

Advanced topics

A brief overview of some advanced features

Lecture overview

- Introduction
- Geometry replication (LATTICE)
- Accessing detailed particle information
- Medical applications
- Special sources
- Other advanced features and conclusion

Advanced topics

- We have covered the fundamental capabilities of FLUKA during this training
- These features are controlled through the default options and can be combined to cover a
 wide variety of problems, including complex geometries, scoring needs etc.
- Still, more complicated/specialised requirements may arise, such as:
 - Replicating geometries
 - Advanced sources
 - Custom scoring, extraction of detailed particle information (on an event-by-event basis)
 - Medical applications
 - And more...
- Some of these advanced capabilities require modified user routines and compilation of custom FLUKA executables
 - Default versions of the user routines can be found in the pathtofluka/src/user/ directory
- We will cover a few examples in this lecture

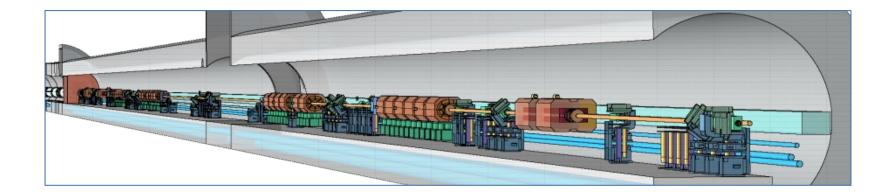


Geometry replication: LATTICE



Geometry with replicated regions

- Complex geometries often exhibit replicated regions
- E.g. in particle accelerator beamlines: dipoles, quadrupoles, beam-loss monitors, etc.

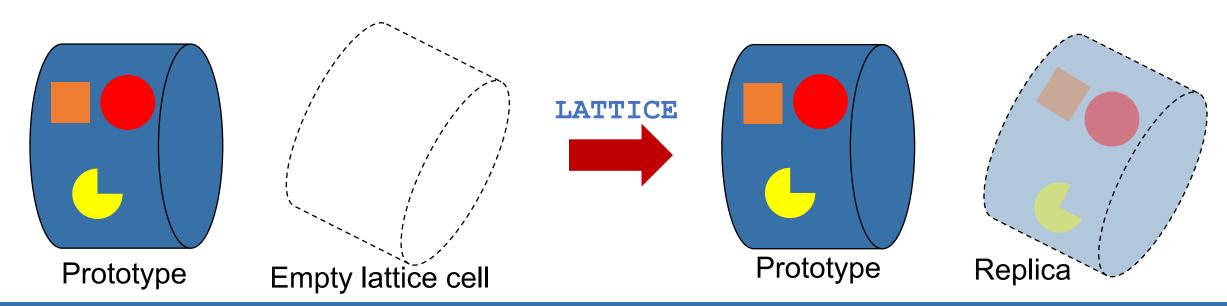


 Instead of defining every replicated region (and their interior complexity) explicitly, one may define a prototype region and then use FLUKA's LATTICE capabilities to replicate it as many times as needed elsewhere in the geometry



Lattice: the concept

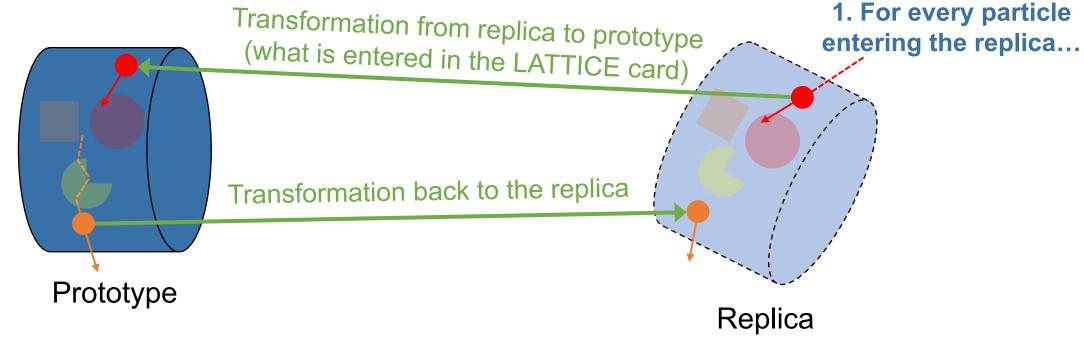
- Prototype region: a region enclosing a complex geometry (full detail of various bodies, regions, material definitions, etc). Typically, the prototype is of a simple shape (RCC, SPH, etc)
- Lattice cell region: empty transformed prototype region where prototype region contents should be effectively replicated (see next slide).
- The **LATTICE** card links lattice cell regions to prototype regions, effectively replicating the prototype region content in each lattice cell (see next slide).
- Replica-prototype link is established via transformation (defined with ROT-DEFI), which exactly transforms the lattice cell into the prototype (!)





Behind the scenes

2. ...its coordinates are transformed back to the prototype, where FLUKA performs the tracking

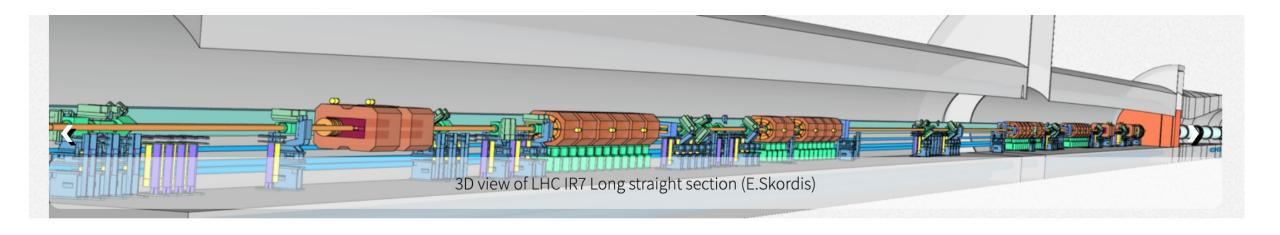


3. A resulting particle exiting the prototype...

4. ...is transformed back to the replica boundary to be tracked in the surrounding geometry

LATTICE example

 Building on the LATTICE functionality, one may construct arbitrarily complex geometries:



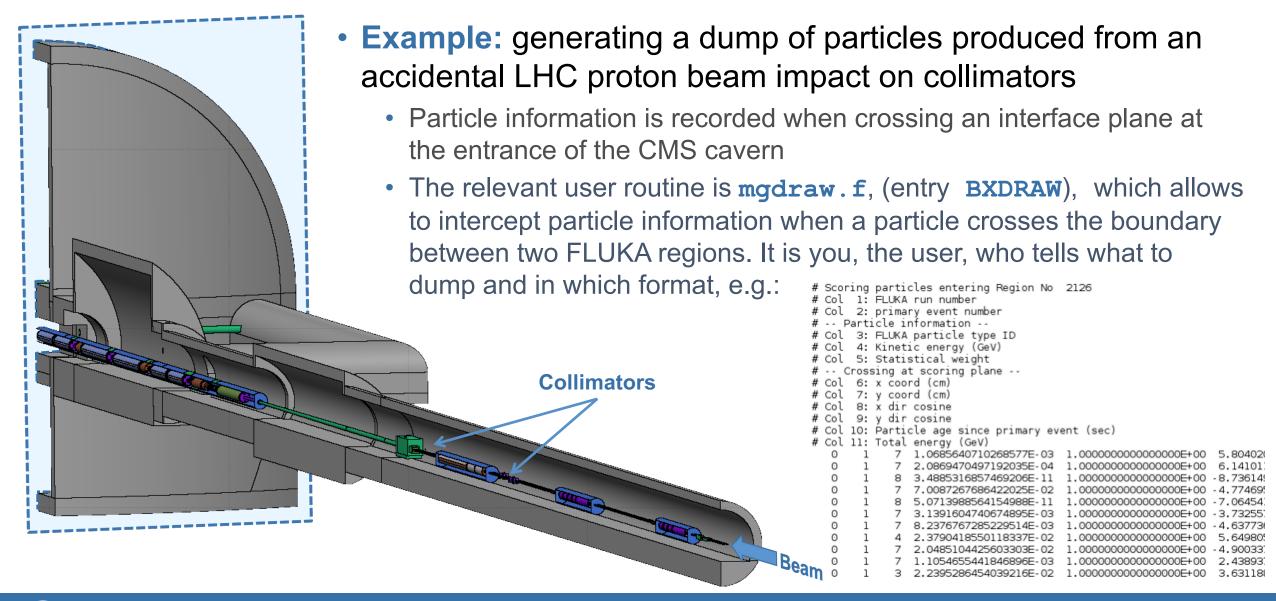
• For accelerator lines: LINE BUILDER (makes heavy use of LATTICE capabilities) [A. Mereghetti et al., IPAC2012, WEPPD071, 2687]



Accessing particle information

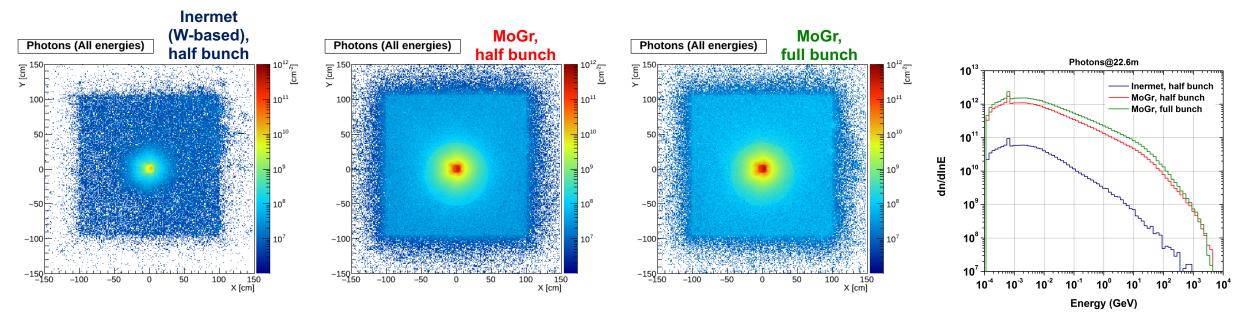


Accessing particle information



Accessing particle information

- This information can be processed to study the properties of the particle population
 - e.g. the spatial distribution near the beam-line and the energy distribution of particles for different choice of collimator materials and beam impacts:

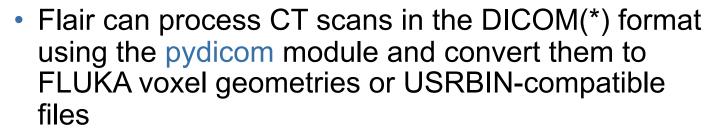


 Note: The great flexibility of this type of scoring comes with the penalty of ad-hoc post-processing with custom tools!

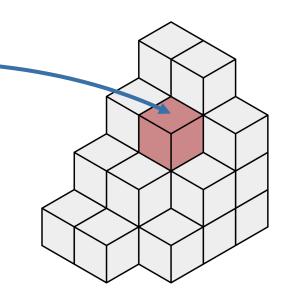


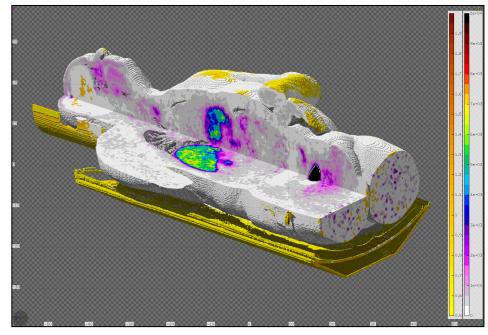


- A geometry can be described in terms of voxels, tiny parallelepipeds of equal size forming a 3dimensional grid
- Voxel geometries are especially useful for importing CT scans, e.g. for dosimetric calculations of radiotherapy treatments



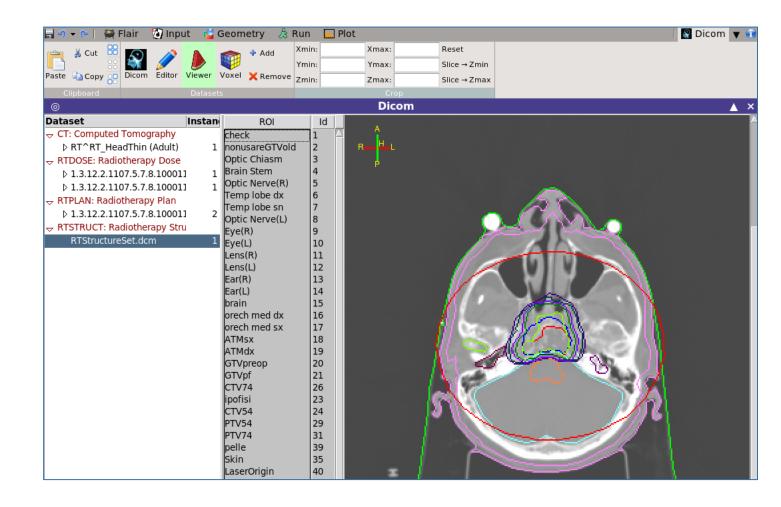
(*) DICOM (Digital Imaging and Communications in Medicine) is a medical standard for distributing any kind of medical image.





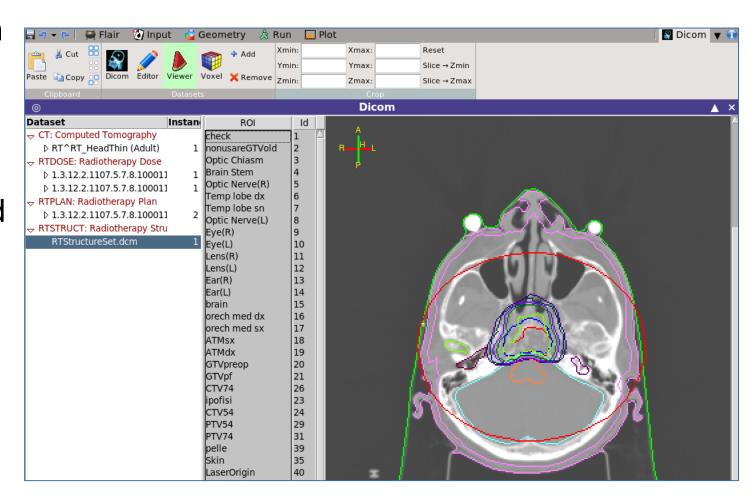


- DICOM files can be browsed, visualised and edited (e.g. anonymised)
- Voxels can be grouped into "organs" or regions of interest (RTSTRUCT in a pre-existing DICOM file)

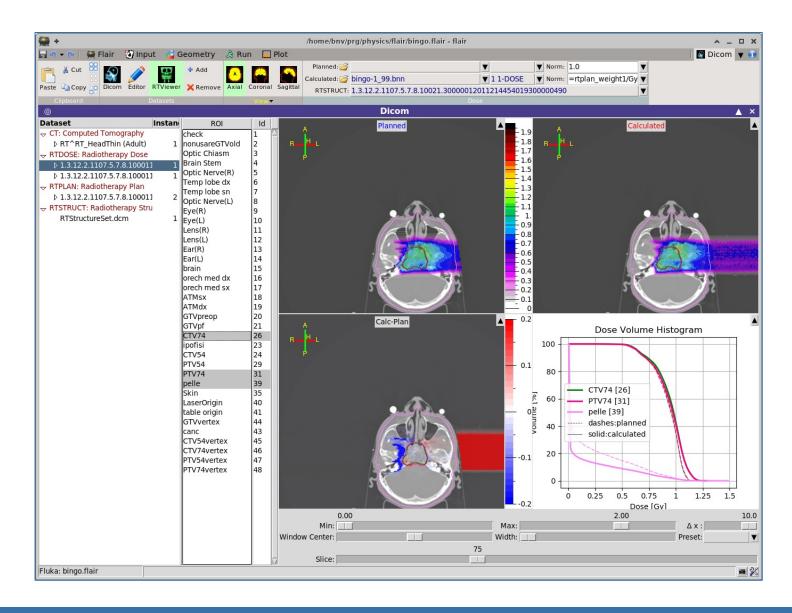


 The DICOM CT file together with RTSTRUCT can be converted to a FLUKA voxel geometry.

 This voxel geometry is contained in an RPP and can be placed within a larger combinatorial FLUKA geometry



- Schneider parametrization to convert HU in the CT file to actual materials (external file in flair website). This conversion is machine-dependent!
- Ranges of HU are assigned to various materials. To smoothly account for density variation with HU, Flair applies additional correction factors. Additionally, one may manually apply dE/dx correction factors
- The DICOM RTPLAN can be converted to a FLUKA beam input
- RTDOSE: the planned dose can be compared to USRBIN scoring
- Automatic generation of DVH (Dose Volume Histogram)
- Relevant cards: VOXELS, CORRFACT, RAD-BIOL, TPSSCORE





Special sources

Type: PPSOURCE, BIN-SOUR



Radiation sources

- Radiation sources we've seen thus far:
 - Particle beams
 - Radioactive isotopes
- FLUKA offers special sources:
 - Hadronic collision as a source term (p-p, p-Pb, etc)
 - USRBIN from another simulation as an extended source
 - SPOTBEAM: multiple beams
 - Cosmic rays
 - •
- And, if you want full customization: source_newgen.f



Input card: SPECSOUR - PPSOURCE

Beam momentum and direction:

Based on the Type, 3 different options

Type: **PPSOURCE**

A SPECSOUR	Type: PPSOURCE ▼			
	P1x:	P1y:	P1z:	
	P2x:	P2y:	P2z:	

 Momentum and direction are defined with the x, y and z components of the total laboratory momentum of either beam [GeV/c]



Input card: SPECSOUR - PPSOURCE

```
SPECSOUR
                              Type: PPSOURCE ▼
                                P1x:
                                                                                        P1z:
                                                            P1y:
                                P2x:
                                                            P2y:
                                                                                        P2z:
                                 σx:
                                                             σy:
                                                                                         σz:
                               Part: ▼
 limit-σ:
                                                                                      σ0 2:
 σθC 1:
                               σ0 1:
                                                         σθC 2:
                         NonElastic: off ▼
                                                         Elastic: off ▼
                                                                            EM dissociation: off ▼
```

Simulates a collision between two beams:

- 1st beam: Hadrons (including protons and heavier nuclei)
- 2nd beam: Only proton or heavier nuclei

Two section of the card:

- **Top**: Beam momenta and directions of the two colliding beams
- Bottom: Volume of interaction, beam particles and divergences, physics interactions



Input card: SPECSOUR - CROSSSYM

Beam momentum and direction:

z crossing angle beam 1 beam 2

Type: CROSSSYM

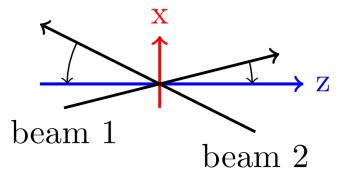
SPECSOUR Type: CROSSSYM ▼

Plab: CrossAng/2: Azimuthal:

- Plab: Total laboratory momentum of beam 1 and 2 [GeV/c]
- CrossAng/2: Half of the crossing angle [radians] (between: $0 \dots \pi/2$)
- Azimuthal: Azimuthal angle defining the crossing plane [degrees (!)]

Input card: SPECSOUR - CROSSASY

Beam momentum and direction:



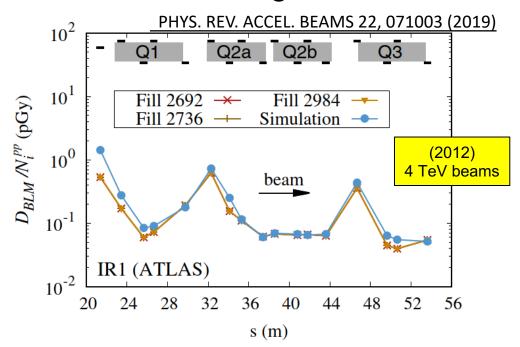
Type: **CROSSASY**

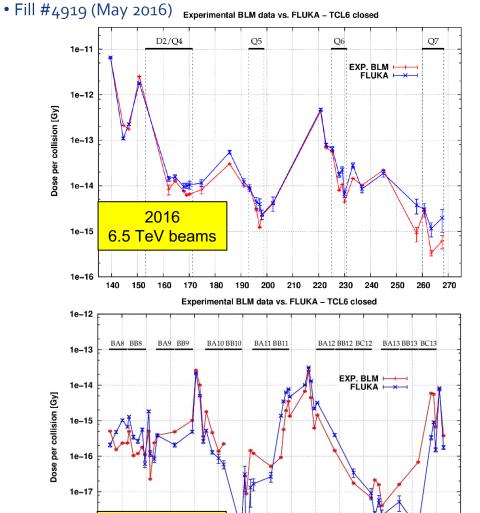
SPECSOUR	Type: CROSSASY ▼			
	P1lab:	Polar1:	Azimuthal:	
	P2-lab:	Polar2:		

- P1lab, P2lab: Total laboratory momentum of beam 1 and 2, respectively [GeV/c]
- **Polar1**, **Polar2**: Polar angle between beam 1 (2) direction and positive (negative) z direction [radians] (between: 0 ... π/2)
- Azimuthal: Azimuthal angle defining the crossing plane [degrees (!)]

Example: LHC BLM signal benchmark

- Source term: p-p collision at several TeV per beam
- Beam-loss monitor (BLM) dose signal vs FLUKA prediction
- At various distances downstream (20 560 m)
- Note the excellent agreement!





Distance from IP [m]

6.5 TeV beams



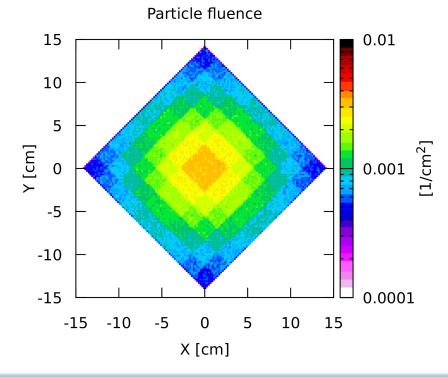
Special sources – USRBIN source

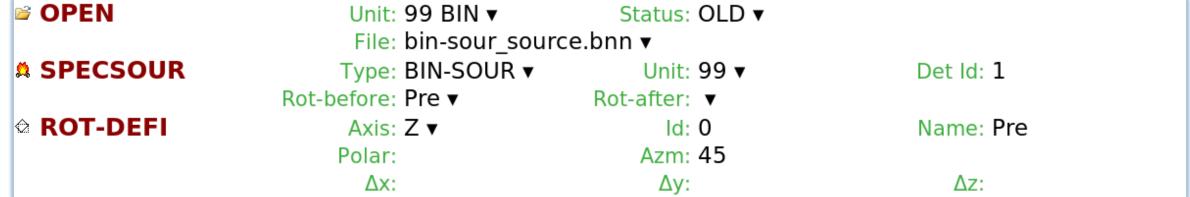
Input card: SPECSOUR

Example:

USRBIN file opened with logical unit 99 and rotated around the Z axis.

(Beam direction along Z axis)



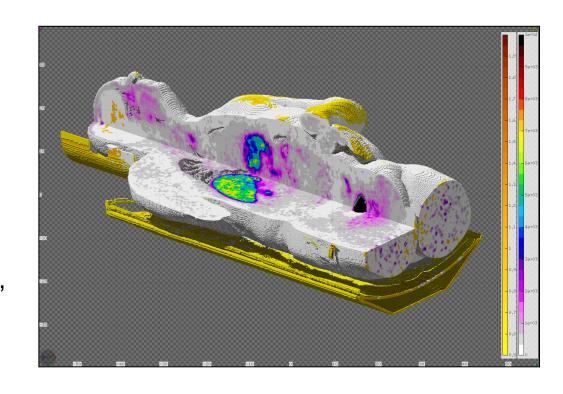




Use case

Suppose you have a PET image from a patient (⁶⁸Ga radionuclide source)

- What about dose to nearby healthy tissue?
- DICOM processed into a FLUKA VOXEL geometry
- RTDOSE can be converted to a USRBIN in Flair
- You can take the resulting USRBIN as a BIN-SOUR for such a simulation, putting ⁶⁸Ga as "beam" particle
- Positions are sampled from the USRBIN weights
- Directions as per **BEAM** card (you'd put isotropic emission)



Conclusion



All good things...

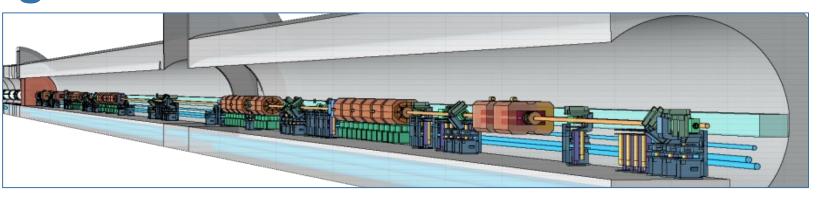
- Many other advanced/specialised topics can be studied with FLUKA...
 - Cosmic rays
 - Neutrino interactions
 - Optical photons
 - Crystal channeling
 - ...and more!
- ...and a lot more flexibility can be achieved via user routines
- This course was meant to get you started on FLUKA (building geometries, defining materials and sources, scoring), while also introducing more advanced concepts and techniques (biasing, advanced geometry features, advanced sources)
 - We would like to hear your feedback, positive and negative!
- Consider following a FLUKA advanced course!
 - Programmable user routines, advanced geometry, tracking, scoring, optical photons, and much more!



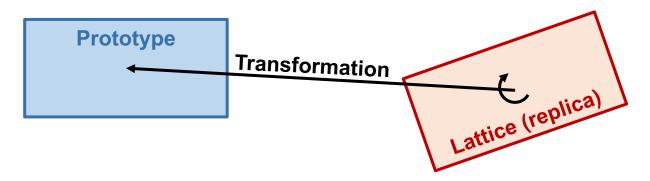


Lattice: basic usage

 Very useful for geometries where models are used multiple times (e.g. beamlines)



- The prototype is defined in detail with all the necessary information (geometry, materials etc.) inside a closed
 container body (RCC, RPP etc.)
- The lattices (replicas) are defined as "empty" regions in their correct location and declared as such via LATTICE card
- The transformations exactly mapping the replicas onto the prototype are defined using ROT-DEFI cards

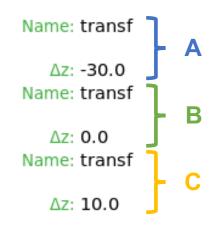


Note: You can load the lattice template in Flair for a simple working example!



Lattice example

② ROT-DEFI	Axis: Z ▼	Id: 0
	Polar: 0.0	Azm: 0.0
	Δx: 0.0	Δy: 0.0
☆ ROT-DEFI	Axis: X ▼	Id: 0
	Polar: 0.0	Azm: 30.0
	Δx: 0.0	∆y: 0.0
☆ ROT-DEFI	Axis: Z ▼	Id: 0
	Polar: 0.0	Azm: 0.0
	Δx: 0.0	∆y: 20.0



Note: if a transformation R consists of 3 rototranslations A, B and C applied in this order, then R = CBA and the inverse transformation is $R^{-1} = A^{-1}B^{-1}C^{-1}$

Here, *R* maps the replica onto the prototype, whereas *R*⁻¹ is used to transform the replica container from the prototype to the replica

