

A Monte Carlo study of target fragmentation in Protontherapy


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Target Fragmentation

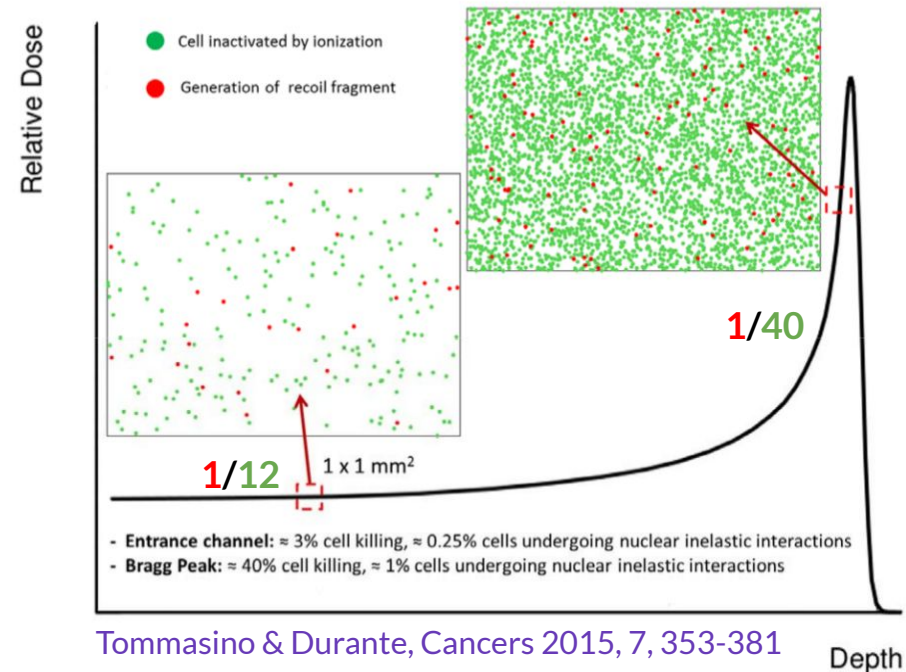
In Protontherapy, secondary particles are produced through primary beam interactions with the patient's body. The fragments created have:

 Low kinetic energy fragments \Rightarrow small range

 High Z \Rightarrow high cell killing effectiveness (dE/dx increase with Z)

 High LET \Rightarrow increase of RBE

 Higher production cross section in the entrance channel \Rightarrow relevant for healthy tissue

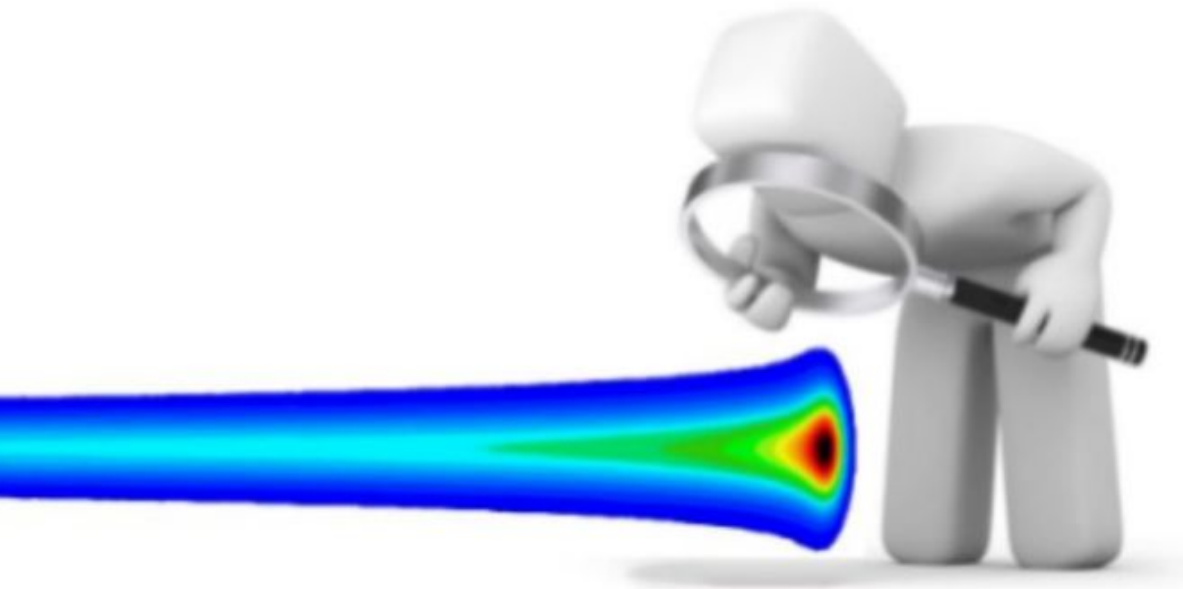


The inclusion of target fragmentation processes can be important for the accurate evaluation of the dose in the treatment.


MC study of target fragmentation

Target fragmentation is not implemented in commercial TPSs.

Monte Carlo simulations were employed to estimate the effect of target fragmentation in Protontherapy.



Fragments characterization:

- Production Cross Section
- Range
- Fluence
-  ρ Ve IT

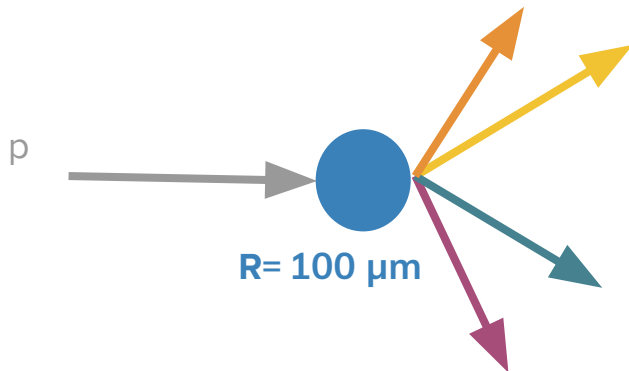
Production Cross Section

Energy spectra

The energy distribution of main fragments produced by proton beam in water have been simulated with FLUKA.

Simulation setup

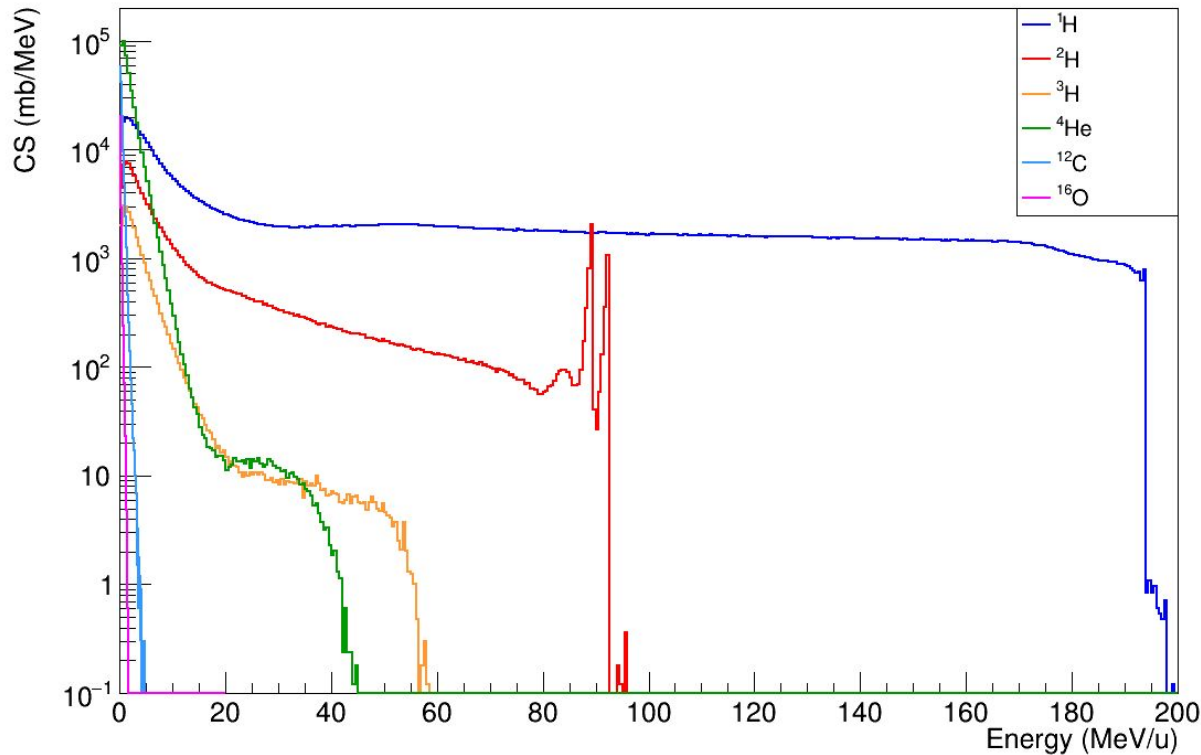
p beam on water target



The energy distribution of main fragments produced by proton beam in water have been analyzed, considering initial kinetic energy $E_p=50, 100, 150, 200$ and 250 MeV.

Energy distribution of fragments

Production CS of inelastic interaction induced by proton beam of 200 MeV in water.



	E_{MAX} (MeV/u)	E_{mean} (MeV/u)
^1H	200	65.4
^2H	95	16.7
^3H	58	3.9
^4He	45	1.7
^{12}C	4.7	0.4
^{16}O	1.7	0.1

Range

Range of fragments

Target fragments can have **high charge** (i.e. high biological effectiveness) and **low residual range**. This means that they will deposit all their energy close to their generation point.

Tommasino & Durante, Cancers 2015, 7, 353-381

Fragment	E (MeV)	LET (keV/ μm)	Range (μm)
^{15}O	1.0	983	2.3
^{15}N	1.0	925	2.5
^{14}N	2.0	1137	3.6
^{13}C	3.0	951	5.4
^{12}C	3.8	912	6.2
^{11}C	4.6	878	7.0
^{10}B	5.4	643	9.9
^8Be	6.4	400	15.7
^6Li	6.8	215	26.7
^4He	6.0	77	48.5
^3He	4.7	89	38.8
^2H	2.5	14	68.9

- Not experimental data
- Average energies of fragments are evaluated using Goldhaber formula
- Range has been estimated starting from average energy

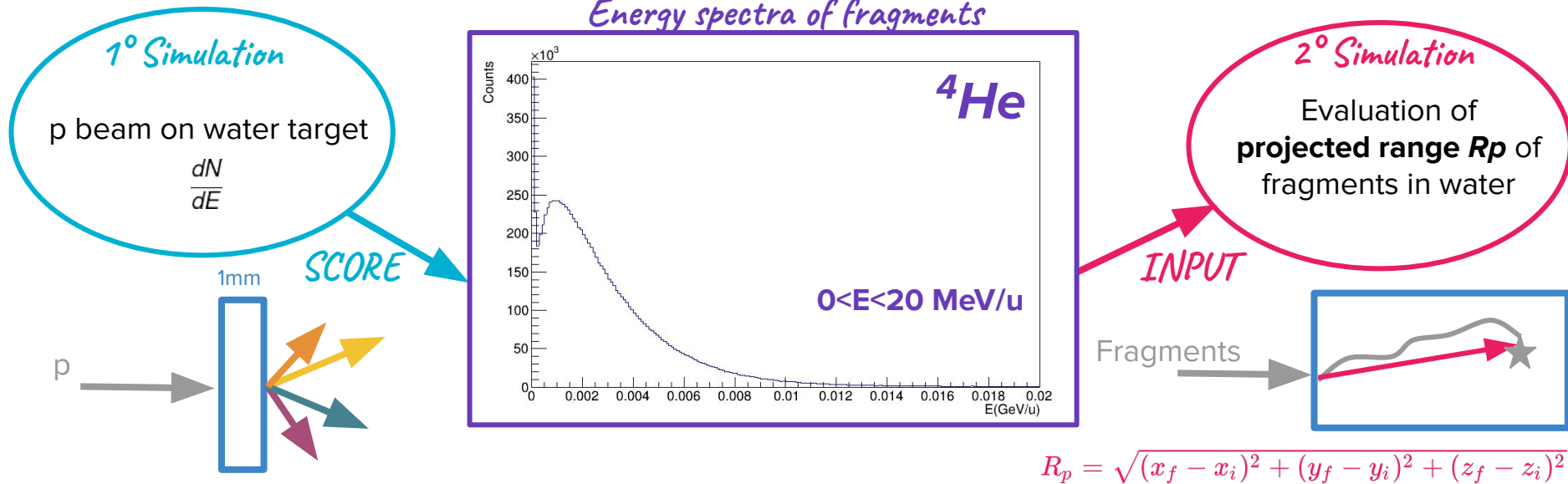


The energy and residual range of target fragments produced in water have been studied using FLUKA MC code.



Simulation setup

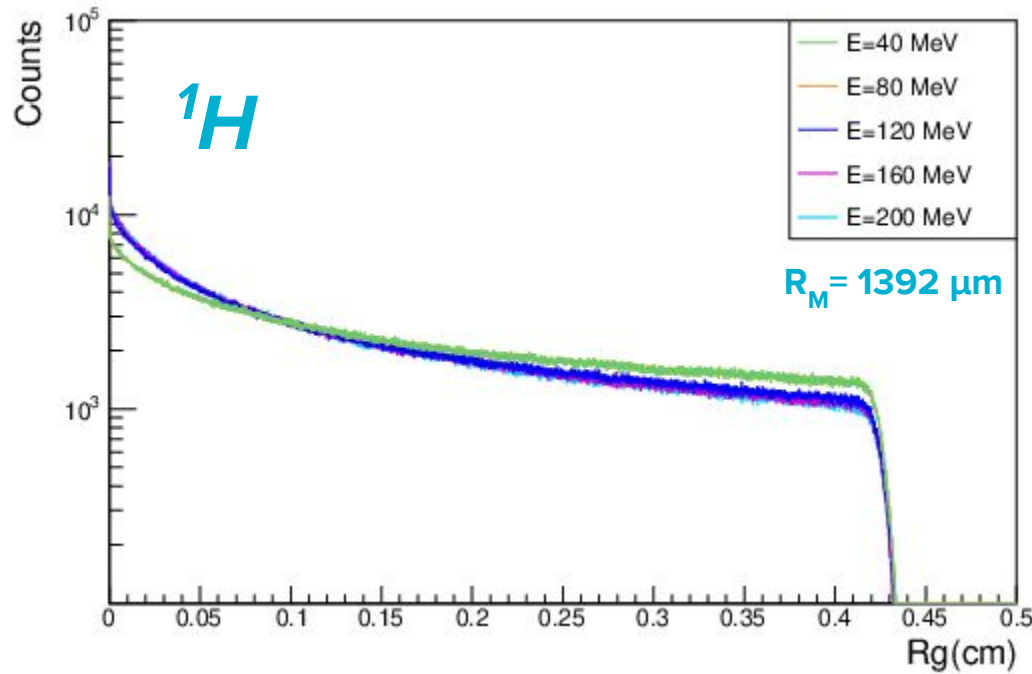
The energy spectrum and range distribution of fragments have been simulated with FLUKA:



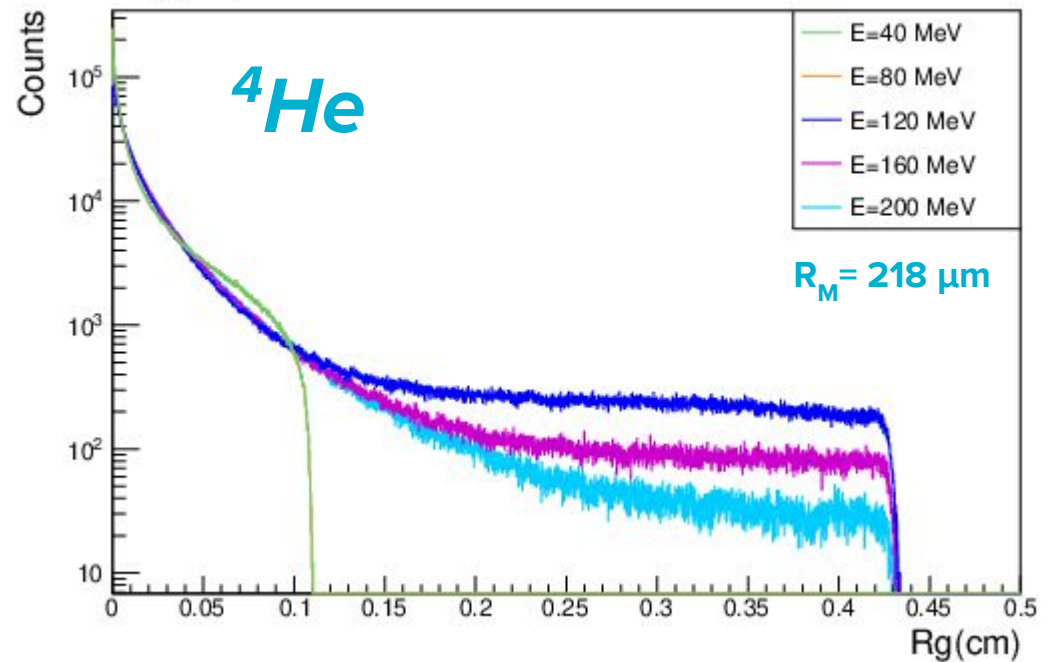
$E_p = 40, 80, 120, 160, 200 \text{ MeV/u}$.

$$R_p = \sqrt{(x_f - x_i)^2 + (y_f - y_i)^2 + (z_f - z_i)^2}$$

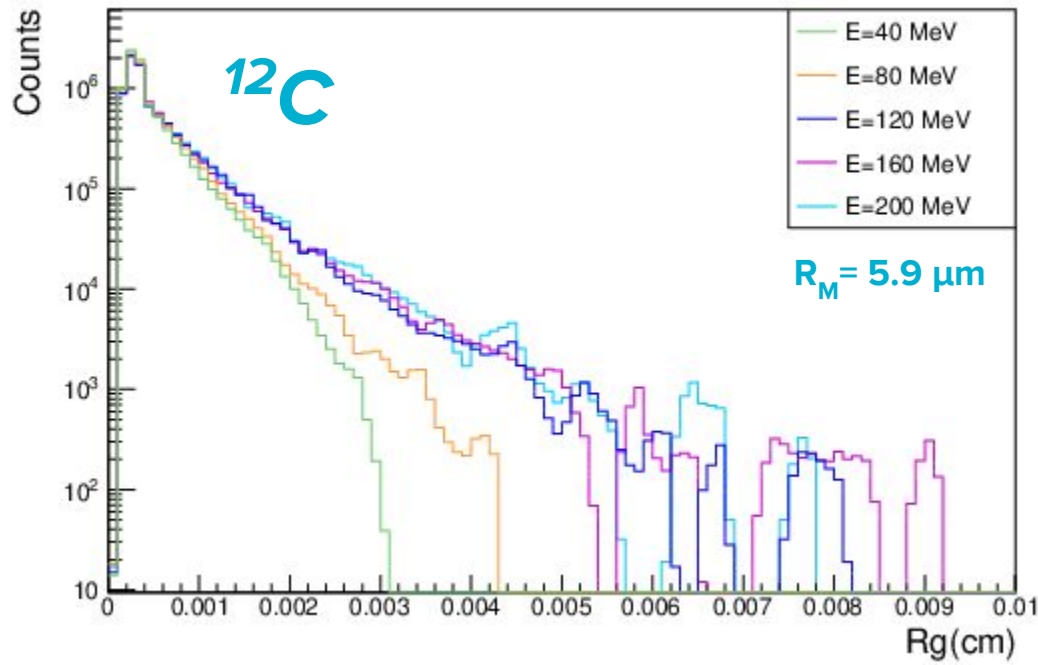
Range distribution of fragments



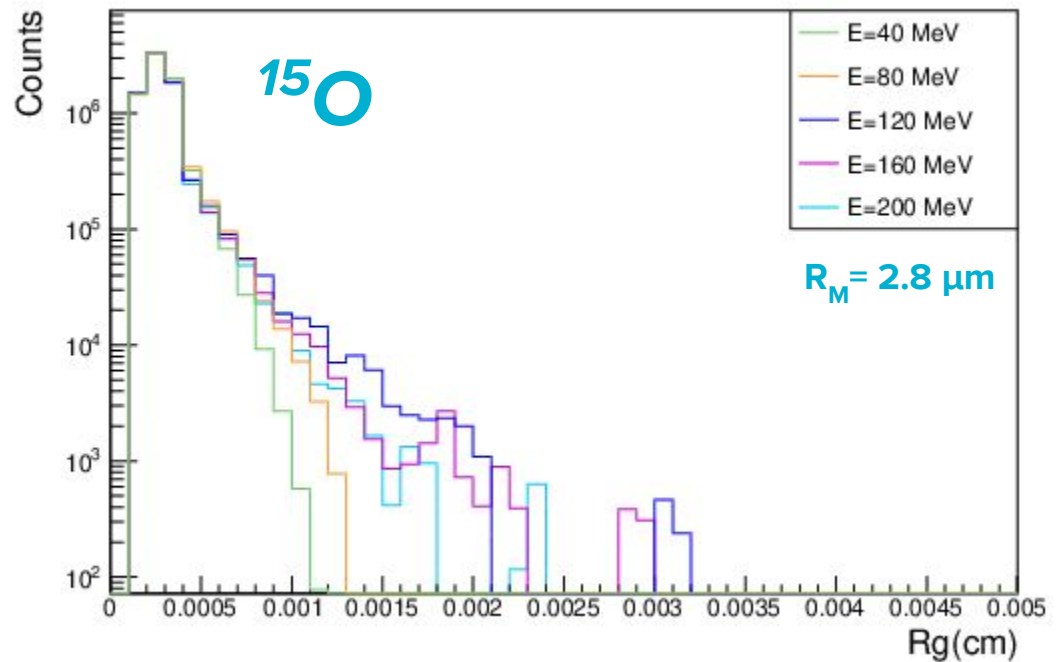
Rg = range
 R_M = mean range



Range distribution of fragments



Rg = range
 R_M = mean range



Range of fragments

Starting from the energy distribution of fragments, the **range distribution** has been evaluated.

For heavier fragments: the mean range obtained from the distribution is similar to the value reported in Cancers.

For low Z fragments: difference between average range in analytical formula and MC distribution.



MC more complete description of physical processes \Rightarrow Range is more reliable

C
O
N
C
L
U
S
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Low Z fragments $R >$ cell nucleus

Heavier fragments $R <$ cell nucleus

Colocalization effects?

FLUKA  Tommasino & Durante 

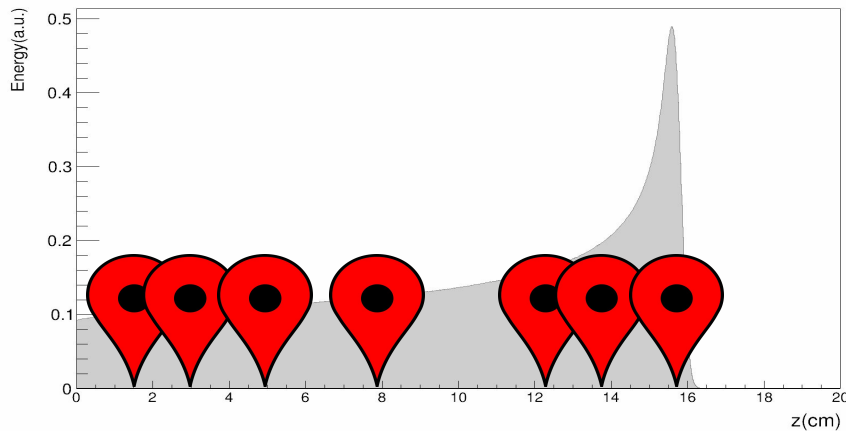
Fragment	$R_p(\mu\text{m})$	$R_{TD}(\mu\text{m})$
^1H	1392	-
^2H	2117	68.9
^3H	1383	-
^3He	465.7	38.8
^4He	218.3	48.5
^6He	229.3	-
^6Li	75.7	26.7
^7Li	74.1	-
^7Be	42.6	-
^9Be	26.1	-
^9B	24.4	-
^{10}B	14.5	9.9
^{11}B	11.2	-
^{10}C	9.9	-
^{11}C	8.1	7
^{12}C	5.9	6.2
^{13}N	4.4	-
^{14}N	4.1	3.6
^{15}O	2.8	2.3
^{16}O	3.5	-

Fragments fluence

Fragments Fluence

To include the impact of fragmentation in the TPS (TRIP98) and estimate the biological effect of fragments, their production in water has been calculated with FLUKA.

→ Fragments fluence as a function of energy evaluated at different depth



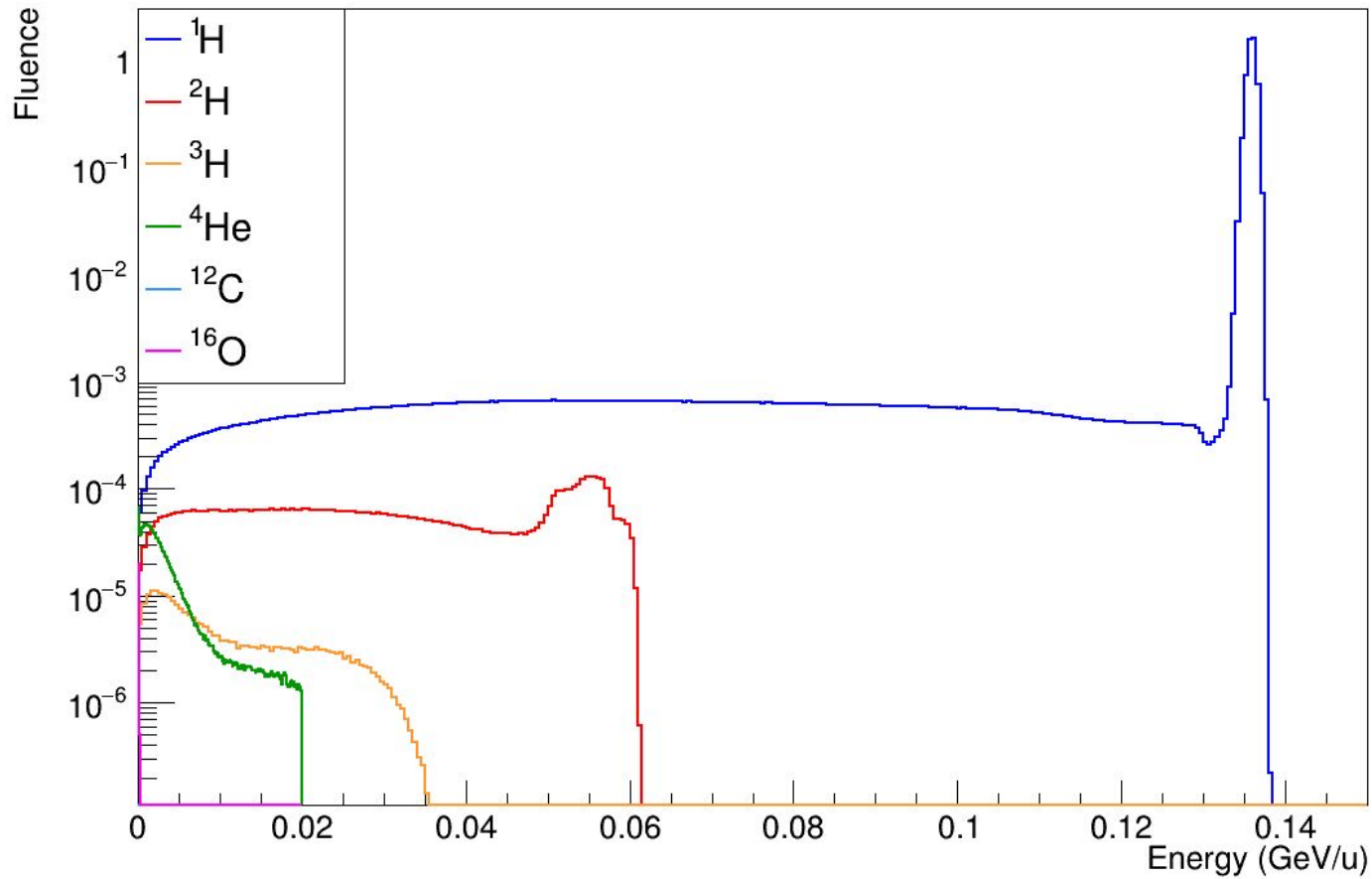
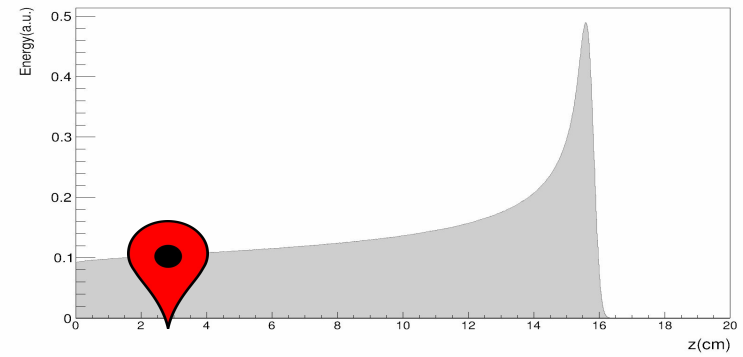
Source:
proton beam $E=150$ MeV

Geometry & Material:
parallelepiped of WATER

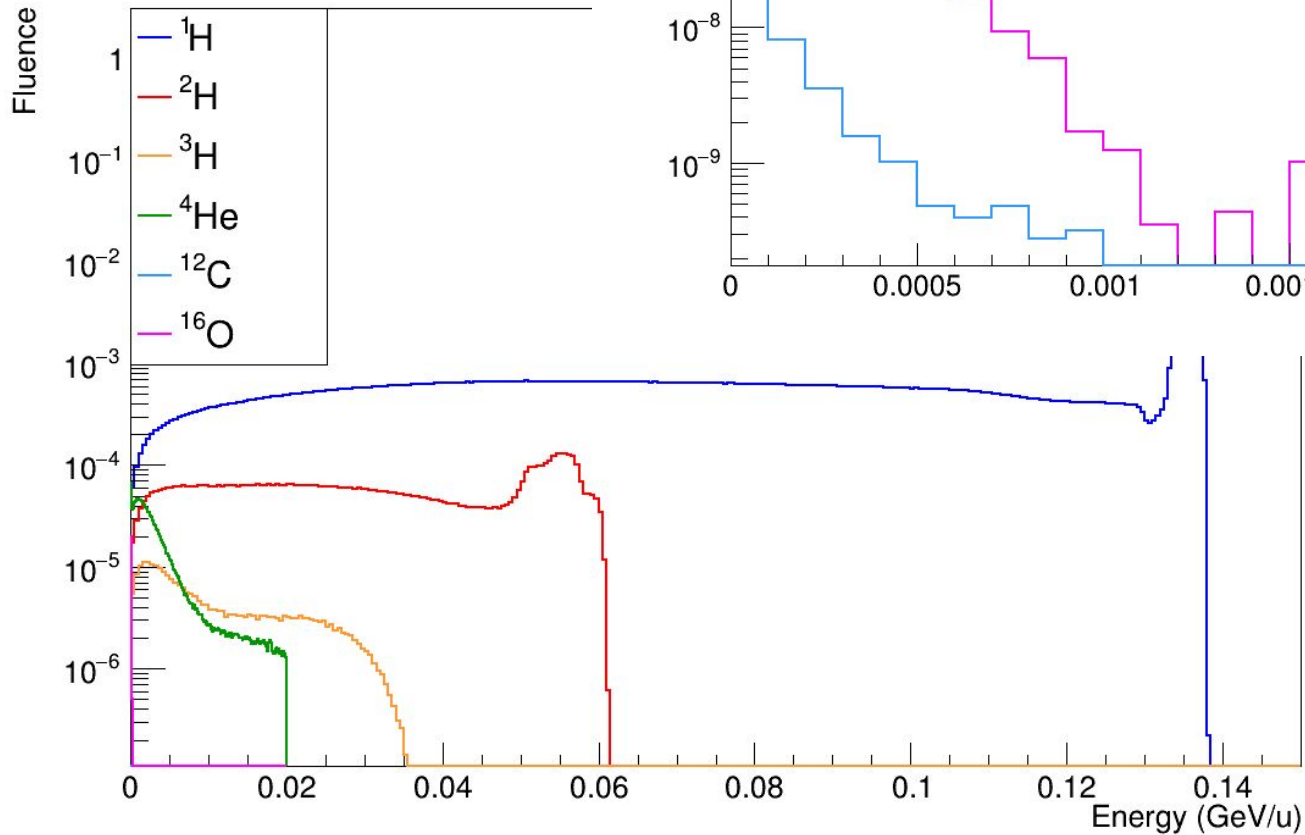
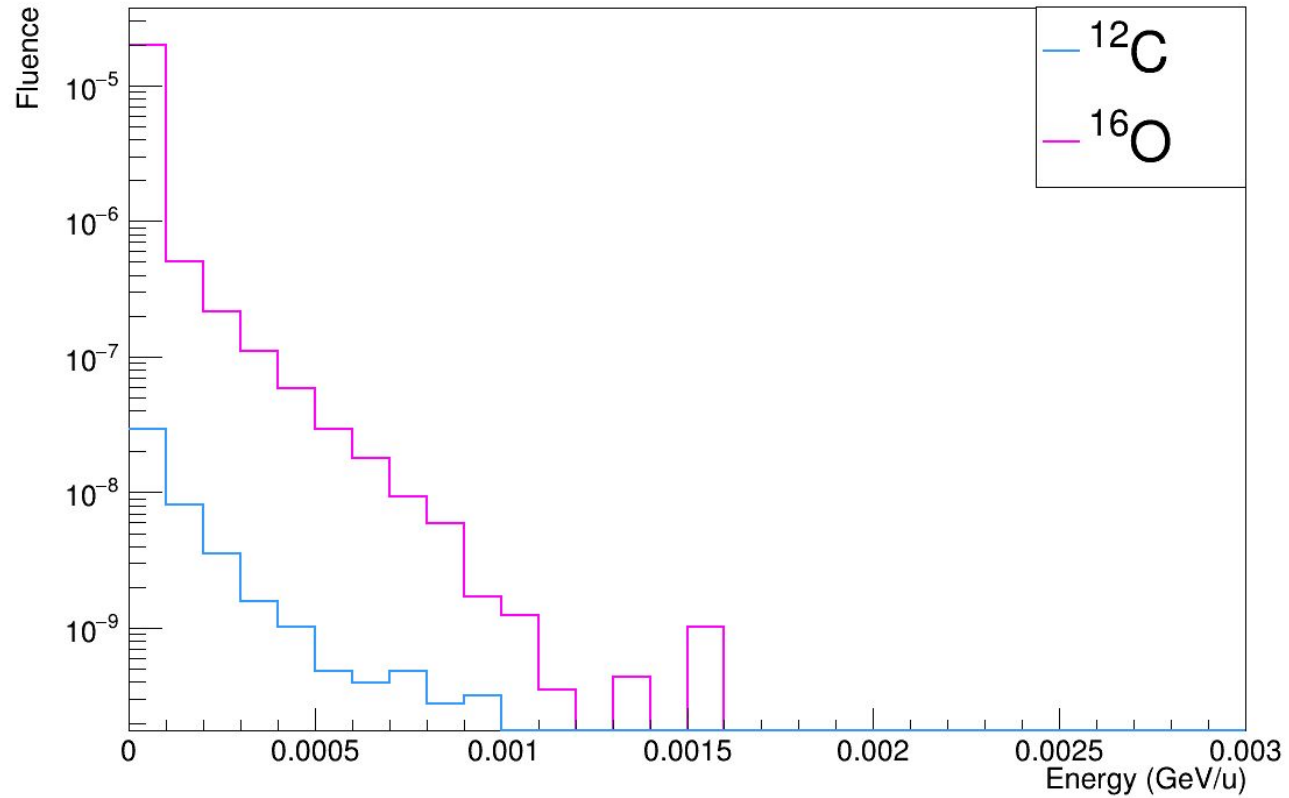
Scored quantity:
Fluence of charged particles
as a function of energy

Statistics: 10^9 primaries

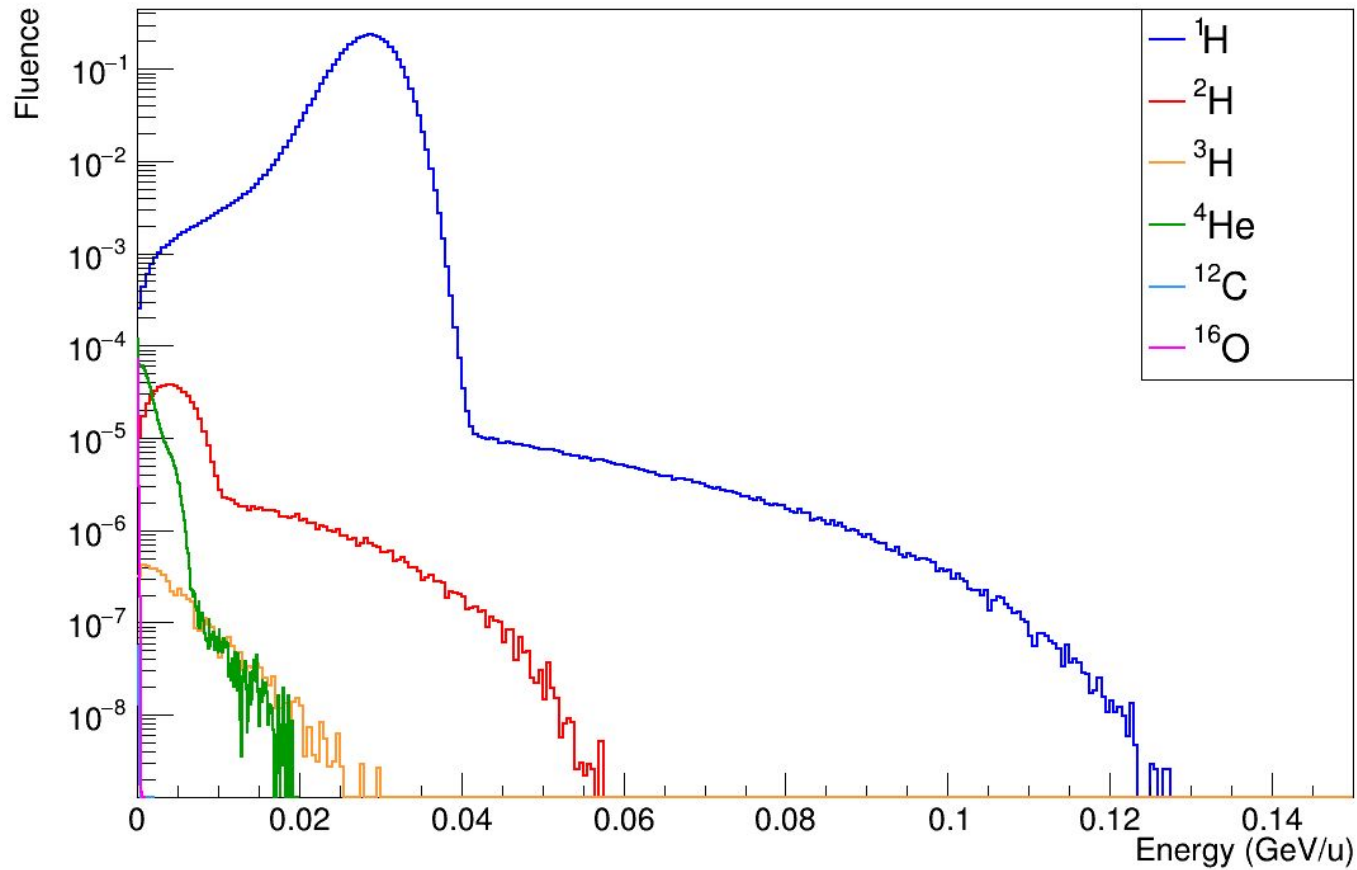
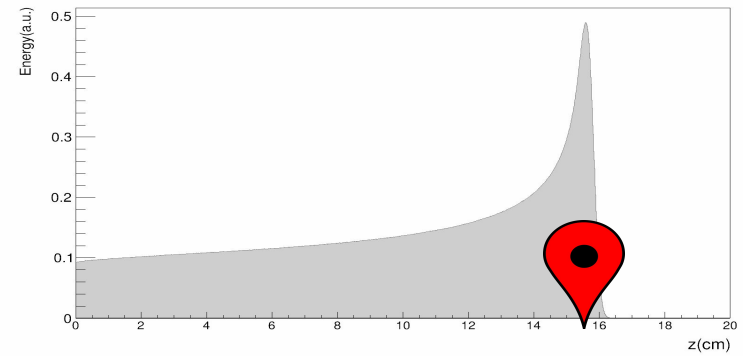
Fragments Fluence



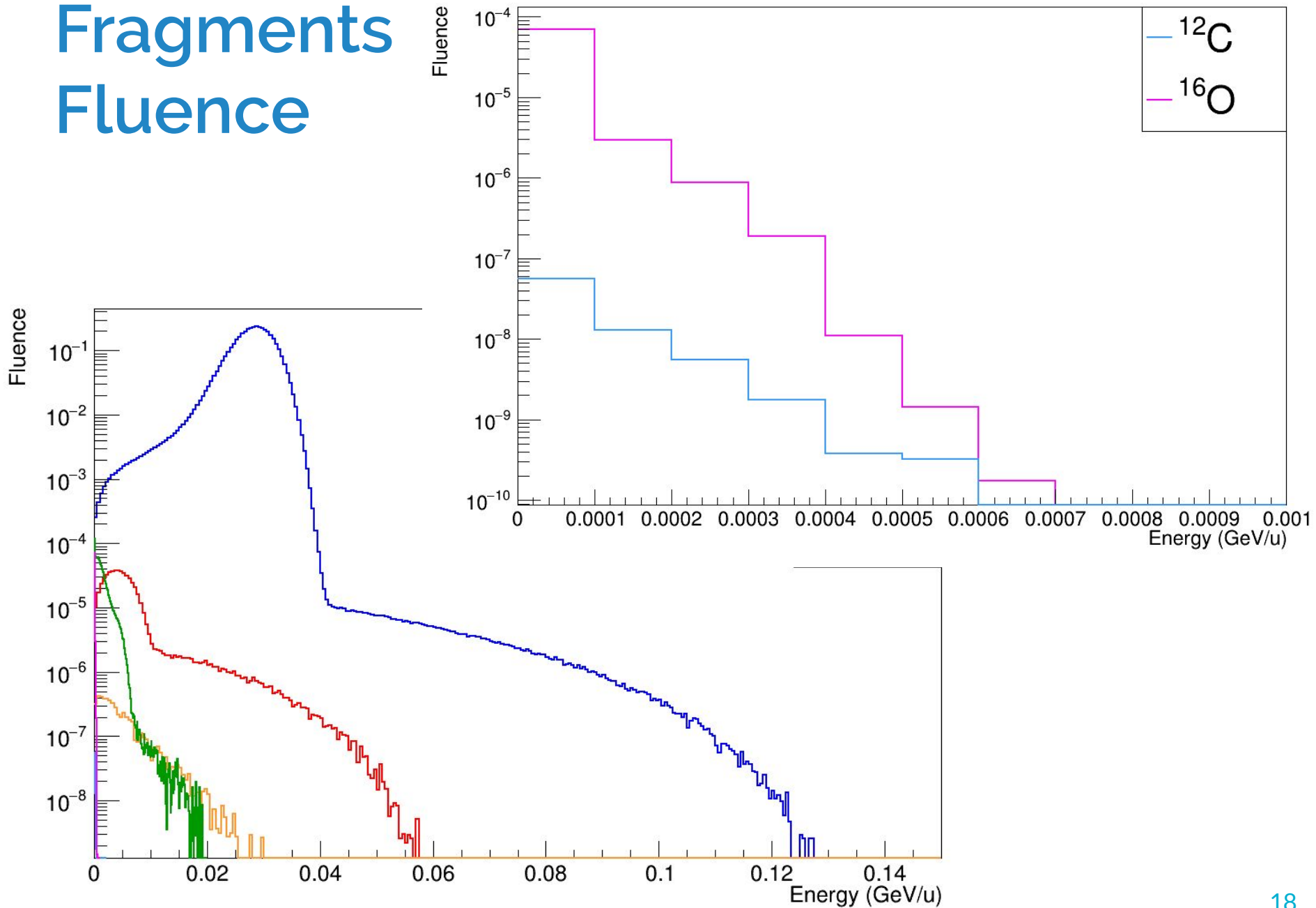
Fragments Fluence



Fragments Fluence



Fragments Fluence

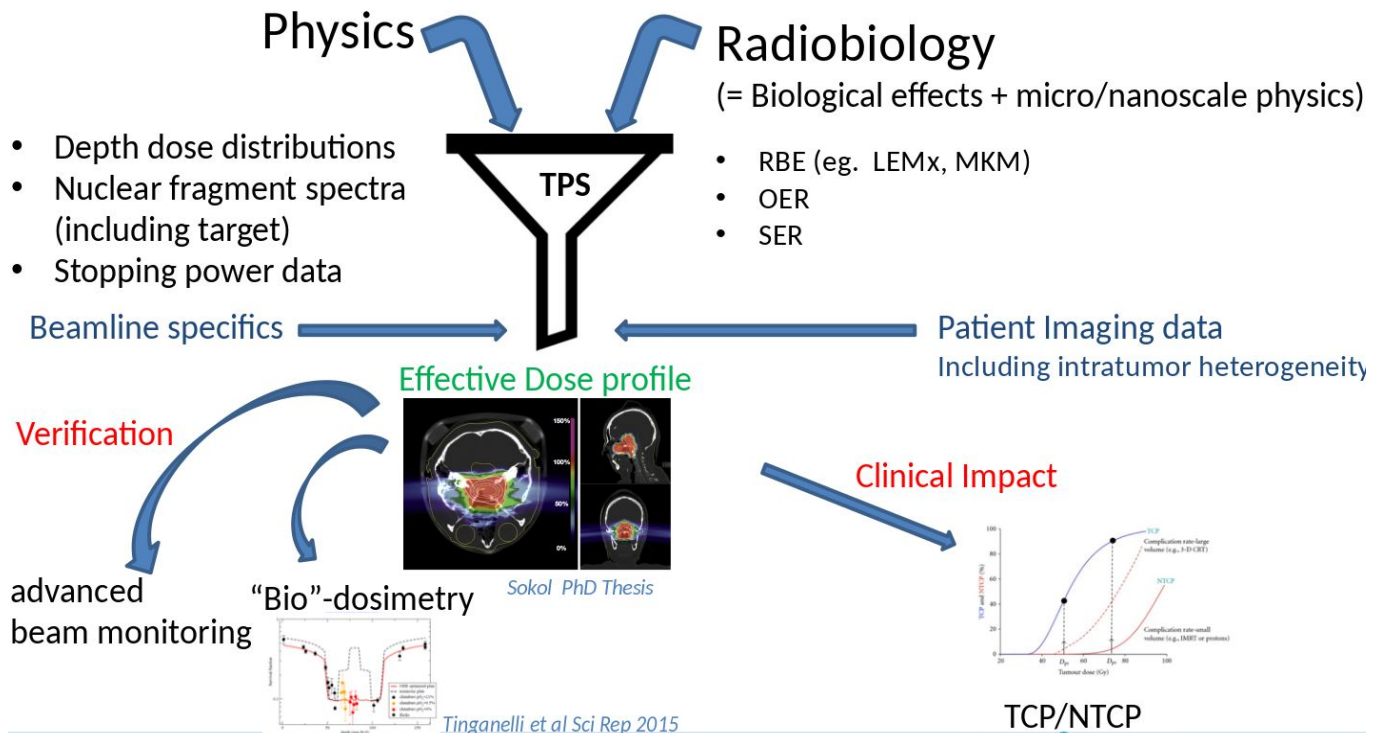


MoVe IT

Modeling and Verification
for Ion beam Treatment
planning

The main effects that will be explored in MoVe-IT project are:

- ▷ biological impact of target nuclei fragmentation
- ▷ relative biological effectiveness (RBE)
- ▷ intra-tumor heterogeneity



Target fragmentation

In order to describe the target fragmentation and estimate the biological effect of fragments, **MoVe IT propose an approach based on a RBE modellization:**



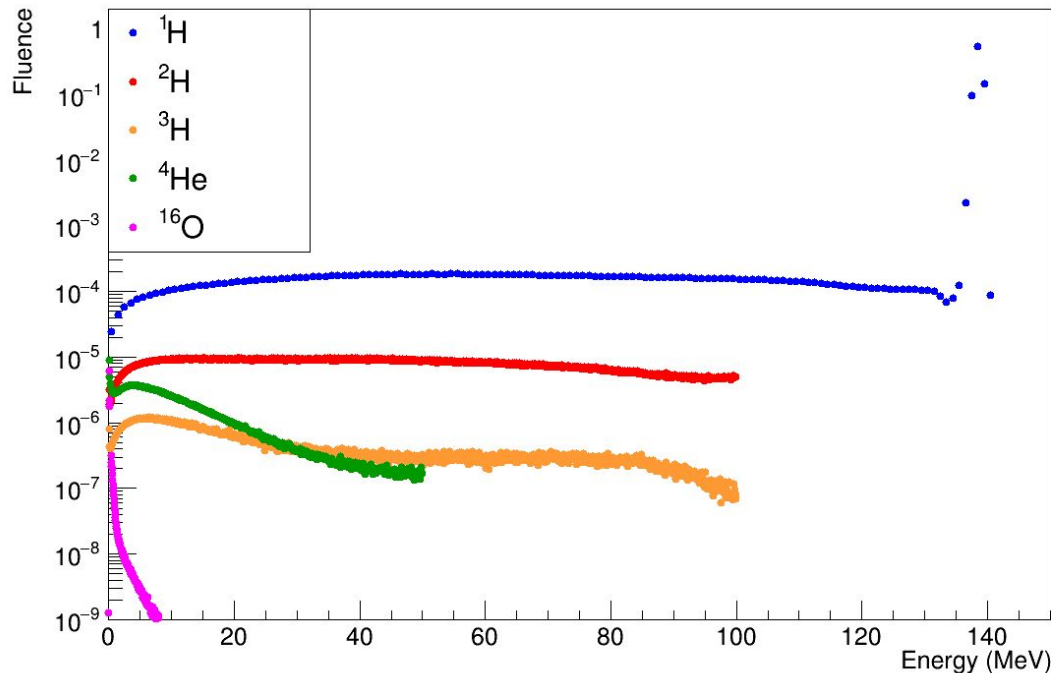
describe the physical dose including the contribution of each **single fragment produced by a proton in water**

evaluate the total RBE by using a **mixed field approach** to weights all the single fragments contributions

import the results in TRiP98 TPS and test the biological dose differences respect to the default calculations

Fragments Fluence

The fluence spectra have been obtained performing Monte Carlo simulations as separate single beams, scoring the fluence of all the produced particles.



Fluence of main fragments produced by proton beam of 150 MeV after 2 cm in water.

Fluence Database

$(E_k, E_f, A, Z, z_{\text{depth}})$



The mixed field model

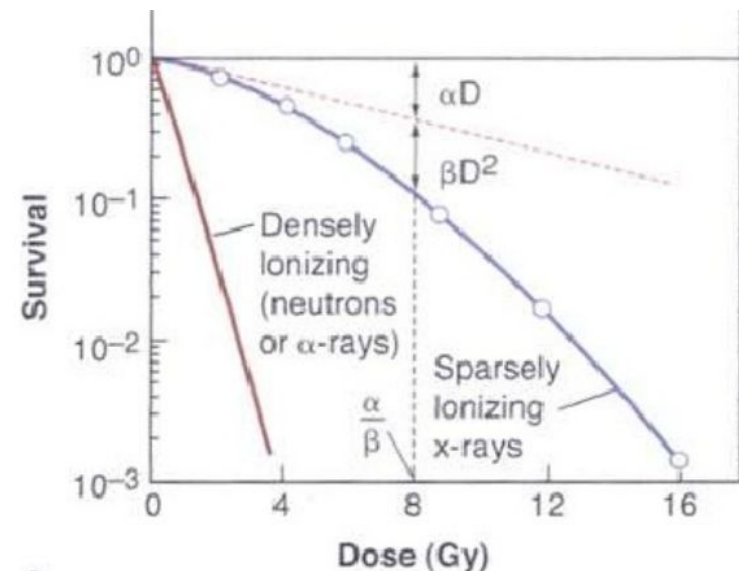
Primary proton's fragments are considered as secondary particles; each single spectra of those fragments is evaluated separately, considering its impact on the RBE.

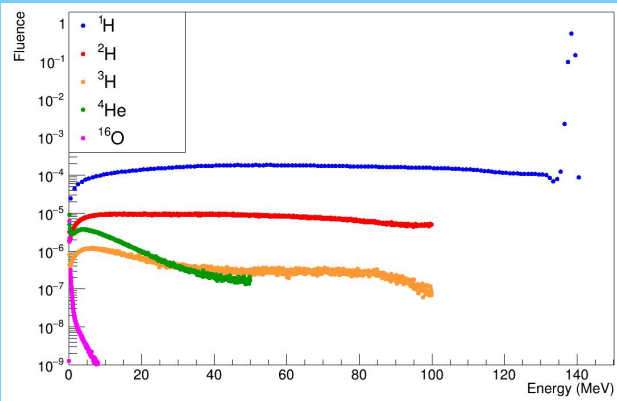
The total RBE is evaluated by using **Zeider & Rossi formula for mixed field irradiation** and **LEM IV model**:

$$\alpha = \frac{\sum_i \alpha_i D_i}{\sum_i D_i}$$

$$\sqrt{\beta} = \frac{\sum_i \sqrt{\beta_i} D_i}{\sum_i D_i}$$

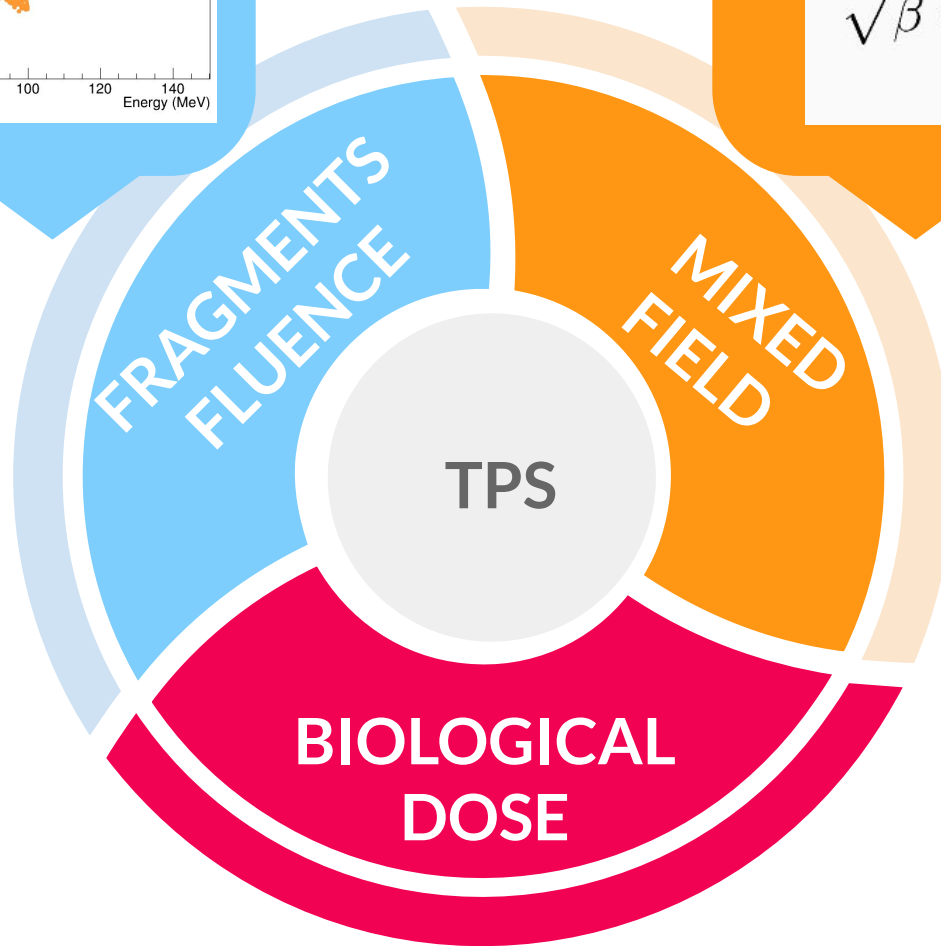
$$S(D) = \exp -(\alpha D + \beta D^2)$$





$$\alpha = \frac{\sum_i \alpha_i D_i}{\sum_i D_i}$$

$$\sqrt{\beta} = \frac{\sum_i \sqrt{\beta_i} D_i}{\sum_i D_i}$$



Biological dose comparison

Preliminary comparison between default TRiP98 data vs new TOPAS-base data

Tissue comparison:

Synthetic tissues obtained from LEM-IV model and characterized by $\alpha/\beta = 2\text{Gy}$ to represent a normal tissue.

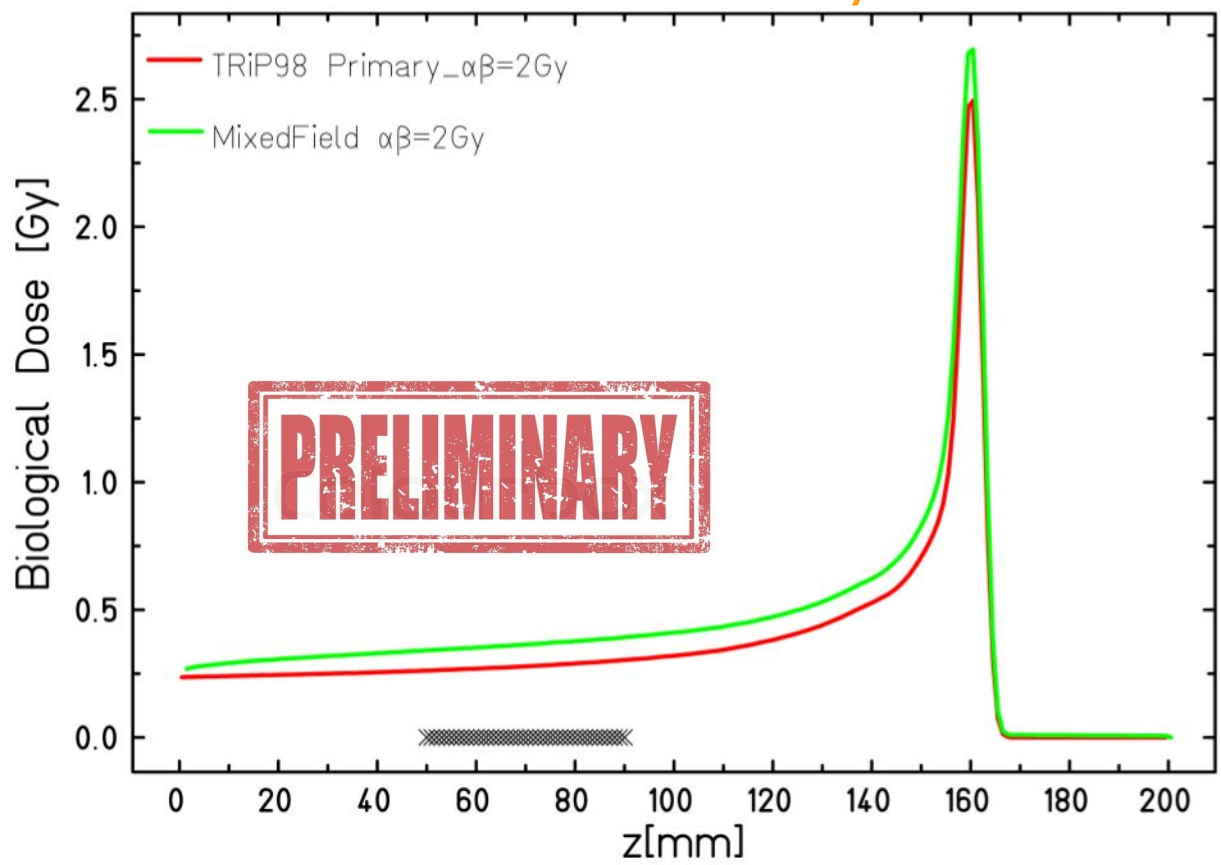
Particles contribution comparison:

- only primaries (Z=1 primary)
- only protons (primary and secondaries) (Z=1)
- All produced fragments

Biological dose comparison

Comparison between default TRiP98 data vs mixed field with TOPAS-base data

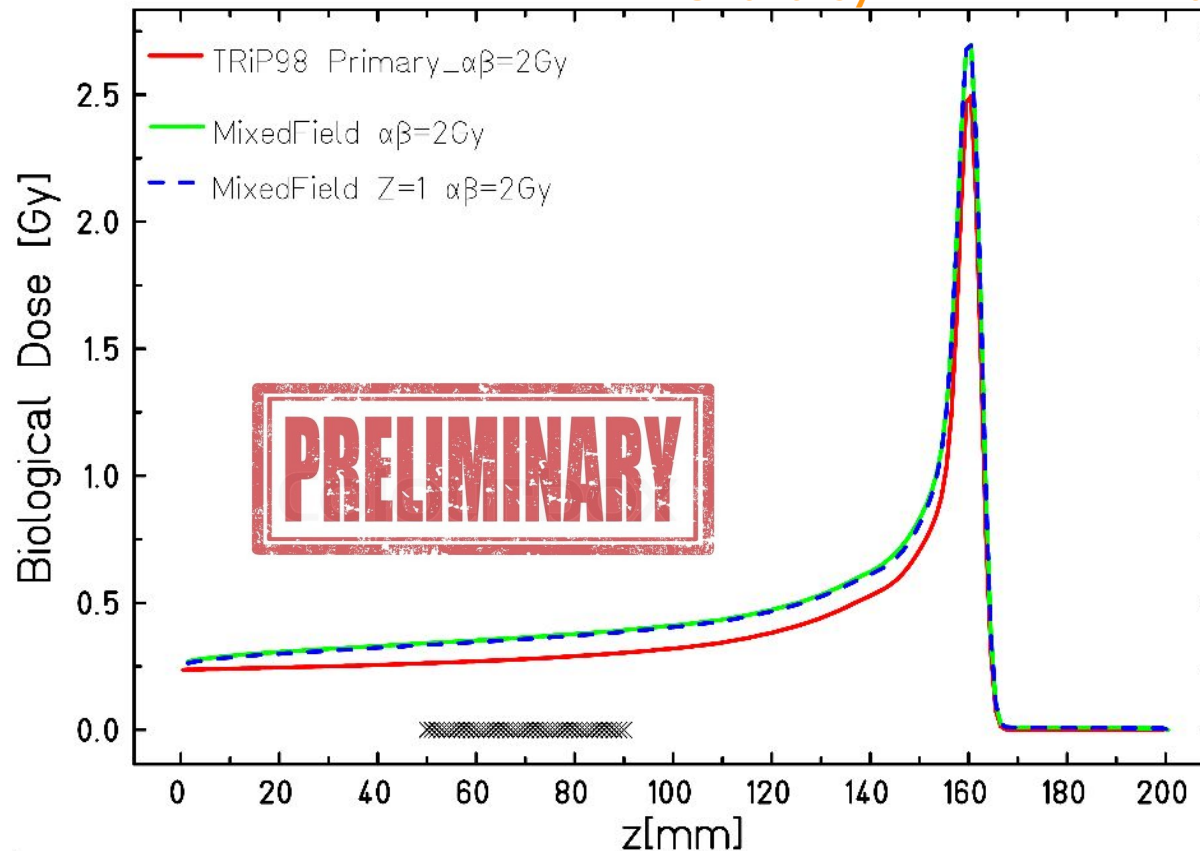
Courtesy of E. Bellinzona



Biological dose comparison

Comparison between default TRiP98 data vs mixed field with TOPAS-base data

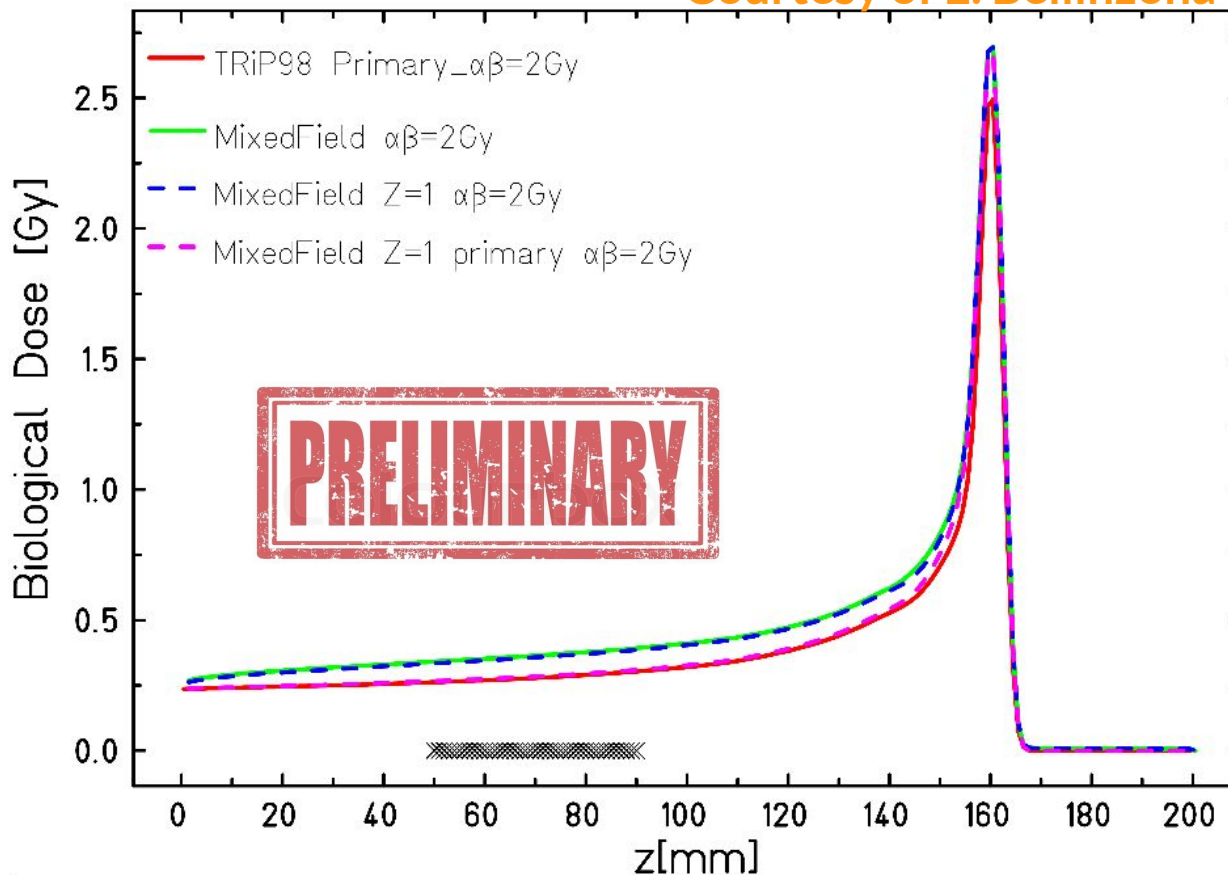
Courtesy of E. Bellinzona



Biological dose comparison

Comparison between default TRiP98 data vs mixed field with TOPAS-base data

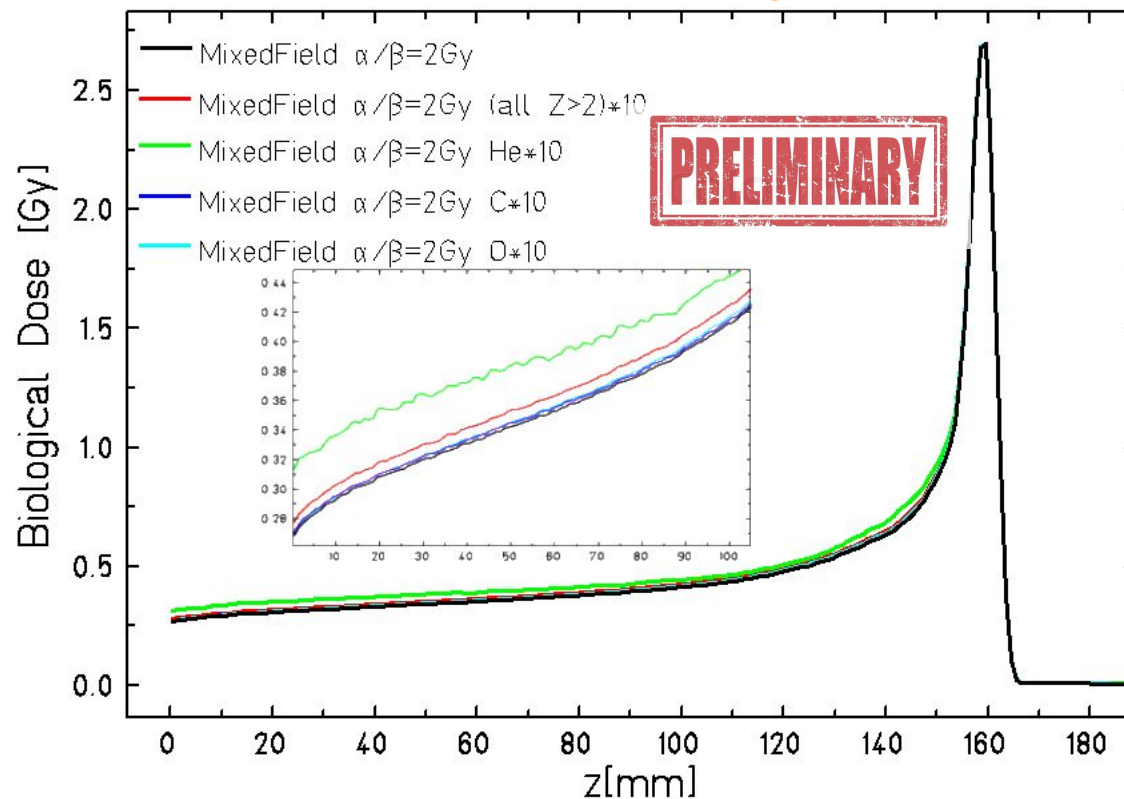
Courtesy of E. Bellinzona



Fragment impact

The fluence contribution of main fragments has been enhanced (by a factor 10) to analyse the dependence of species contribution on the biological dose.

Courtesy of E. Bellinzona



MoVe IT Future perspective

- Creation of fragments fluence database ($E_k, E_f, A, Z, z_{\text{depth}}$)
- Results validation with different MC simulations (FLUKA, GEANT4, TOPAS)
- Biological dose evaluation of SOPB with TRIP98, starting from MC databases, in water at two depths (entrance and distal region)

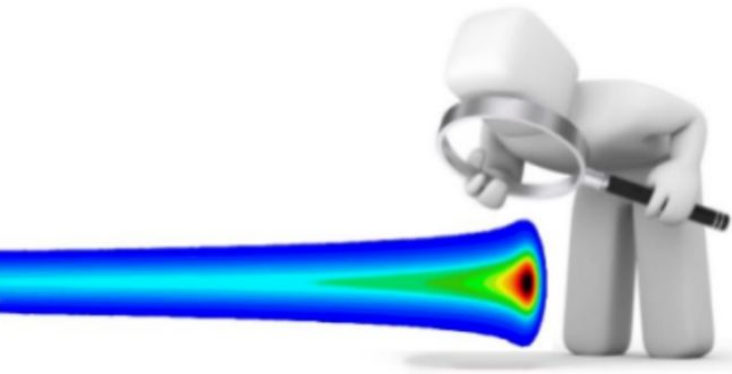


WAITING FOR FOOT EXPERIMENTAL DATA

(see poster #8:
FOOT FragmentatiOn Of Target)

Conclusions

Target fragmentation has been studied using *FLUKA MC code*.



- The energy spectra of fragments: $0 < E < 20$ MeV/u

LOW ENERGY and **HIGH Z**

↳ **ENHANCED RBE**

- The range of fragments is of the order of 10-100 μm **SMALL RANGE**

↳ **Relevant for normal tissues in the entrance region**

MoVe IT

Starting from these preliminary results, the next step is the creation of the fragments database (fluence, energy, Z, A) for the inclusion of target fragmentation in the TPS.

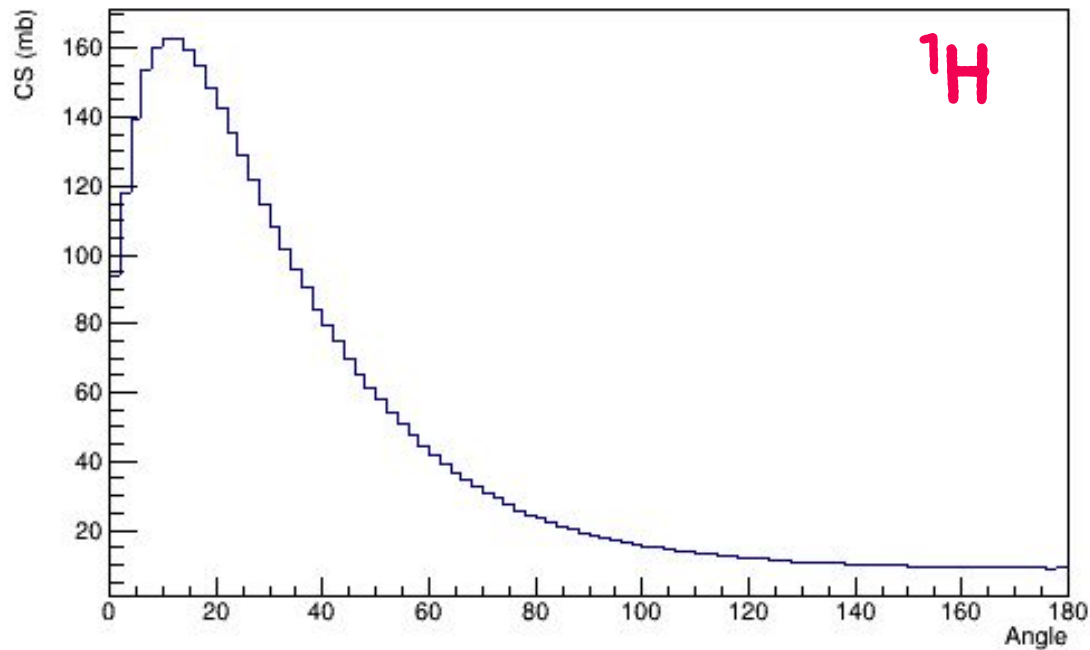
Two horizontal bars are positioned above the text. The left bar is light blue and dark blue. The right bar is orange and red.

**THANK
YOU!**

BACKUP SLIDE

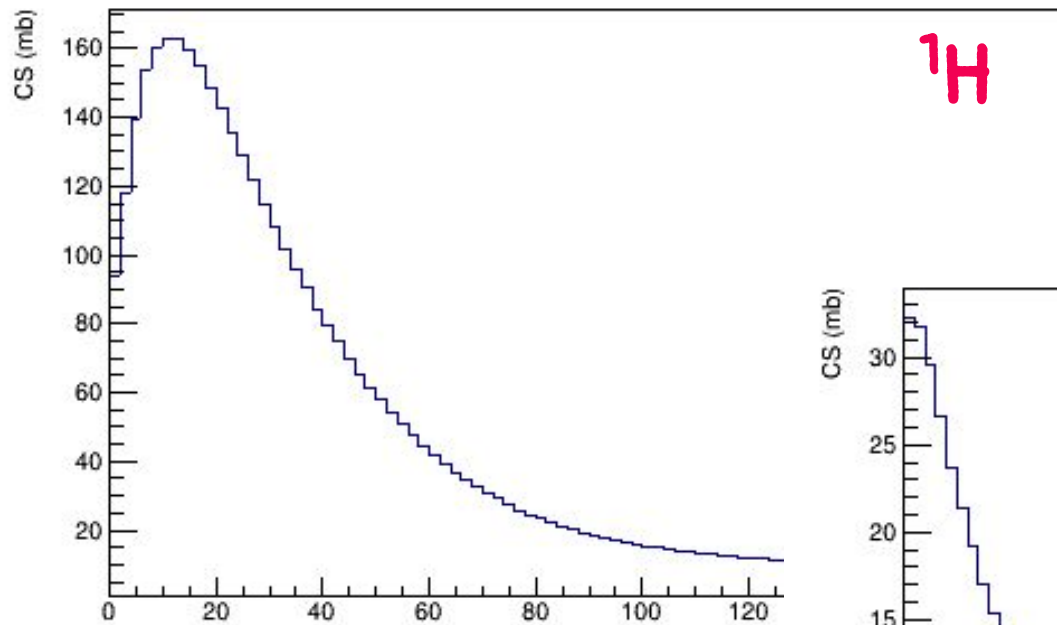
Angular distribution of fragments

Production CS of inelastic interaction induced by proton beam of 200 MeV in water.



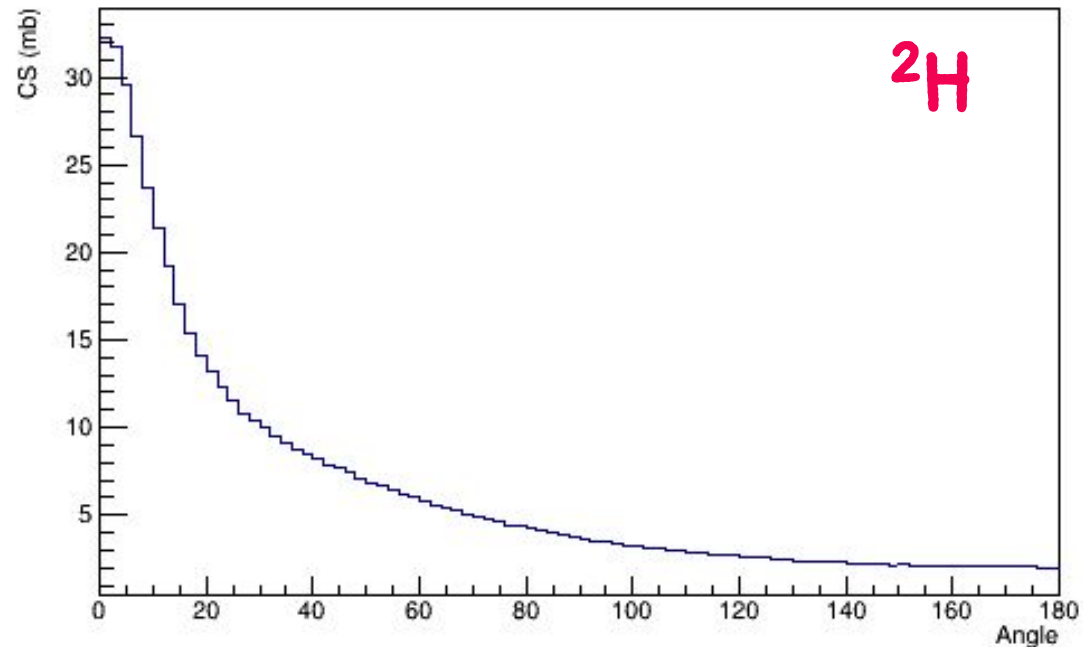
Angular distribution of fragments

Production CS of inelastic interaction induced by proton beam of 200 MeV in water.



^1H

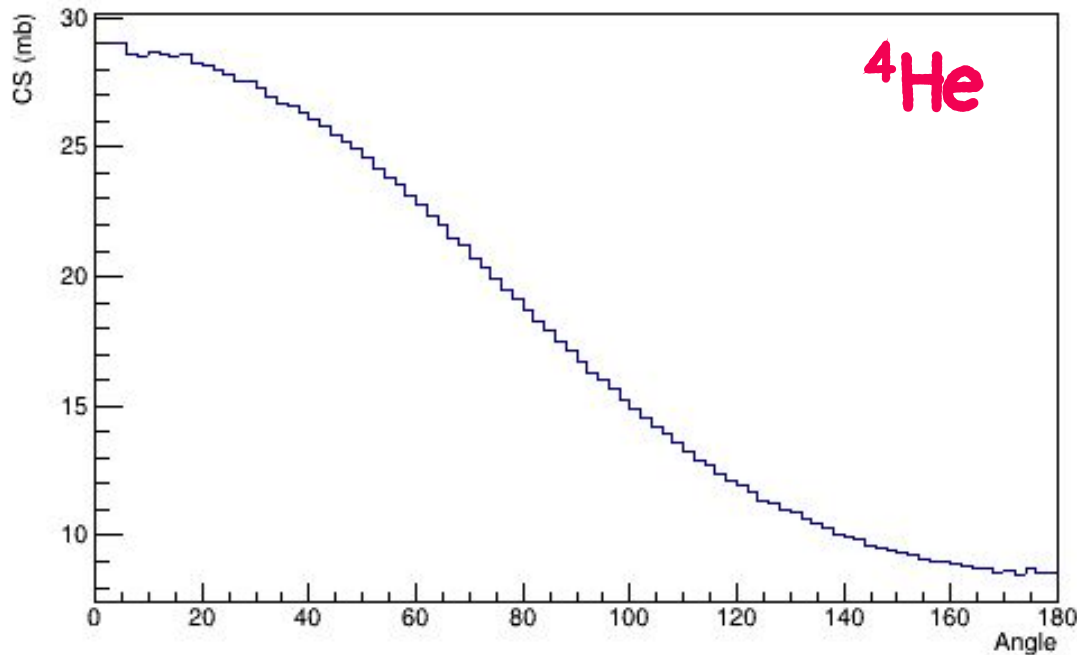
The emission of Hydrogen fragments is focused forward.



^2H

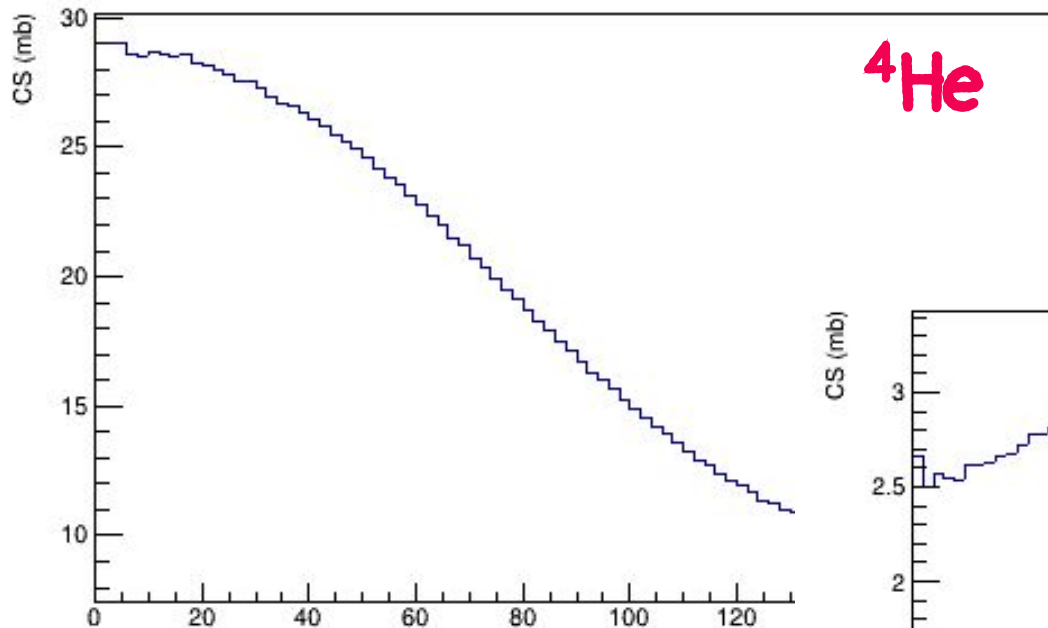
Angular distribution of fragments

Production CS of inelastic interaction induced by proton beam of 200 MeV in water.



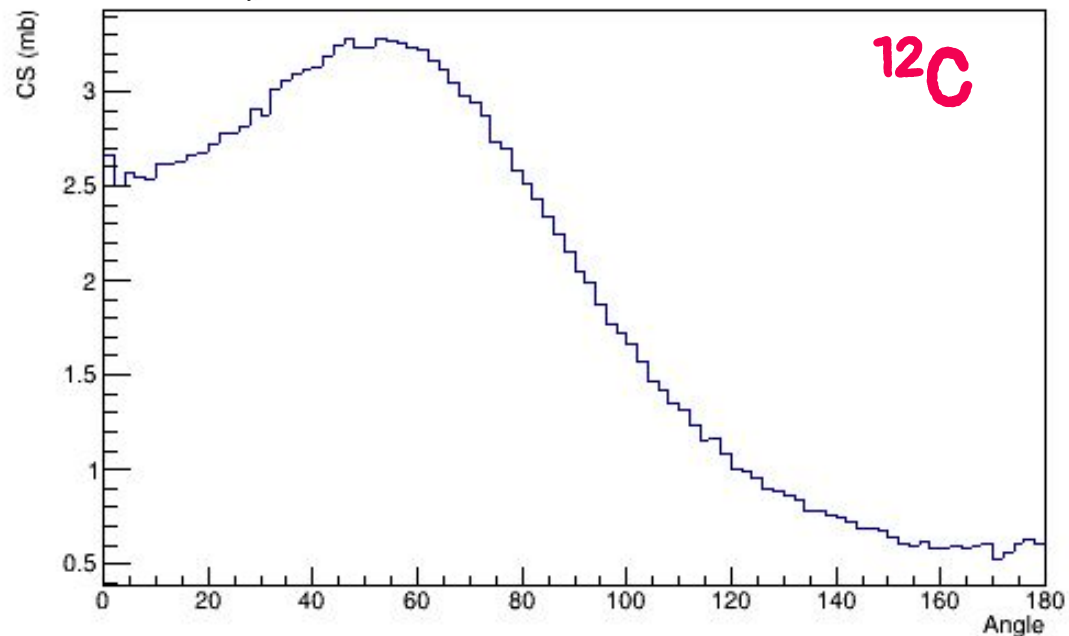
Angular distribution of fragments

Production CS of inelastic interaction induced by proton beam of 200 MeV in water.



^4He

For heavy fragments, the angular distribution present a mean value of 60 degree.



^{12}C

Range of fragments

The range distribution of main fragments produced by proton beam in water have been analyzed, considering different energies $E_p=40,80,120,160,200$ MeV/u.

The energy spectra of fragments are in a range $0 < E < 20$ MeV/u.

Starting from the energy distribution of fragments, **the range distribution** has been evaluated.

For heavier fragments: the mean range obtained from the distribution is similar to the value reported in Cancers.

For low Z fragments: difference between average range in analytical formula and MC distribution.

The table compares the range distribution of fragments produced by a proton beam in water, comparing results from FLUKA (red box) and Tommasino & Durante (purple box). The columns represent the range R_p (in μm) and the range R_{TD} (in μm). The rows list various fragments from ^1H to ^{16}O . Arrows point from the labels 'FLUKA' and 'Tommasino & Durante' to their respective columns.

Fragment	R_p (μm)	R_{TD} (μm)
^1H	1392	-
^2H	2117	68.9
^3H	1383	-
^3He	465.7	38.8
^4He	218.3	48.5
^6He	229.3	-
^6Li	75.7	26.7
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^{11}C	8.1	7
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^{15}O	2.8	2.3
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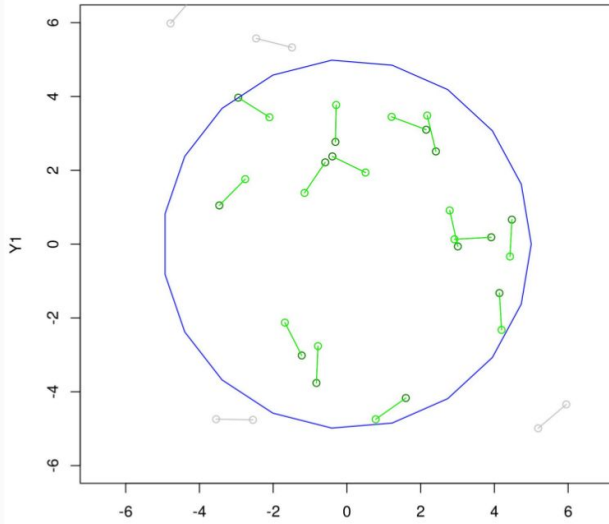
MC more complete description of physical processes

Range is more reliable

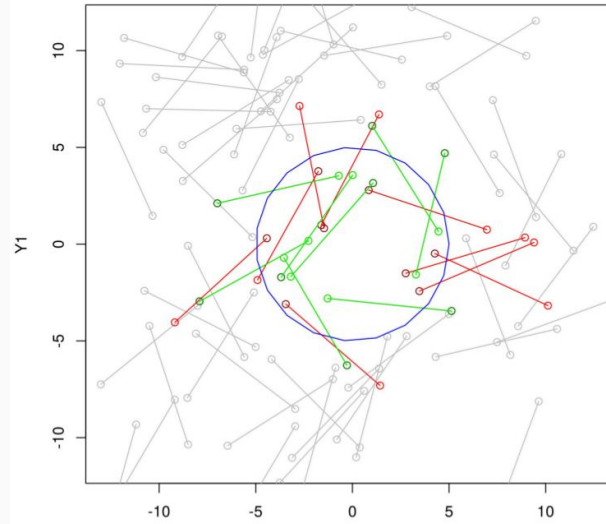
Fragments Range

Courtesy of A. Attili

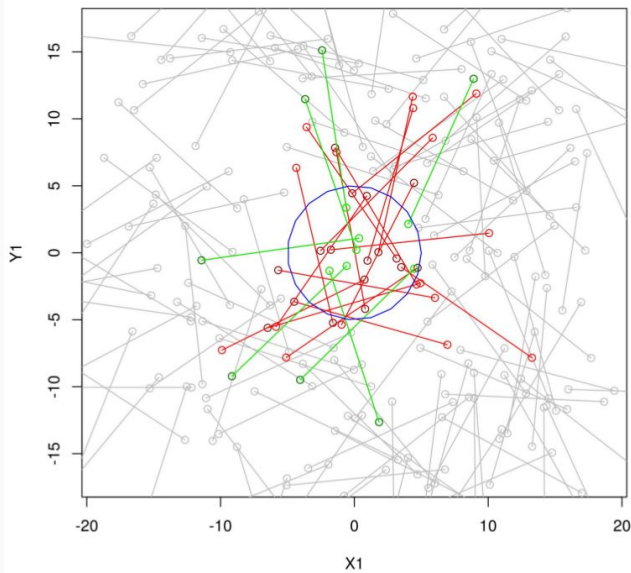
RN = 5, R = 1



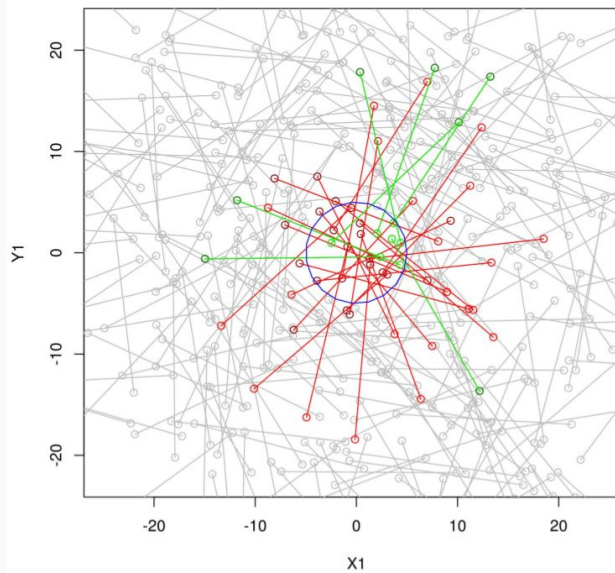
RN = 5, R = 6.44



RN = 5, R = 11.89



RN = 5, R = 17.33



Fragments Range

Courtesy of A. Attili

SRIM evaluation of projected range as a function of energy:

Low Z fragments
 $R >$ cell nucleus

Heavier fragments
 $R <$ cell nucleus
Colocalization effects

