 scores all isotopes.
3. **MEDIUM**: an integer specifying the medium for LET weighted scoring. If left unspecified, the default is the medium material that the bin is made of. Any other integer from the medium list specified in mat.dat is valid here.

4. unused.
5. unused.
6. unused.
3.5.4 MSH - Cartesian scoring

**MSH**: Cartesian mesh scoring card. This card requires a second succeeding card. Arguments for the first card:

1. $X_{\text{min}}$: lowest X position.
2. $Y_{\text{min}}$: lowest Y position.
3. $Z_{\text{min}}$: lowest Z position.
4. $X_{\text{max}}$: highest X position.
5. $Y_{\text{max}}$: highest Y position.
6. $Z_{\text{max}}$: highest Z position.

Arguments for the second card:

1. $X_{\text{bin}}$: number of bins in X direction.
2. $Y_{\text{bin}}$: number of bins in Y direction.
3. $Z_{\text{bin}}$: number of bins in Z direction.
4. JPART: particle type. See list in section A.3. Setting this value to -1 scores all particles.
   - If heavy ions are scored (JPART=25) then a third continuation card is required, where the first argument is an integer representing the particle charge $Z$, and the second argument is an integer representing the particle mass $A$.
5. DETECTOR: quantity that should be scored. A list of available detectors is given in section 3.5.10.
6. outputfile: output file name.

Analogue to ZONE and CYL a continuation card is only required if JPART=25 or if DETECTOR is specified as LETFLU (and only LETFLU). Arguments for the third card are:

1. $Z$: an integer specifying the charge of the particle to be scored.
2. $A$: an integer specifying the number of nucleons of the particle to be scored. Setting this value to -1 scores all isotopes.
3. MEDIUM: an integer specifying the medium for LET weighted scoring. If left unspecified, the default is the medium material which the bin is made of. Any other integer from the medium list specified in *mat.dat* is valid here.
4. unused.
5. unused.
6. unused.
3.5.5 PLANE - Scoring by 2D plane

PLANE: Scoring by a single 2-D infinite plane \( s \), independently of \( geo.dat \). The plane is specified by a random point which must lie in the plane, and with the normal vector of that plane. The normal can either be specified in this card, or - if omitted - the normal vector will automatically calculated from the beam direction specified by the \textsc{beamtheta} and \textsc{beamphi} cards in \textit{beam.dat}.

Arguments are:

1. \( S_x \): X coordinate of point in plane.
2. \( S_y \): Y coordinate of point in plane.
3. \( S_z \): Z coordinate of point in plane.
4. \( n_x \): x component of normal vector.
5. \( n_y \): y component of normal vector.
6. \( n_z \): z component of normal vector.

Arguments for the second card:

1. unused.
2. unused.
3. unused.
4. \textsc{Jpart}: particle type. See list in section 4.3. Setting this value to -1 scores all particles. If heavy ions are scored (\textsc{Jpart}=25) then a third continuation card is required, where the first argument is an integer representing the particle charge \( Z \), and the second argument is an integer representing the particle mass \( A \).
5. \textsc{detector}: quantity that should be scored. A list of available detectors is given in section 3.5.10.
6. \textit{outputfile}: output file name.

Analogue to \textsc{cylic} and \textsc{mshe}, a continuation card is only required if \textsc{Jpart}=25 or if \textsc{detector} is specified as \textsc{letflu} (and only \textsc{letflu}). Arguments for the continuation card are:

1. \( Z \): an integer specifying the charge of the particle to be scored.
2. \( A \): an integer specifying the number of nucleons of the particle to be scored. Setting this value to -1 scores all isotopes.
3. \textsc{medium}: an integer specifying the medium for LET weighted scoring. If left unspecified, the default is the medium material that the bin is made of. Any other integer from the medium list specified in \textit{mat.dat} is valid here. In that case, scoring will apply the stopping powers of the medium specified here. Transport will irrespective of the scoring always be done in the as medium specified in \textit{geo.dat}. This is useful for calculating stopping power ratios needed by various cavity theories in radiation dosimetry.

4. unused.
5. unused.
6. unused.
3.5.6 DZONE, DCYL, DMSH and DPLANE - Differential scoring

DZONE, DCYL, DMSH and DPLANE: Differential zone, cylindrical and mesh scoring. Same as ZONE, CYL, MSH and PLANE respectively, but with an extra card that specifies the differential dimension of the scoring. The differential scoring card must be inserted before the heavy ion selection card, if that is used as well.

Arguments for the extra card:

1. $\Delta_{\text{start}}$: Start value of differential binning.
2. $\Delta_{\text{stop}}$: Stop value of differential binning.
3. $\Delta_n$: Number of bins. If this value is negative, then binning is done in logarithmic steps, and the $\Delta_{\text{start}}$ parameter must be larger than zero.
4. internal use, keep this field clear.
5. Type: Type of differential scoring. Currently, ENERGY, LET and ANGLE are available, being differential in MeV/n, MeV/cm or radians, respectively.
6. unused.

The optional third card for CYL and MSH is the fourth card when using DCYL and DMSH.

If applying differential scoring on pions, their energies will be converted to specific energies (MeV/n) by multiplying with 0.1498 ($\pi^+\text{ and } \pi^-\text{ mass divided by proton mass}$).

\[\text{In the case of } \pi^0\text{ it is } 0.1448 \text{ which the } \pi^0\text{ mass divided by proton mass. However, SHIELD-HIT12A does not transport } \pi^0\text{ as it decays instantaneously.}\]
3.5.7 DMSHZ and DCYLZ - Spectral files in TRiP98 format

DMSHZ and DCYLZ: Creates a .spc file for TRiP98 format using either mesh or cylindrical scoring. The SPC format is described in http://bio.gsi.de/DOCS/TRiP98/PRO/DOCS/trip98fmtspc.html.

DMSHZ and DCYLZ is similar to DMSH DCYL, but requires the special detector SPC in the DETECTOR field of the second card. The mandatory third card specifies differential dimensions of the scoring:

1. $\Delta_{\text{start}}$: Start of differential binning in energy (MeV/A).
2. $\Delta_{\text{stop}}$: Stop of differential binning in energy (MeV/A).
3. $\Delta_n$: Number of bins. If this value is negative, then binning is done in logarithmic steps, and the $\Delta_{\text{start}}$ parameter must be larger than zero.
4. $Z_{\text{peak}}$: Peak position for placement of scoring grid along Z. If negative the peak position will be estimated by SHIELD-HIT12A in CSDA approximation, assuming material NUMMED=1.
5. Type: ENERGY, only.
6. unused.

For DMSHZ and DCYLZ scoring, the first medium NUMMED=1 specified in mat.dat can be any material, but most likely one may use water (e.g. for generating proper TRiP spec files).

The peak position $Z_{\text{peak}}$ is needed to place the adaptive scoring grid in Z direction which has the highest resolution at the Bragg-peak, and will be more coarse the further away the scoring is from the peak position. If a negative number is given, SHIELD-HIT12A will estimate the peak position using the continuous slowing down approximation (CSDA) in water. This may be insufficient when a true beam line is modeled, which might significantly shift the true $Z_{\text{peak}}$ upstream. In this case, the center of the grid can be placed manually by specifying a positive $Z_{\text{peak}}$.

The $Z_{\text{max}}$ position specified in the first card of the DMSHZ estimator will be used by SHIELD-HIT12A for setting up the Z interval wherein the Bragg-peak position will be found. For the scoring of the spectrum, however, $Z_{\text{max}}$ will be then overwritten by SHIELD-HIT12A by twice the $Z_{\text{peak}}$ position. Currently only $Z_{\text{min}} = 0$ is possible.

The specified number of Z bins, $\Delta_n$, is not propagated to the output SPC-files, but internally used by SHIELD-HIT12A to determine the Bragg peak position along the Z axis. This should not be confused with $Z_{\text{peak}}$ specified by the user, as the result of Bragg-peak position calculation is only available after transportation. The exact position of the Bragg-peak location is then written in the SPC file, needed by TRiP.

When handling divergent beams, the distance to the isocenter becomes important. For DDD and SPC file generation, a constant source to surface distance (SSD) is assumed, thus the isocenter is at a fixed position in the target, irrespective of the penetration depth of the ions. However, in treatment planning in TRiP the isocenter is located in the treatment volume, which means varying SSDs. While unimportant for non-diverging beams, the solution for divergent beams can be twofold.

1. Newer versions of TRiP provide a new parameter for the scancap command which solves this issue.
2. Older versions of TRiP one should recalculate a new set of base data for each field, which uses a proper SSD and isocenter depth corresponding to this particular field.
Contrary to the behaviour of DMSH and DCYL, DMSHZ and DCYLZ will score all (primary or secondary) pions and antiprotons as protons with the same energy per mass. The reason is that TRiP cannot handle pion or antiprotons per default. Yet one can approximate that the antiproton (when treating the secondaries from annihilation separately) resembles a proton in terms of dose deposition and stopping power. Pions have also unity charge, but are 7 times lighter than protons. A 70 MeV pion is therefore scored as a 10 MeV proton, since these two particles have comparable stopping powers at these energies.

The resulting .spc files can be inspected using the SPCinspector program found on GitHub: [https://github.com/pytrip/SPCinspector](https://github.com/pytrip/SPCinspector)
3.5.8 **TRACE - Dump particle tree**

**TRACE**: A single card, which instructs SHIELD-HIT12A to dump all transport steps of all particles into a comma delimited file. This estimator is useful for debugging and visualizing particle trajectories across the simulated universe. Each line contains the start and stop position of a particle step [cm], the JPART number, charge (Z) and nucleon (A) numbers, the kinetic energy [MeV/nucleon] and the energy loss $\Delta E$ for this step [MeV/nucleon]. If the start and stop positions are equal, then a point-wise scoring happened (e.g. at the end of a particle trajectory).

Only one **TRACE** card is allowed within *detect.dat*.

Since a lot of output is generated which is written to disk, it will slowdown the calculation massively. It is thus not recommended to record more than 100 particle trajectories.

Arguments are:

1. first particle to be scored
2. last particle to be scored
3. unused.
4. unused.
5. unused.
6. **outputfile**: output file name. `.csv` suffix will be appended automatically.

Analogue to **CYL** and **MSH**, a continuation card is only required if JPART=25 or if **DETECTOR** is specified as **LETFLU** (and only **LETFLU**). Arguments for the continuation card are:

1. **Z**: an integer specifying the charge of the particle to be scored.
2. **A**: an integer specifying the number of nucleons of the particle to be scored. Setting this value to -1 scores all isotopes.
3. **MEDIUM**: an integer specifying the medium for LET weighted scoring. If left unspecified, the default is the medium material that the bin is made of. Any other integer from the medium list specified in *mat.dat* is valid here. In that case, scoring will apply the stopping powers of the medium specified here. Transport will irrespective of the scoring always be done in the as medium specified in *geo.dat*. This is useful for calculating stopping power ratios needed by various cavity theories in radiation dosimetry.

4. unused.
5. unused.
6. unused.
3.5.9 VOXSCORE - Scoring within loaded CT-cube

VOXSCORE : Scoring within a voxelized structure, which is loaded with the VOX card in geo.dat. VOX functions similar to MSH, however the Cartesian mesh will always be aligned with the loaded CT-cube, automatically taking care of any translation and rotation of the CT-cube.

Arguments are:

1. unused.
2. unused.
3. unused.
4. JPART: particle type. See list in section A.3. Setting this value to -1 scores all particles. If heavy ions are scored (JPART=25) then a continuation card is required, where the first argument is an integer representing the particle charge Z, and the second argument is an integer representing the particle mass A.
5. DETECTOR: quantity that should be scored. Important: Zone scoring works straightforwardly only with the detectors ENERGY, CROSSFLU, DLET, COUNTER and PET as described in section 3.5.10. All other detectors can be applied as well, but require knowledge of the zone volume, which is not known by SHIELD-HIT12A (unlike the MSH and CYL scorer). Therefore the user must divide the scoring result with the corresponding zone volume (in cm³) for all other detectors than those listed above. This may be changed in a future release (see ticket #177).

6. outputfile: output file name.

Analogue to CYL and MSH, a continuation card is only required if JPART=25 or if DETECTOR is specified as LETFLU (and only LETFLU). Arguments for the continuation card are:

1. Z: an integer specifying the charge of the particle to be scored.
2. A: an integer specifying the number of nucleons of the particle to be scored. Setting this value to -1 scores all isotopes.
3. MEDIUM: an integer specifying the medium for LET weighted scoring. If left unspecified, the default is the medium material that the bin is made of. Any other integer from the medium list specified in mat.dat is valid here. In that case, scoring will apply the stopping powers of the medium specified here. Transport will irrespective of the scoring always be done in the as medium specified in geo.dat. This is useful for calculating stopping power ratios needed by various cavity theories in radiation dosimetry.

4. unused.
5. unused.
6. unused.
3.5.10 Available detectors for scoring

Different detector types are available for the auxiliary scoring. For each geometrical estimator invoked in the file `detect.dat` one type of detector has to be specified for DETECTOR in the second card. But not every geometrical estimator type can be combined with each of the available detectors. A detector type may itself invoke an additional card. The output is given in SHIELD-HIT12A standard units (see table A.1). The quantities that scale with particle number are normalized per simulated primary particle. The available detectors are:

**ENERGY**: Absolute energy deposited in each bin.

**FLUENCE**: Track length fluence \( \Phi \) in each bin.

**CROSSFLU**: Absolute number of particles entering each bin.

**FLU-CHAR**: Charged particle track length fluence \( \Phi_{charged} \) in each bin.

**FLU-NET**: Neutral particle track length fluence \( \Phi_{neutral} \) in each bin.

**DOSE**: Energy deposited in each bin divided by total mass (volume \( \times \) density) of the bin.

**LETFLU**: Sum of the products between fluence and unrestricted linear energy transfer (LET) over particle type \( i \),

\[
\overline{\Phi}_i \overset{\text{LET}}{=} \sum_i \phi_i(E, Z) \frac{dE}{dx}(E, Z)_i,
\]

for a medium specified by card 3, argument 3 which might be different from the material of the bin. This argument is a number \( m \) that corresponds to the \( m \)th material in `mat.dat`. The product for one specific particle \( i \) can be selected with arguments 1 and 2 in card 3. All particles are considered if these two arguments are left blank.

**DLET**: Dose (\( D \)) averaged unrestricted stopping power (LET) in MeV/cm. Since SHIELD-HIT12A 0.7.0 the scoring method used, is the algorithm “C” (equation 12) mentioned in [34]. Only charged particles are considered, since neutral particles have no electronic stopping power. For each particle \( i \):

\[
\overline{LET}_D = \frac{\sum_i D_i(E, Z) \frac{dE}{dx}(E, Z)_i}{\sum_i D_i(E, Z)}.
\]

**TLET**: Track length (\( \tau \)) - averaged unrestricted stopping power (LET) in MeV/cm. TLET Uses similar scoring method as for DLET, but for each charged particle \( i \):

\[
\overline{LET}_\tau = \frac{\sum_i \tau_i(E, Z) \frac{dE}{dx}(E, Z)_i}{\sum_i \tau_i(E, Z)}.
\]

which equals the fluence (\( \Phi \)) averaged unrestricted stopping power, since \( \Phi = \tau / V \).

**AVG-ENERGY**: Track length averaged energy for charged and neutral particles. For each particle \( i \):

\[
\overline{E}_\tau = \frac{\sum_i E_i(Z) \tau_i(E, Z)}{\sum_i \tau_i(E, Z)}.
\]

---

6Earlier versions of SHIELD-HIT12A employed the algorithm “A” from the same reference.
AVG-BETA: Track length averaged relativistic velocity $\beta = v/c$, applied to charged and neutral particles. For each particle $i$:

$$\bar{\beta}_r = \frac{\sum_i \beta_i(Z) \tau_i(E, Z)}{\sum_i \tau_i(E, Z)} \quad \ldots \quad (3.5)$$

Q: Armin’s Q-value. $Q$ is a radiation quality specifier similar to LET which is useful to express variations of observables such as the relative biological effectiveness (RBE) or relative effectiveness (RE or $\eta$) of detectors. Contrary to LET, the observables will be independent on particle charge, when expressed as a function of $Q$. The quantity is dose-averaged ($D$), and thus not defined in vacuum or for neutral particles. Only particles with a kinetic energy of 100 keV/nucleon or more will be scored. **Experimental implementation. $Q$ is not fully defined yet, and may change in future releases.**

$$Q_D = \frac{1}{\rho} \frac{1}{z_i^2} \frac{1}{E} \sum_i D_i(E, Z) \quad \ldots \quad (3.6)$$

where $z$ is the projectile charge, $\rho$ is the material density, and $E$ is the energy divided by the integer nucleon number (i.e. not divided by the atomic mass). $Q$ is per definition dimensionless.

DDD: Only available for CYL. See section 3.5.3. This detector is used for generating depth dose curve kernels for TRiP98.

SPC: Only available for DMSHZ and DCYLZ. See section 3.5.7

ALANINE: Response-equivalent dose for the alanine detector. The medium in which this detector is applied must be alanine (ICRU ID = 105) in order to make sense. For each charged particle $i$:

$$D_{\text{al}, \text{photon}} = \sum_i \eta(Z_i, E_i) D_{i, \text{alanine}, \text{ion}} \quad \ldots \quad (3.7)$$

The relative effectiveness $\eta$ is calculated using an implementation [35] of the Hansen and Olsen model [36,37] which only takes pions, kaons, and ions with $Z < 7$ into account. For heavier ions ($Z > 6$) $\eta$ is set to 1.0. For energies below 1 keV/nucleon $\eta$ is fixed to 0.3 for all ions.

COUNTER: A counter which counts the number of particles crossing a PLANE estimator.

### 3.5.11 Uncertainties

SHIELD-HIT12A does not record the sum of squares of the results in order to minimize the memory footprint of the program at runtime. Therefore the output from the detectors will not include any statement of the precision of the calculation.

Instead the user is encouraged to run the process several times in order to derive a best estimate of the precision. For $n$ independent runs (i.e. with different random seeds), but with equal number of primaries and each with the result $x_i$, the best estimate of precision $s_n$ can be obtained by

$$s_n = \frac{\sum (x_i - X_n)^2}{(n - 1)^{1/2}} \quad \ldots \quad (3.8)$$

where $X_n$ is the mean of all obtained $x_i$. Note that the underlying distributions may not be normal distributed, so the estimate can be misleading.
3.5.12 Examples - Scoring

Example 1: 1D and 2D maps

This example shows how to score a 1-D energy deposition of all particles in a cylindrical volume which is 20 cm long (from 35 cm to 55 cm). The output will be written into the file foo. In addition, a 2-D fluence map (1×20×300 bins) of carbon-12 ions (Z = 6 and A = 12) is generated and written to the file bar. Finally, the average energy deposited by all particles per primary particle in the zones 2 to 12 defined in geo.dat is outputted to foobarlongfilename.

Example 2: SPC file generation

The second example shows how to produce two .spc files in TRiP98 [33] format. The first file lin.spc contains the particle spectrum at different depths resolved by particle on a linear energy scale while for the second file, log.spc, a logarithmic energy scale is used.

Example 3: Angular histogram

Here is an example, how to record the angular histogram of particles crossing a plane. A X-Y plane is placed at Z = 10.0 cm. Differential angle scoring is invoked, scoring all from 0.0 to 5.0 mrad in 20 bins.

Example 4: DOSE and LET in CT-cubes, with particle tracks

This example shows how to score the Dose and Dose-averaged LET inside a CT-cube, which was loaded using the VOX card in geo.dat. A TRACE card is added, to visualize a few of the simulated transport events in the CT-cube.
Note, that for a 512x512x172 cube, this example allocates order of 4 GB of RAM for scoring. While this is normally not a problem in single-threaded mode, in parallel mode you may want to make sure that you have sufficient RAM available in order to prevent SHIELD-HIT12A to use swap space.
Chapter 4

Output files

SHIELD-HIT12A provides several output files, depending on the configuration in geo.dat and detect.dat.

Programs to convert the binary *.bdo files to human readable ASCII exist, where pymchelper is recommended to use.

See [https://pymchelper.readthedocs.io/](https://pymchelper.readthedocs.io/)

Finally a run-specific logfile is created shieldhit.log which will cover all details of the run.

4.1 Auxiliary output files

These files are not intended for further processing. They contain auxiliary information for the user in the manner of man-readable ASCII coding. There are no scripts for handling these files in an parallelized environment.

*for017*: This file is created and used by the geometry parser. If parsing of geo.dat failed you may find the file for017 which may contain additional diagnostics. Else this file can safely be deleted.

*shieldhit.log*: Contains input and output data of the current run. It is recommended to keep this file, if you need your run to be documented. Also found here are the stopping powers used as well as dependencies range-energy, macroscopic cross sections and other useful information for all particles/fragments and all materials in the task. Also found here are multigroup neutron cross sections below 14.5 MeV for all chemical elements used within this run.

*for028*: This is an optional file which can be requested by the **MAKELN** card in beam.dat. It contains a list of secondary neutrons below 14.5 MeV with all individual parameters: the birth point XYZ, direction of fly, kinetic energy. This file can be used as a neutron source for transportation with MCNP [21] like programs.
Chapter 5

Auxiliary programs

5.1 Helper scripts

5.2 Format conversion

5.2.1 convertmc - Convert .bdo files to ASCII or similar

`convertmc` converts the output of an estimator specified in `detect.dat` (see section 3.5). If run in a parallel environment, and multiple outputs exist of the same estimator, then `convertmc` will average over all files specified in the argument and estimate the error. `convertmc` is the most important script for postprocessing, and is not provided by the SHIELD-HIT12A installation, but is provided by the `pymchelper` package. The user is referred to [https://github.com/DataMedSci/pymchelper](https://github.com/DataMedSci/pymchelper) where this package is documented. For installation, make sure `pip` is installed (`sudo apt-get install pip` on Ubuntu/Debian systems) and run

```bash
$ pip install pymchelper
```

5.2.2 shield2fluka - Convert geo.dat to FLUKA format

`shield2fluka` converts the geometry file to a FLUKA formatted input file. Only the geometry file is converted, not the beam settings or scoring. The produced fluka formatted `output.inp` cannot be run in FLUKA, but it can be read by “SimpleGeo” which is a windows program for viewing geometries in 3D.


5.2.3 fluka2shield - Convert FLUKA .inp files to SHIELD-HIT12A input files

`fluka2shield` converts a FLUKA formatted input file to SHIELD-HIT12A input files. The resulting files are by no means complete, and must be manually edited by the user, however it may still serve as a help for converting more complex geometries.

`fluka2shield` is about to be obsolete, and the user is instead referred to `pymchelper` [https://github.com/DataMedSci/pymchelper](https://github.com/DataMedSci/pymchelper) which is more complete than `fluka2shield`. 
5.2.4 \textit{shield\textunderscore dEdx} - Prepare an external stopping power table

\textit{shield\textunderscore dEdx} generates a properly formatted stopping power table which can be read by SHIELD-HIT12A. \textit{shield\textunderscore dEdx} requires the \textit{libdedx} software package installed \cite{3,26}, which can be downloaded for free at \url{https://github.com/APTG/libdedx}.

The \textit{libdEdx} library features access to a range of alternative stopping power tables and a comprehensive list of target materials. An output files is generated according to the nomenclature found in table A.4. The generated ASCII file is human readable and contains a header with information on creation date, version number etc.

Example of usage:

\$ ./processing/dEdx/shield\textunderscore dEdx 1 223
Preparing stopping power table for PMMA using PSTAR + MSTAR.
Wrote Lucite.txt.
Chapter 6

Parallelization

SHIELD-HIT12A can run in a parallelized environment in several ways, ranging from single PC which features multiple cores, or massive computer clusters. SHIELD-HIT12A runs on CPUs only, currently no GPU code is implemented. In this chapter several ways of parallelizing SHIELD-HIT12A is explained. SHIELD-HIT12A will here run in “embarrassingly parallel” mode, where simple independent jobs are submitted to various cores.

Each independent job needs it’s own random number seed. The command line option -N can be used to add an offset in form of an integer value to the RNDSEED card specified in the beam.dat file. If RNDSEED was not specified in the beam.dat the offset is simply added to the default seed.

In other words, assuming a project is setup in the example/ directory, the command

```
$ shieldhit example/ -N17
```

will add 17 to the RNDSEED value specified in example/beam.dat and run SHIELD-HIT12A with this new seed.

All output results example/*.bdo will carry a four digit zero-padded string with the given offset. That is, following the aforementioned example, in the form of example/*0017.bdo. Beware that the offset specified by the -N option is limited to 9999 and negative values are not accepted.

Example scripts for parallelization are only shipped with the Linux distribution. These are rtshield.sh for Torque systems, rbshield.sh for Platform LSF based systems and rcshield.py for Condor based systems.

### 6.1 Torque

Torque is a widely used system for job submission across multiple nodes. Here a very simple tutorial is given for using multiple nodes on a PC running SHIELD-HIT12A processes. It is assumed that Torque and the PBS scheduler is running and correctly configured. Furthermore it is assumed that SHIELD-HIT12A was fully installed.

A simple PBS submission script rtshield.sh can look like this:

```
#!/bin/sh
#
PBS -N SHIELDHIT_JOB
PBS -M bassler@phys.au.dk
#
shieldhit $1 -N$PBS_ARRAYID
```

53
Simply create this file in the directory you wish to run. Step into the same directory and run:

```
$ qsub -V -t 0-9 -d . rtshield.sh
```

The `-V` forwards the local environment variables to the submission script. This way we do not need to specify the path of the `shieldhit` command, since it is referred to by the `PATH` environment variable.

The `-t` option specifies the nodes to be run, in this case from 0 to 9, meaning 10 instances in total. The output files will be suffixed accordingly, so they will not overwrite each other.

Finally, the `-d .` option specifies the working directory. `qsub` expects to find all the input files as well as the `rtshield.sh` script here.

You can check the status of your job submission with `qstat`. A good manual on Torque can be found on [http://www.adaptivecomputing.com/support/documentation-index/torque-resource-manager](http://www.adaptivecomputing.com/support/documentation-index/torque-resource-manager).

After the run completes, each output file (which can be merged by the `convertmc` script) will be appended with an integer number corresponding to the instance number. Thus in the example above, the output files will be suffixed with integers ranging from 0000 to 0009.

### 6.2 LSF

LSF system is quite similar to torque. You will have to adapt the example script `rbshield.sh` to your local environment, and then do the submission by e.g:

```
$ bsub -q test -J jobname[1-2] < ./rbshield.sh
```

which will run two jobs in the `test` queue.

### 6.3 Condor

Another job submission and scheduling system is CONDOR[1]. Here submission can be handled by the `rcshield.py` script.

```
$ rcshield.py -c -M100 examples/simple
```

will run the job found in the `examples/simple` directory 100 times. All results, debugging output is transferred back to the directory, and each result file is suffixed with a number in order to distinguish the results. `rcshield.py` accepts a range of options:

- `-h` - print all options with a short description.

- `-f` - should be followed by a comma separated (without space) file list, which will be transmitted along with the usual mandatory files. The usual input files which always will be transmitted are: `mat.dat geo.dat beam.dat Air.txt Water.txt detect.dat`. Additional files could be a ripple filter file `rifi.dat`, source or a files which overrides the default model parameters (see section 3.3.3).

- `-M` - number of simulations to be performed. Random seed is automatically changed for each run.

- `-N` - first run will have the number followed by this option. Useful if you want to add more statistics to a previous batch or runs. If the last completed run had number 99, you should set `-M100`.

-a - after each batch of runs, rcfluka.py will wait 30 seconds, for any file transfers to complete. This grace time is necessary if you run any postprocessing scripts or the -c option. For small runs and quick testing, the 30 secs may be annoying, and can therefore be deactivated with the -a option.

-n - the jobs are submitted as nice jobs to the cluster, backfilling it if no other tasks are pending. Use this option if you are not in a hurry to get your data.

-c - clean up some of the condor files which were generated, which are only of interest if something goes wrong. Purpose of this option is to keep your run directory tidy.

-C - big clean: use this if you want to start from scratch in your directory. Nothing is submitted to the cluster, but the directory is emptied for any data generated by previous runs. DANGEROUS, files which may resemble generated data are deleted as well. Basically anything which matches rcshield*, rcshield*[0-9][0-9][0-9][0-9], fort[0-9][0-9][0-9] and fort.24 are removed.

-t - test script, but do not submit jobs to condor cluster.

-S - use standard universe instead of vanilla (see condor manual). Standard universe will only work if you have linked in the appropriate condor libraries for checkpointing into the SHIELD-HIT12A executable.

-o - followed by a string which will be added to the condor submit file.

-i - followed by a filename of a file which will be appended to the condor submit script.

-m - followed by an email address, which will be notified when all jobs are completed (or failed).

-x - single submission mode. This overrides -M and -m option. Simply one job is submitted, and rcshield.py immediately exits after submission. This option is useful if you just want the raw output from a single run, and dont want the rcshield.py script block the terminal while running. Entire control is within condor, which will transfer files upon job completion.
Appendix A

User’s reference tables and lists

A.1 Command line options and arguments

SYNOPSIS

```
shieldhit [OPTIONS]... [WORKDIR]
```

DESCRIPTION

The `shieldhit` command takes exactly one optional argument, and that is the working directory. All input files are expected to be placed in the working directory, and all output files will be placed in the same directory. Additional options may be given, as described below.

OPTIONS

- `-h`, `--help`: Print all available options and exit.
- `-V`, `--version`: Print version and exit.
- `-v`, `--verbose`: Be verbose.
- `-n`, `--nstat=NUM`: Override NSTAT using NUM amount of particles instead. Useful for testing with short runs or dry runs, `-n0`.
- `-t`, `--time="HH:MM:SS"`: Set maximum total simulation time in HH:MM:SS (including initialization).
- `-N`, `--seedoffset=NUM`: Offset added to RNDSEED specified in `beam.dat`. Useful for parallelization, as described in chapter 6.
- `-b`, `--beamfile=FILE`: Load FILE instead of WORKDIR/beam.dat.
- `-g`, `--geofile=FILE`: Load FILE instead of WORKDIR/geo.dat.
- `-m`, `--matfile=FILE`: Load FILE instead of WORKDIR/mat.dat.
- `-d`, `--detectfile=FILE`: Load FILE instead of WORKDIR/detect.dat.

SIGNALS

- `SIGUSR1`: (Linux only) Saves current results.
- `SIGINT`: Saves current results and exits.
A.2 Units

Table A.1 provides an overview of the standard units in SHIELD-HIT12A. They apply at any time unless explicitly stated otherwise.

Table A.1: Table of default units in SHIELD-HIT12A.

<table>
<thead>
<tr>
<th>Value</th>
<th>Default unit</th>
</tr>
</thead>
<tbody>
<tr>
<td>Distance; position</td>
<td>cm</td>
</tr>
<tr>
<td>Track length density; fluence</td>
<td>cm⁻²</td>
</tr>
<tr>
<td>Volume</td>
<td>cm³</td>
</tr>
<tr>
<td>Density</td>
<td>g/cm³</td>
</tr>
<tr>
<td>Atomic density</td>
<td>atoms/cm³</td>
</tr>
<tr>
<td>Projectile energy (ions)</td>
<td>MeV/nucleon</td>
</tr>
<tr>
<td>Projectile energy (non-ions)</td>
<td>MeV</td>
</tr>
<tr>
<td>Deposited energy</td>
<td>MeV</td>
</tr>
<tr>
<td>Dose</td>
<td>MeV/g</td>
</tr>
<tr>
<td>Stopping power; LET</td>
<td>MeV/cm</td>
</tr>
<tr>
<td>Mass stopping power</td>
<td>MeV cm²/g</td>
</tr>
<tr>
<td>Mean excitation potential</td>
<td>eV</td>
</tr>
</tbody>
</table>
A.3 Particle codes

Table A.2: List of available particle identifiers $\text{JPART}$ and the corresponding particle names.

<table>
<thead>
<tr>
<th>$\text{JPART}$</th>
<th>Particle</th>
<th>$\text{JPART}$</th>
<th>Particle</th>
</tr>
</thead>
<tbody>
<tr>
<td>-1</td>
<td>All particles</td>
<td>13</td>
<td>Electron</td>
</tr>
<tr>
<td>1</td>
<td>Neutron</td>
<td>14</td>
<td>Positron</td>
</tr>
<tr>
<td>2</td>
<td>Proton</td>
<td>15</td>
<td>Muon $\mu^-$</td>
</tr>
<tr>
<td>3</td>
<td>Pion $\pi^-$</td>
<td>16</td>
<td>Muon $\mu^+$</td>
</tr>
<tr>
<td>4</td>
<td>Pion $\pi^+$</td>
<td>17</td>
<td>$e^-$-Neutrino $\nu_e$</td>
</tr>
<tr>
<td>5</td>
<td>Pion $\pi^0$</td>
<td>18</td>
<td>$e^-$-Anti-neutrino $\bar{\nu}_e$</td>
</tr>
<tr>
<td>6</td>
<td>Anti-neutron</td>
<td>19</td>
<td>$\mu^-$-Neutrino $\nu_\mu$</td>
</tr>
<tr>
<td>7</td>
<td>Anti-proton</td>
<td>20</td>
<td>$\mu^-$-Anti-neutrino $\bar{\nu}_\mu$</td>
</tr>
<tr>
<td>8</td>
<td>Kaon $\kappa^-$</td>
<td>21</td>
<td>Deuteron</td>
</tr>
<tr>
<td>9</td>
<td>Kaon $\kappa^+$</td>
<td>22</td>
<td>Triton</td>
</tr>
<tr>
<td>10</td>
<td>Kaon $\kappa^0$</td>
<td>23</td>
<td>He-3</td>
</tr>
<tr>
<td>11</td>
<td>Kaon $\kappa^-$</td>
<td>24</td>
<td>He-4</td>
</tr>
<tr>
<td>12</td>
<td>Gamma ray</td>
<td>25</td>
<td>Heavy ions</td>
</tr>
</tbody>
</table>

Note, the generalized particle $\text{JPART}=-1$ is only available within a scoring estimator, and not as a particle source. For a heavy ion $\text{JPART}=25$ the charge $Z$ and atomic number $A$ have usually to be defined additionally. As particle source, $Z > 2$ is expected.
### A.4 Nuclear targets with neutron cross sections

Table A.3 shows a list of elements and isotopes for which neutron cross sections are included in SHIELD-HIT12A. Natural isotope mixtures are marked with a "*".

Table A.3: List of nuclear target identifiers with available neutron cross sections \texttt{NUCLID} and the corresponding names used by SHIELD-HIT12A.

<table>
<thead>
<tr>
<th>Z</th>
<th>NUCLID</th>
<th>Isotope</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>H-1 - Hydrogen</td>
<td>JEFF-3.1</td>
</tr>
<tr>
<td>1</td>
<td>101</td>
<td>H-2 - Deuterium</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>1</td>
<td>102</td>
<td>H-3 - Tritium</td>
<td>BROND</td>
</tr>
<tr>
<td>2</td>
<td>104</td>
<td>He-3</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>He-4</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>3</td>
<td>105</td>
<td>Li-6</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>3</td>
<td>3</td>
<td>Li-7</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>4</td>
<td>4</td>
<td>Be-9</td>
<td>ENDFB7.1</td>
</tr>
<tr>
<td>5</td>
<td>106</td>
<td>B-10</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>5</td>
<td>5</td>
<td>B-11</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>6</td>
<td>6</td>
<td>C-*=</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>7</td>
<td>7</td>
<td>N-*=</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>8</td>
<td>8</td>
<td>O-*=</td>
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<tr>
<td>9</td>
<td>9</td>
<td>F-19</td>
<td>BROND-2</td>
</tr>
<tr>
<td>11</td>
<td>11</td>
<td>Na-23</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>12</td>
<td>12</td>
<td>Mg-*=</td>
<td>CENDL-2</td>
</tr>
<tr>
<td>13</td>
<td>13</td>
<td>Al-27</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>14</td>
<td>14</td>
<td>Si-*=</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>15</td>
<td>15</td>
<td>P-31</td>
<td>BROND-2</td>
</tr>
<tr>
<td>16</td>
<td>16</td>
<td>S-*=</td>
<td>JEFF-3</td>
</tr>
<tr>
<td>17</td>
<td>17</td>
<td>Cl-*=</td>
<td>BROND-2</td>
</tr>
<tr>
<td>18</td>
<td>18</td>
<td>Ar-*=</td>
<td>JENDL40</td>
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<td>19</td>
<td>19</td>
<td>K-*=</td>
<td>CENDL-2</td>
</tr>
<tr>
<td>20</td>
<td>20</td>
<td>Ca-*=</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>22</td>
<td>22</td>
<td>Ti-*=</td>
<td>JEFF-3</td>
</tr>
<tr>
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<td>23</td>
<td>V-51</td>
<td>CENDL-2</td>
</tr>
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<td>24</td>
<td>Cr-*=</td>
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<tr>
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<td>25</td>
<td>Mn-55</td>
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<td>26</td>
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<td>27</td>
<td>27</td>
<td>Co-59</td>
<td>ANL/NDM</td>
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<tr>
<td>28</td>
<td>28</td>
<td>Ni-*=</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>29</td>
<td>29</td>
<td>Cu-*=</td>
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</tr>
<tr>
<td>30</td>
<td>30</td>
<td>Zn-*=</td>
<td>BROND-2</td>
</tr>
<tr>
<td>31</td>
<td>31</td>
<td>Ga-*=</td>
<td>JENDL</td>
</tr>
<tr>
<td>32</td>
<td>32</td>
<td>Ge-*=</td>
<td>JEFF-3.1</td>
</tr>
<tr>
<td>33</td>
<td>33</td>
<td>As-75</td>
<td>JENDL</td>
</tr>
<tr>
<td>41</td>
<td>41</td>
<td>Nb-93</td>
<td>CENDL-2</td>
</tr>
</tbody>
</table>

Continued on Next Page...
<table>
<thead>
<tr>
<th>$Z$</th>
<th>NUCLID</th>
<th>Isotope</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
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<td>42</td>
<td>42</td>
<td>Mo-*</td>
<td>JENDL</td>
</tr>
<tr>
<td>47</td>
<td>47</td>
<td>Ag-*</td>
<td>JENDL</td>
</tr>
<tr>
<td>48</td>
<td>48</td>
<td>Cd-*</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>50</td>
<td>50</td>
<td>Sn-*</td>
<td>CENDL-2</td>
</tr>
<tr>
<td>63</td>
<td>63</td>
<td>Eu-*</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>64</td>
<td>64</td>
<td>Gd-*</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>68</td>
<td>68</td>
<td>Er-*</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>73</td>
<td>73</td>
<td>Ta-181</td>
<td>JEFF</td>
</tr>
<tr>
<td>74</td>
<td>74</td>
<td>W-*</td>
<td>CENDL31</td>
</tr>
<tr>
<td>75</td>
<td>75</td>
<td>Re-*</td>
<td>JEFF-3.1</td>
</tr>
<tr>
<td>79</td>
<td>79</td>
<td>Au-187</td>
<td>BROND-2</td>
</tr>
<tr>
<td>80</td>
<td>80</td>
<td>Hg-*</td>
<td>JEFF</td>
</tr>
<tr>
<td>82</td>
<td>82</td>
<td>Pb-*</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>83</td>
<td>83</td>
<td>Bi-209</td>
<td>ENDFB7.1</td>
</tr>
<tr>
<td>90</td>
<td>90</td>
<td>Th-232</td>
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<td>U-238</td>
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<td>94</td>
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<td>Pu-239</td>
<td>BNAB-81</td>
</tr>
<tr>
<td>94</td>
<td>107</td>
<td>Pu-239</td>
<td>BNAB-81</td>
</tr>
</tbody>
</table>
### A.5 Material ICRU_ID

Table A.4 lists all material numbers as defined by ICRU which are known to SHIELD-HIT12A. The ICRU ID number can optionally be specified in the material input file *mat.dat* which then will trigger SHIELD-HIT12A to load stopping power data from an external ASCII file (see section [3.2](#)). The file containing the stopping power data must exactly be named as the according file name in the table below.

#### Table A.4: List of ICRU ID and file names used by SHIELD-HIT12A.

<table>
<thead>
<tr>
<th>ICRU_ID</th>
<th>File name</th>
<th>Material name and remarks</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>H.txt</td>
<td>HYDROGEN</td>
</tr>
<tr>
<td>2</td>
<td>He.txt</td>
<td>HELIUM</td>
</tr>
<tr>
<td>3</td>
<td>Li.txt</td>
<td>LITHIUM</td>
</tr>
<tr>
<td>4</td>
<td>Be.txt</td>
<td>BERYLLIUM</td>
</tr>
<tr>
<td>5</td>
<td>B.txt</td>
<td>BORON</td>
</tr>
<tr>
<td>6</td>
<td>C.txt</td>
<td>AMORPHOUS CARBON (density 2.0 g/cm³)</td>
</tr>
<tr>
<td>906</td>
<td>Graphtie.txt</td>
<td>GRAPHITE</td>
</tr>
<tr>
<td>7</td>
<td>N.txt</td>
<td>NITROGEN</td>
</tr>
<tr>
<td>8</td>
<td>O.txt</td>
<td>OXYGEN</td>
</tr>
<tr>
<td>(...)</td>
<td>(...)</td>
<td>(...)</td>
</tr>
<tr>
<td>98</td>
<td>Cf.txt</td>
<td>CALIFORNIA</td>
</tr>
<tr>
<td>99</td>
<td>A-150.txt</td>
<td>A-150 TISSUE-EQUIVALENT PLASTIC</td>
</tr>
<tr>
<td>100</td>
<td>Acetone.txt</td>
<td>ACETONE</td>
</tr>
<tr>
<td>101</td>
<td>Acetylene.txt</td>
<td>ACETYLENE</td>
</tr>
<tr>
<td>102</td>
<td>Adenine.txt</td>
<td>ADENINE</td>
</tr>
<tr>
<td>103</td>
<td>Adipose.txt</td>
<td>ADIPOSE TISSUE (ICRP)</td>
</tr>
<tr>
<td>104</td>
<td>Air.txt</td>
<td>AIR, DRY (NEAR SEA LEVEL)</td>
</tr>
<tr>
<td>105</td>
<td>Alanine.txt</td>
<td>ALANINE</td>
</tr>
<tr>
<td>106</td>
<td>Al2O3.txt</td>
<td>ALUMINUM OXIDE</td>
</tr>
<tr>
<td>107</td>
<td>Amber.txt</td>
<td>AMBER</td>
</tr>
<tr>
<td>108</td>
<td>Ammonia.txt</td>
<td>AMMONIA</td>
</tr>
<tr>
<td>109</td>
<td>Aniline.txt</td>
<td>ANILINE</td>
</tr>
<tr>
<td>110</td>
<td>Anthracene.txt</td>
<td>ANTHRACENE</td>
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<tr>
<td>111</td>
<td>B-100.txt</td>
<td>B100</td>
</tr>
<tr>
<td>112</td>
<td>Bakelite.txt</td>
<td>BAKELITE</td>
</tr>
<tr>
<td>113</td>
<td>BaF2.txt</td>
<td>BARIUM FLUORIDE</td>
</tr>
<tr>
<td>114</td>
<td>BaSO4.txt</td>
<td>BARIUM SULFATE</td>
</tr>
<tr>
<td>115</td>
<td>Benzene.txt</td>
<td>BENZENE</td>
</tr>
<tr>
<td>116</td>
<td>BeO.txt</td>
<td>BERYLLIUM OXIDE</td>
</tr>
<tr>
<td>117</td>
<td>BiGeO.txt</td>
<td>BISMUTH GERMANIUM OXIDE</td>
</tr>
<tr>
<td>118</td>
<td>BloodICRP.txt</td>
<td>BLOOD (ICRP)</td>
</tr>
<tr>
<td>119</td>
<td>BoneICRU.txt</td>
<td>BONE, COMPACT (ICRU)</td>
</tr>
<tr>
<td>120</td>
<td>BoneICRP.txt</td>
<td>BONE, CORTICAL (ICRP)</td>
</tr>
<tr>
<td>121</td>
<td>B4C.txt</td>
<td>BORON CARBIDE</td>
</tr>
<tr>
<td>122</td>
<td>BoronOxide.txt</td>
<td>BORON OXIDE</td>
</tr>
<tr>
<td>123</td>
<td>BrainICRP.txt</td>
<td>BRAIN (ICRP)</td>
</tr>
<tr>
<td>124</td>
<td>Butane.txt</td>
<td>BUTANE</td>
</tr>
<tr>
<td>125</td>
<td>N-ButylAlcohol.txt</td>
<td>N-BUTYLALCOHOL</td>
</tr>
</tbody>
</table>

Continued on Next Page...

1 All elements up to Californium (Z = 98) are supported following this naming scheme.
<table>
<thead>
<tr>
<th>ICRU_ID</th>
<th>File name</th>
<th>Material name and remarks</th>
</tr>
</thead>
<tbody>
<tr>
<td>126</td>
<td>C-552.txt</td>
<td>C-552 AIR-EQUIVALENT PLASTIC</td>
</tr>
<tr>
<td>127</td>
<td>CdTe.txt</td>
<td>CADMIUM TELLURIUM</td>
</tr>
<tr>
<td>128</td>
<td>CdWO4.txt</td>
<td>CADMIUM TUNGSTATE</td>
</tr>
<tr>
<td>129</td>
<td>CaCO3.txt</td>
<td>CALCIUM CARBONATE</td>
</tr>
<tr>
<td>130</td>
<td>CaF2.txt</td>
<td>CALCIUM FLUORIDE</td>
</tr>
<tr>
<td>131</td>
<td>CaO.txt</td>
<td>CALCIUM OXIDE</td>
</tr>
<tr>
<td>132</td>
<td>CaSO4.txt</td>
<td>CALCIUM SULFATE</td>
</tr>
<tr>
<td>133</td>
<td>CaWO4.txt</td>
<td>CALCIUM TUNGSTATE</td>
</tr>
<tr>
<td>134</td>
<td>CO2.txt</td>
<td>CARBON DIOXIDE</td>
</tr>
<tr>
<td>135</td>
<td>CCl4.txt</td>
<td>CARBON TETRACHLORIDE</td>
</tr>
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<td>136</td>
<td>Cellophane.txt</td>
<td>CELLULOSE ACETATE, CELLOPHANE</td>
</tr>
<tr>
<td>137</td>
<td>CAB.txt</td>
<td>CELLULOSE ACETATE BUTYRATE</td>
</tr>
<tr>
<td>138</td>
<td>CelNitrates.txt</td>
<td>CELLULOSE NITRATE</td>
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<tr>
<td>139</td>
<td>CeSO4Dos.txt</td>
<td>CERIC SULFATE DOSIMETER SOLUTION</td>
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<td>140</td>
<td>CsF.txt</td>
<td>CESIUM FLUORIDE</td>
</tr>
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<td>141</td>
<td>CsI.txt</td>
<td>CESIUM IODIDE</td>
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<td>ClBenzene.txt</td>
<td>CHLOROBENZENE</td>
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<td>143</td>
<td>CHCl3.txt</td>
<td>CHLOROFORM</td>
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<td>144</td>
<td>Concrete.txt</td>
<td>CONCRETE PORTLAND</td>
</tr>
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<td>Cyclohexane.txt</td>
<td>CYCLOHEXANE</td>
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<tr>
<td>146</td>
<td>DCIBenzene.txt</td>
<td>1,2-DICHLOROBENZENE</td>
</tr>
<tr>
<td>147</td>
<td>DCIDEtEth.txt</td>
<td>DICHLORODIETHYL ETHER</td>
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<td>DCIEthane.txt</td>
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<td>DEtyEther.txt</td>
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<td>DMF.txt</td>
<td>N,N-DIMETHYL FORMAMIDE</td>
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<td>DIMETHYL SULFOXIDE</td>
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<td>EthylCell.txt</td>
<td>ETHYL CELLULOSE</td>
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<td>C2H4.txt</td>
<td>ETHYLENE</td>
</tr>
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<td>Eyelens.txt</td>
<td>EYELENS (ICRP)</td>
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<td>Fe2O3.txt</td>
<td>FERRIC OXIDE</td>
</tr>
<tr>
<td>158</td>
<td>FeB.txt</td>
<td>FERRO BORIDE</td>
</tr>
<tr>
<td>159</td>
<td>FeO.txt</td>
<td>FERROUS OXIDE</td>
</tr>
<tr>
<td>160</td>
<td>FeSO4Dos.txt</td>
<td>FERROUS SULFATE DOSIMETER SOLUTION</td>
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Table A.4 – Continued

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Table A.4 – Continued

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<td>278</td>
<td>Xylene.txt</td>
<td>XYLENE</td>
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A.6 Bodies

Each body is described with an identifier, that is a three-letter code preceded by two empty spaces. Another integer assigns a number to the body in the next 5 columns. Then 6 blocks each 10 columns wide specify the arguments $P_1$ to $P_6$. If additional arguments are required, a second card is added holding the arguments $P_7$ to $P_{12}$, but leaving the first 10 columns empty. Even further cards follow the format of the second card. The arguments $P_1$ can specify a point, a vector or a radius, depending on the body described. The bodies available in SHIELD-HIT12A are listed below:

**SPH**: A sphere defined by the center coordinates $(P_1,P_2,P_3)$ and the radius $P_4$. The remaining arguments $P_5$ and $P_6$ are ignored.

**WED**: A wedge spanned by three vectors. The origin is the point $(P_1,P_2,P_3)$ from where the three vectors $(P_4,P_5,P_6)$, $(P_7,P_8,P_9)$ and $(P_{10},P_{11},P_{12})$ start.

**ARB**: Arbitrary convex polyhedron. This body is described by 8 points, i.e. 24 arguments. The points must be specified in the order as indicated by the figure to the left: Point 1 is $(P_1,P_2,P_3)$, point 2 $(P_4,P_5,P_6)$ … point 8 $(P_{22},P_{23},P_{24})$.

**RPP**: Rectangular parallelepiped. $(P_1,P_2)$ marks minimum and maximum X coordinates. $(P_3,P_4)$ is min and max values for Y, and $(P_5,P_6)$ for Z.
BOX : A box spanned by 3 vectors. All vectors start in the point \((P_1, P_2, P_3)\) and point to the points \((P_4, P_5, P_6)\), \((P_7, P_8, P_9)\) and \((P_{10}, P_{11}, P_{12})\). They must be orthogonal on each other.

RCC : Right circular cylinder. One end of the cylinder is described by a circle with the center at \((P_1, P_2, P_3)\). The arguments \((P_4, P_5, P_6)\) hold a vector measured from the center to the opposite end of the cylinder. \(P_7\) marks the radius of the cylinder.

REC : Right elliptical cylinder. Similar to RCC, but instead of a radius, two additional vectors \((P_7, P_8, P_9)\) and \((P_{10}, P_{11}, P_{12})\) describe the minor and major axis. All three vectors must be orthogonal to each other.

TRC : A cone spanned by two circles. First circle is centered at point \((P_1, P_2, P_3)\) and has radius \(P_7\). The other end of the cone is described by the circle centered at the end of a vector \((P_4, P_5, P_6)\) starting from the center of the first circle. The second circle has a radius of \(P_8\). The radius in \(P_7\) must be larger than the radius in \(P_8\).

ELL : Ellipsoid. Centered at point \((P_1, P_2, P_3)\), three orthogonal vectors \((P_4, P_5, P_6)\), \((P_7, P_8, P_9)\) and \((P_{10}, P_{11}, P_{12})\) span the body. This body is differently defined than in FLUKA.

VOX : Prepares a voxelized structure in shape of a box. \((P_1, P_2, P_3)\) marks the SHIELD-HIT12A universe \((0, 0, 0)\) position inside the CT-cube. The \((P_1, P_2, P_3)\) coordinates are therefore in terms of the CT-cube universe, and all are in [cm]. \(P_4, P_5\) are the couch and gantry angles, respectively. A continuation card is needed for specifying the filename to be loaded. The filename must be formatted in VOXELPLAN format, used also by the treatment planning system TRiP98. SHIELD-HIT12A will read both the header and the Hounsfield unit data cube, having the \\".hed" \\".ctx" suffixes, respectively. Therefore, no suffix must be specified when loading the cube. See also section 3.4.3.

MOV : Copies a body number \(P_1\) to a new position translated by the vector \((P_2, P_3, P_4)\). The columns 6-10 in the MOV card must hold an integer specifying a new and unique body number, just like a normal body specification.
### A.7 Input and output files.

Table A.5 provides an overview of input (I) and output (O) files used by SHIELD-HIT12A at run time. They have either ASCII (A) or binary (B) format. Some files are optional.

<table>
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<tr>
<th>Filename</th>
<th>I/O</th>
<th>ASCII/Bin</th>
<th>Description</th>
<th>Remarks</th>
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<td>beam.dat</td>
<td>I</td>
<td>A</td>
<td>Seed, projectile, stats, etc.</td>
<td></td>
</tr>
<tr>
<td>detect.dat</td>
<td>I</td>
<td>A</td>
<td>Auxiliary scoring</td>
<td></td>
</tr>
<tr>
<td>geo.dat</td>
<td>I</td>
<td>A</td>
<td>Geometry description</td>
<td></td>
</tr>
<tr>
<td>mat.dat</td>
<td>I</td>
<td>A</td>
<td>Target medium chemical composition</td>
<td></td>
</tr>
<tr>
<td>*.bdo</td>
<td>O</td>
<td>B</td>
<td>Result of scoring</td>
<td>1), 2)</td>
</tr>
<tr>
<td>shieldhit.log</td>
<td>O</td>
<td>A</td>
<td>Logfile</td>
<td></td>
</tr>
<tr>
<td>for017</td>
<td>O</td>
<td>A</td>
<td>GEMCA parser log</td>
<td></td>
</tr>
<tr>
<td>for028</td>
<td>O</td>
<td>A</td>
<td>Secondary neutrons production&lt; 14.5 MeV enabled with MAKLN card</td>
<td></td>
</tr>
<tr>
<td>parlev.dat</td>
<td>I</td>
<td>A</td>
<td>External parameter file for PARLEV parameters</td>
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<tr>
<td>rifi.dat</td>
<td>I</td>
<td>A</td>
<td>Description of the ripple filter</td>
<td>1), 3)</td>
</tr>
<tr>
<td>sobp.dat</td>
<td>I</td>
<td>A</td>
<td>External particle source file</td>
<td>1), 3)</td>
</tr>
</tbody>
</table>

1) Optional.
2) File name is not predefined but set by user in `detect.dat`.
3) File name is not predefined but set by user in `beam.dat`.
The PARLEV parameters listed in table A.6 can be used for tuning the nuclear models in SHIELD-HIT. Up to 40 numeric parameters are set in SHIELD-HIT12A which may affect the calculated results. The default values are carefully selected and benchmarked against experimental data, and should therefore not be changed. However, for research purposes they can be changed using an external input file as described in section 3.3.3. In this case, a message is displayed for all modified settings in the output file for024.

Values which are marked “for high energies” in the table below, are only used at high energies (well above 1 GeV/A) and will therefore not influence calculations in normal particle radiotherapy settings.

A short description of the individual PARLEV parameters is given below. Details on the individual nuclear fragmentation models are given in the references [13–15, 17–19, 38–41].

**PARLEV(1) – PIDABS** : The probability of absorption of pion in nucleus by the quasi-deuteron pair.

**PARLEV(2) – TLIMIT** : Time of simulation of the cascade in nucleus (does not affect the strong decays of the resonances).

**PARLEV(3) – MMES** : The time of formation of mesons in their rest frame: \( \tau = \frac{1}{5.06 \cdot \text{MMES}} \) [fm/c].

**PARLEV(4) – MBAR** : The time of formation of baryons in their rest frame: \( \tau = \frac{1}{5.06 \cdot \text{MBAR}} \) [fm/c].

**PARLEV(5) – EPS1, EPS2** : Separation energy of a nucleon from the nucleus to the cascade stage of the nuclear reaction. The energy is the same for the target nucleus (EPS2) and the projectile (EPS1).

**PARLEV(6) – VP** : The depth of the potential well for pion inside the nucleus.

**PARLEV(7) – C1, C2** : The parameter of diffuseness of the Saxon-Woods distribution of nuclear density \( \rho_{WS}(r) = \rho_0 / [1 + \exp(r - R)/c] \), \( R = r_0 A^{1/3} \). The parameter is the same for the target nucleus (C2) and the projectile (C1).

**PARLEV(8) – D1, D2** : The parameter that determines the maximal radius \( R_{\text{max}} \) of the distribution of nuclear density \( \rho(r) \) from the condition \( \rho(0) / \rho(0) = D \). This parameter is used for both the Saxon-Woods distribution \( \rho_{WS}(r) \) and for light nuclei: \( \rho(r) = \rho_0 \exp(-ar^2) \). The parameter is the same for the target nucleus (D2) and the projectile (D1).

**PARLEV(9) – R0N1, R0N2** : The parameter \( r_0 \), which determines the radius of the nucleus by the formula \( R = r_0 A^{1/3} \). For light nuclei this parameter directly sets the radius of nucleus. The parameter is the same for the target nucleus (R0N2) and the projectile (R0N1). PARLEV(9) is applicable to all isotopes, except for those specified in PARLEV(10) – PARLEV(21).

**PARLEV(10) – PARLEV(21)** : Individual values of R0N1, R0N2 for light nuclei. SHIELD-HIT12A does not discriminate between R0N1 and R0N2. In some cases groups of isotopes are affected. For instance, PARLEV(20) affects all nitrogen isotopes with nucleon number of 10 or less. Any isotope which are not affected by PARLEV(10) – PARLEV(21) are treated with PARLEV(9).
PARLEV(22) – PRECOM: This parameter determines the contribution of pre-equilibrium emission of the lightest nuclei during transition from the cascade stage of nuclear reactions toward the equilibrium de-excitation (0 ≤ PRECOM ≤ 1).

PARLEV(23) – ICAN: This parameter determines the number of evaporation channels at equilibrium deexcitation. If ICAN = 0 then only 6 channels are evaporated (n,p,d,t,He^3,He^4+fission). This mode should be used when mainly dealing with Fermi-Breakup of lighter nuclei. If ICAN = 1 then there are 32 channels (+ fission). The latter should only be used when working with heavy excited nuclei.

PARLEV(24) – IMA: Maximal number of fragments at multifragmentation of nuclei with excitation below 3 MeV/nucleon.

PARLEV(25) – EPSIL0: Inverse level density parameter for internal bulk excitation of fragments in multifragmentation.

PARLEV(26) – FKACOL: Freeze-out volume for Coulomb interaction of fragments in multifragmentation.

PARLEV(27) – RNCL: The parameter r_0, which determines the radius of the excited nucleus by the formula R = r_0 A^{1/3} at the stage of equilibrium de-excitation.

PARLEV(28) – AMS: Density level parameter of the excited nucleus at evaporation.

PARLEV(29) – AMFS: Density level parameter of the excited nucleus at fission. Changing this parameter has no effect, as it is overridden internally.

PARLEV(30): The border for the Fermi break-up on the number of nucleons: for heavier nuclei the multifragmentation/evaporation/fission model is used.

PARLEV(31) – ILEVRA: The border for the multifragmentation on the excitation energy per nucleon: for more low excitations only the evaporation/fission model is used. (do not set higher than 2 MeV/nucleon).

PARLEV(32) – ILEVRA: Products of Fermi break-up are produced only in the ground state (ILEVRA=1), or these products can be formed in an excited state (ILEVRA=2).


PARLEV(35) – PARLEV(38): Reserved.

PARLEV(39) – SIGION: The parameter for renormalization the total \( \sigma_{\text{tot}}(E) \) and inelastic \( \sigma_{\text{in}}(E) \) cross sections of nucleus-nucleus interactions in the cascade inside a macroscopic target.

PARLEV(40) – MICROD: The parameter for renormalization the total \( \sigma_{\text{tot}}(E) \) and inelastic \( \sigma_{\text{in}}(E) \) cross sections of hadron-nucleus interactions in the cascade inside a macroscopic target.
Table A.6: List of PARLEV parameters with their default values in SHIELD-HIT12A.

<table>
<thead>
<tr>
<th>PARLEV#</th>
<th>Description</th>
<th>Default</th>
<th>Units</th>
<th>Remarks</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>AGT: PIDABS</td>
<td>0.01</td>
<td>probability</td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>AGT: TLIMIT</td>
<td>25.0</td>
<td>fm / c</td>
<td>1)</td>
</tr>
<tr>
<td>3</td>
<td>AGT: MMES</td>
<td>0.2</td>
<td>GeV</td>
<td>1)</td>
</tr>
<tr>
<td>4</td>
<td>AGT: MBAR</td>
<td>999.999</td>
<td>GeV</td>
<td>1)</td>
</tr>
<tr>
<td>5</td>
<td>AGT: EPS1,EPS2</td>
<td>0.007</td>
<td>GeV</td>
<td></td>
</tr>
<tr>
<td>6</td>
<td>AGT: VPI</td>
<td>0.025</td>
<td>GeV</td>
<td></td>
</tr>
<tr>
<td>7</td>
<td>AGT: C1,C2</td>
<td>0.545</td>
<td>fm</td>
<td>Wood-Saxon param.</td>
</tr>
<tr>
<td>8</td>
<td>AGT: D1,D2</td>
<td>0.05</td>
<td>(N/A)</td>
<td></td>
</tr>
<tr>
<td>9</td>
<td>AGT: R0N1,R0N2</td>
<td>1.097</td>
<td>fm</td>
<td></td>
</tr>
<tr>
<td>10</td>
<td>AGT: R0N H(2,1)</td>
<td>2.704</td>
<td>fm</td>
<td></td>
</tr>
<tr>
<td>11</td>
<td>AGT: R0N H(3,1)</td>
<td>2.55</td>
<td>fm</td>
<td></td>
</tr>
<tr>
<td>12</td>
<td>AGT: R0N He(A,2)</td>
<td>2.55</td>
<td>fm</td>
<td>$A \neq 4$</td>
</tr>
<tr>
<td>13</td>
<td>AGT: R0N He(4,2)</td>
<td>2.157</td>
<td>fm</td>
<td></td>
</tr>
<tr>
<td>14</td>
<td>AGT: R0N Li(6,3)</td>
<td>3.317</td>
<td>fm</td>
<td></td>
</tr>
<tr>
<td>15</td>
<td>AGT: R0N Li(A,3)</td>
<td>3.11</td>
<td>fm</td>
<td>$A \neq 6$</td>
</tr>
<tr>
<td>16</td>
<td>AGT: R0N Be(A,4)</td>
<td>3.25</td>
<td>fm</td>
<td>all Be isotopes</td>
</tr>
<tr>
<td>17</td>
<td>AGT: R0N B(10,5)</td>
<td>3.16</td>
<td>fm</td>
<td></td>
</tr>
<tr>
<td>18</td>
<td>AGT: R0N B(4,5)</td>
<td>2.48</td>
<td>fm</td>
<td>$A &lt; 10$</td>
</tr>
<tr>
<td>19</td>
<td>AGT: R0N C(A,6)</td>
<td>2.48</td>
<td>fm</td>
<td>$A \leq 10$</td>
</tr>
<tr>
<td>20</td>
<td>AGT: R0N N(4,7)</td>
<td>2.48</td>
<td>fm</td>
<td>$A \leq 10$</td>
</tr>
<tr>
<td>21</td>
<td>AGT: R0N O(A,8)</td>
<td>2.48</td>
<td>fm</td>
<td>$A \leq 10$</td>
</tr>
<tr>
<td>22</td>
<td>PRECO: PRECOM</td>
<td>1.0</td>
<td>(N/A)</td>
<td></td>
</tr>
<tr>
<td>23</td>
<td>DEEX: ICAN</td>
<td>0.0</td>
<td>flag: 0 or 1</td>
<td></td>
</tr>
<tr>
<td>24</td>
<td>DEEX: IMA</td>
<td>3.0</td>
<td>(N/A)</td>
<td></td>
</tr>
<tr>
<td>25</td>
<td>DEEX: EPSIL0</td>
<td>16.0</td>
<td>MeV</td>
<td></td>
</tr>
<tr>
<td>26</td>
<td>DEEX: FKACOL</td>
<td>2.0</td>
<td>(N/A)</td>
<td></td>
</tr>
<tr>
<td>27</td>
<td>DEEX: RNCL</td>
<td>0.0</td>
<td>fm</td>
<td></td>
</tr>
<tr>
<td>28</td>
<td>DEEX: AMS</td>
<td>0.125</td>
<td>MeV$^{-1}$</td>
<td>level density</td>
</tr>
<tr>
<td>29</td>
<td>DEEX: AMFS</td>
<td>0.125</td>
<td>MeV$^{-1}$</td>
<td>level density</td>
</tr>
<tr>
<td>30</td>
<td>DEEX: PARLEV(30)</td>
<td>16.0</td>
<td>Mass number $A_{\text{max}}$ for Fermi density</td>
<td></td>
</tr>
<tr>
<td>31</td>
<td>DEEX: PARLEV(31)</td>
<td>0.002</td>
<td>GeV/nucleon</td>
<td>min of $U_{\text{mut}}/A$</td>
</tr>
<tr>
<td>32</td>
<td>DEEX: ILEVRA</td>
<td>2.0</td>
<td>(flag: 1 or 2)</td>
<td>ground or excited</td>
</tr>
<tr>
<td>33</td>
<td>DEEX: FKAP1-Coulomb</td>
<td>0.65</td>
<td>(N/A)</td>
<td></td>
</tr>
<tr>
<td>34</td>
<td>DEEX: FKAP2-Volume</td>
<td>18.0</td>
<td>(N/A)</td>
<td></td>
</tr>
<tr>
<td>35</td>
<td>(Reserved)</td>
<td>0.0</td>
<td>-</td>
<td></td>
</tr>
<tr>
<td>36</td>
<td>(Reserved)</td>
<td>0.0</td>
<td>-</td>
<td></td>
</tr>
<tr>
<td>37</td>
<td>(Reserved)</td>
<td>0.0</td>
<td>-</td>
<td></td>
</tr>
<tr>
<td>38</td>
<td>(Reserved)</td>
<td>0.0</td>
<td>-</td>
<td></td>
</tr>
<tr>
<td>39</td>
<td>SIGION</td>
<td>1.00</td>
<td>(N/A)</td>
<td>Renorm $\sigma_{AA}(E)$</td>
</tr>
<tr>
<td>40</td>
<td>MICROD</td>
<td>1.00</td>
<td>(N/A)</td>
<td>Renorm $\sigma_{hA}(E)$</td>
</tr>
</tbody>
</table>

1) For high energies
1 fm = $10^{-13}$ cm
$c = 3 \cdot 10^{10}$ cm/sec
N/A : Non-applicable, dimensionless
Appendix B

Additional background information

B.1 Optional antiproton annihilation corrections APCORR

When negatively charged antimatter particles traverse matter come to rest they will get captured to an atom replacing an orbital electron and eventually annihilate on the atomic nucleus.

In a compound target the projectile (in SHIELD-HIT12A the possible negative antiparticles are $p^-$, $K^-$ and $\bar{p}$) can get captured to each of the atoms in the compound. The decision which atom the projectile is captured to is based on the calculation of the capture probability for the individual atoms in the compound target. Currently the default calculation is based on the Fermi-Teller $Z$-law [29], which states that the capture probability is proportional to the $Z$-value of the target atom. But this law has shown not to give the correct description of experimental data (see e.g. [42]). Physically more correct theories have been developed, and a combination of the formulas found by Daniel [43] and by Ponomarev [42] and extended by Pawlevicz [44] (called the PPD-law here) has been implemented into SHIELD-HIT12A – for $p^-$, $K^-$ and $\bar{p}$ projectiles. The PPD-law is enabled by setting the flag APCORR to 1 in beam.dat.

The two calculation routines have been tested against experimental data for the antiproton depth dose curve in water. It was found that the current calculation routine using the Fermi-Teller $Z$-law agreed well with the experimental data, whereas the new implementation of the PPD-law makes the simulations overestimate the dose in the top of the Bragg peak, but this can be resolved by scaling the antiproton cross sections by a factor of 1.08. This scaling is enabled along with the PPD-law, when setting APCORR to 1 in beam.dat. [Article in preparation.]

Therefore nearly the same result should be seen when transporting antiprotons setting APCORR to either 0 or 1 (this is tested having water as target medium), but a better physically justified theory is enabled by setting APCORR to 1. This new implementation is for research and development purposes and are subject to change as more experimental data become available.
B.2 History

This is a short ChangeLog that describes the most noteworthy changes in the A-branch of SHIELD-HIT.

**SHIELD-HIT12A**

SHIELD-HIT12A is the first version of SHIELD-HIT that was ever released in public.

- All changes from SHIELD-HIT10 were merged into SHIELD-HIT12A; these were mostly minor bugfixes.
- `for022.dat`, `for023.dat` and `pasin.dat` were renamed into `mat.dat`, `beam.dat` and `geo.dat`.
- Comments allowed in `geo.dat`.
- `mat.dat` is now in free format, making extension with new cards possible without breaking compatibility. Comments are now allowed, too.
- `beam.dat` is now in free format, making extension with new cards possible without breaking compatibility. Comments are now allowed, too.
- Flat circular and flat square beams are now possible.
- Transportation in any direction, specified with the `BEAMDIR` card.
- Implemented Gaussian or flat optional beam divergence and beam focus model using the `BEAMDIV` card.
- Updated inelastic antiproton cross sections. This is still experimental.
- Implementation of a new Vavilov straggling function, which is 5-6 times faster than the old version taken from GEANT3.21. This speeds up SHIELD-HIT12A by 30-40% for a typical C-12 depth dose calculation.
- New Moliere scattering function.
- Removed code that was taken from GEANT3.21 thereby liberating SHIELD-HIT12A from GPL license and Copyright issues.
- No need for `atab.dat` and `tabnuc.dat` files as they are hardcoded now.
- Bug fixes, most prominently the missing fluence in vacuum issue (ticket #27).
- Script for generating stopping power data files from libdEdx.
- Scoring by zone is now provided by the new scoring system. Zone-scoring was also possible with the old scorer, however, incompatible with parallelization.
- New detector: alanine dosimeter response model is now included following [35].
- Implementation of neutron cross sections (below 14.5 MeV) for Argon. Update of neutron cross sections for Tungsten.
- Update of antiproton total and inelastic inflight annihilation cross section, following Sychev’s data book.
- `TRACE` scorer added for displaying particle trajectories.
- Voxelized structures in the form of CT-cubes can be loaded.
SHIELD-HIT10A

SHIELD-HIT10A was never released as such, but was subject to continuous development. SHIELD-HIT10A started as a fork from the SHIELD-HIT08 code. Changes to SHIELD-HIT10 were still migrated to SHIELD-HIT10A though during the development process.

- This manual was initiated.
- The shieldhit executable accepts a directory as argument, and -n option provides an iterator for multiple runs in the same directory, which is useful for parallelization.
- Arbitrary mesh / cylindrical scoring, with several estimators.
- External beam files can be read, i.e. Spread out Bragg-peak data files derived from GSI raster scan files.
- Ripple filters can be defined and loaded.
- RANLUX random number generator replaced by RANSHE.
- Added script shield_detect2ascii which can average results from multiple parallelized runs.
- Baraschenkov’s cross sections updated for heavy ion - heavy ion reactions.
- Fermi-Breakup: free coulomb energy and free volume parameters changed from 1.0 to 0.65 and 18.0 respectively (see PARLEV33 and 34).
Appendix B. Additional background information
Bibliography


[23] GEANT, Detector Description and Simulation Tool; 1994. Section PHYS325: Moliere scattering.


Index

alanine detectors, 45
AMASS, 14
APCORR, 18
average beta detectors, 44
average energy detectors, 44
beam modulator, 23
beam.dat, 8, 17
example, 26
examples, 26
BEAMDIR, 18
BEAMDIV, 18
BEAMPOS, 19
BEAMSIGMA, 19
BMODMC, 19
BMODTRANS, 19
bodies, 27
convertmc, 5, 9, 51
counter detectors, 45
DDD detectors, 45
DELTAE, 19
DEMIN, 19
detect.dat, 8, 34
example, 46
detectors, 44
alanine, 44
average beta, 44
average energy, 44
charged particle fluence, 44
counter, 45
crossing fluence, 44
DDD, 44
dose, 44
dose-averaged LET, 44
energy, 44
fluence, 44
LET-fluence, 44
neutral particle fluence, 44
Q, 15
SPC, 15
track-averaged LET, 44
differential example, 46
dose-averaged LET detectors, 44
EMTRANS, 17
END mat.dat, 14
examples beam.dat, 26
detect.dat, 46
differential, 46
geo.dat, 24, 30, 31, 33
mat.dat, 15
scoring, 16
SPC, 16
voxscore, 46
EXTSPEC, 19
fluence charged particle, 44
neutral particle, 44
for*, 19
geo.dat, 8, 27
example, 24
examples, 30, 31, 38
HIPROJ, 20
ICRU mat.dat, 14
IVALUE, 14
JPART0, 20

80
LET
  dose-averaged, 44
  track-averaged, 44
Linux
  installation, 5
LOADDEDX, 14
LOENT, 8
MAKELN, 20
mat.dat, 7
  example, 15
media, 30
MEDIUM, 14
MSCAT, 20

NEUTRFAST, 20
NEUTRLCUT, 20
NSTAT, 20
NUCLID, 14
NUCRE, 20
output files, 49

parameter
  load external, 22
pymchelper, 5

Q
  detectors, 43

RHO, 14
RiFi, 4
ripple filter, 4
RNDSEED, 20

scoring, 3
  cylindrical, 56
  examples, 46
  geomap, 51
  voxscore, 43
  zone, 35
shield_dEdx, 52
source
  load external, 22

SPC
  detectors, 43
  example, 16

STATE, 14
stopping power
  load external, 12
stopping power, 52

STRAGG, 21

TLE, 4
TMAX0, 21
TRACE, 22
Track Length Estimation, 4
  track-averaged LET
detectors, 44

uncertainties, 45
USEBMOD, 21
USECBEAM, 21
USEPARLEV, 21

VOX, 28
voxel
  materials, 13
  scoring, 13
  voxel body, 28
VOXMED, 14
voxscore
  example, 16

Windows
  installation, 5

zones, 29
  geomap, 34
  scoring, 35