

Development and implementation of new Geant4 QMD model and its validation

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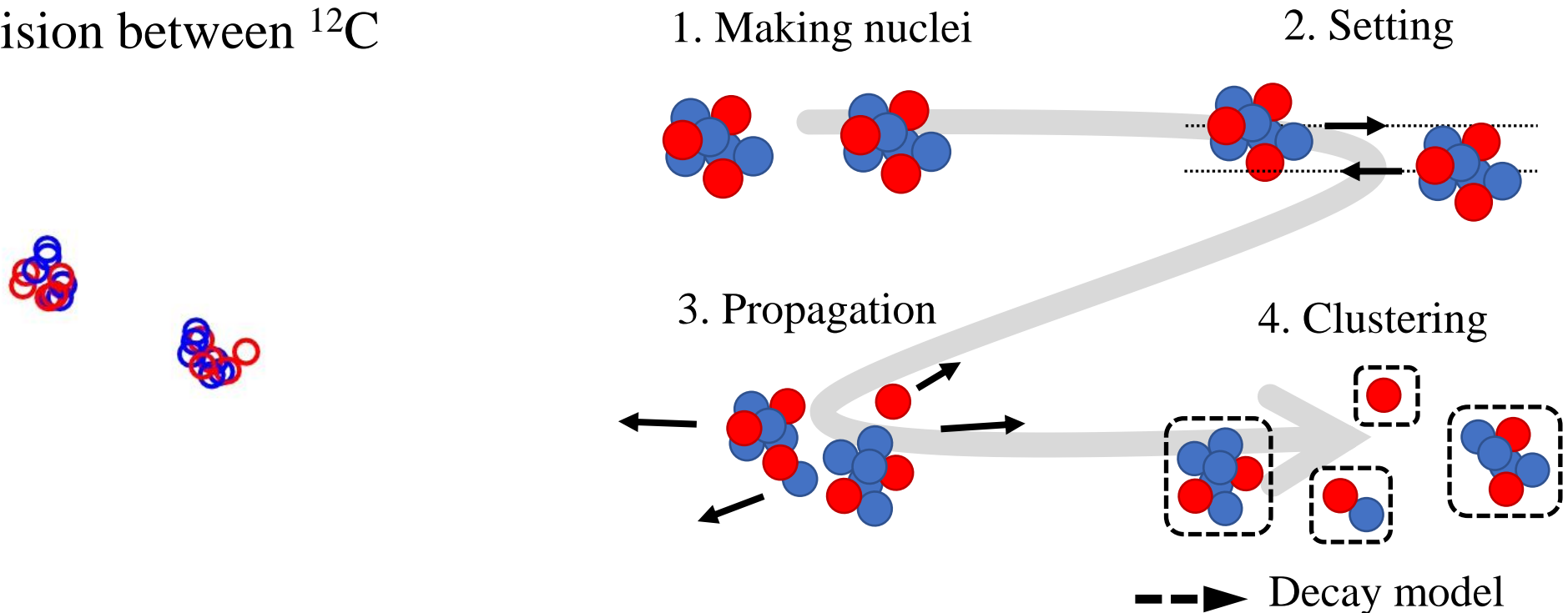
28th Geant4 Collaboration Meeting at Hokkaido University,
Sep. 26th, 2023

Quantum Molecular Dynamics (QMD)

Introduction

The QMD model is a quantum extension of the classical molecular-dynamics model and can describe hadronic processes, especially **inelastic processes**, in Geant4.

Collision between ^{12}C



Development of more accurate QMD model for hadron therapy

Introduction

IOP Publishing *Phys. Med. Biol.* 67 (2022) 225001 <https://doi.org/10.1088/1361-6560/ac9a9a>

Physics in Medicine & Biology

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Institute of Physics and
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PAPER

Development of a more accurate Geant4 quantum molecular dynamics model for hadron therapy

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11 July 2022

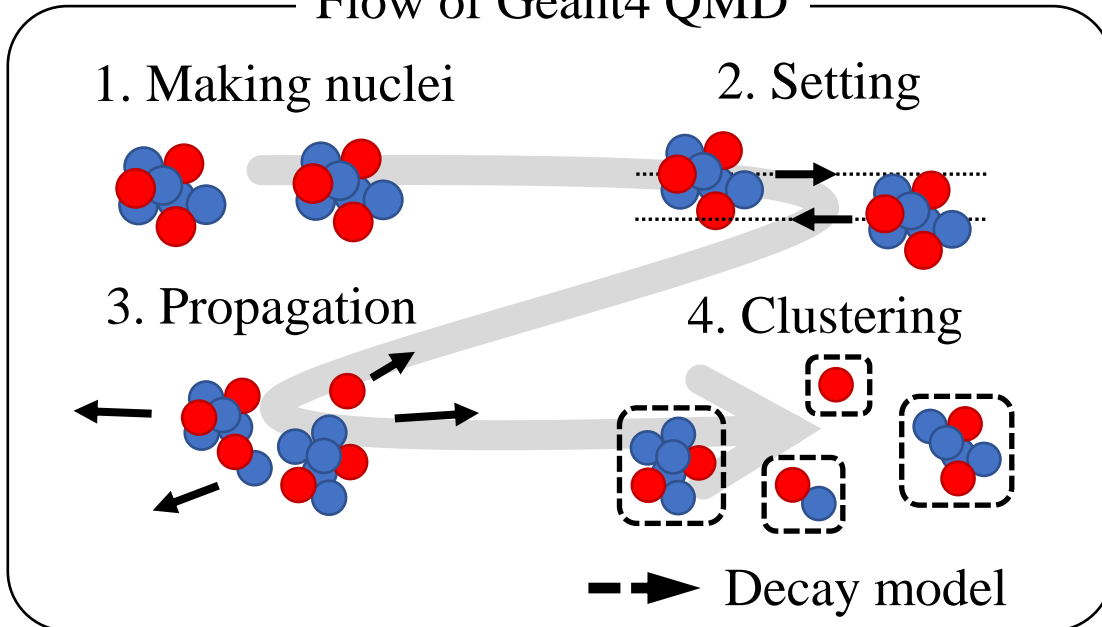
REVISED
6 October 2022

ACCEPTED FOR PUBLICATION
14 October 2022

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Sato, Y.-H. *et al.* Development of a more accurate Geant4 quantum molecular dynamics model for hadron therapy. *Phys. Med. Biol.* **67**, (2022)

Flow of Geant4 QMD

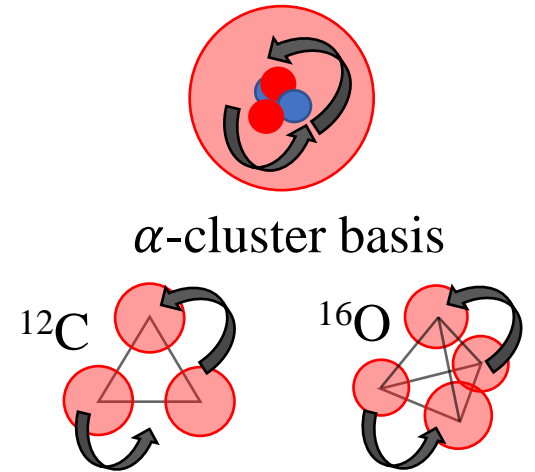


Three improvements

1. Modern interaction

	JQMD ^b (G4QMD)	SLy4 ^c	SkM ^{*c}	SIII ^c
A [MeV]	-219.4	-297.82	-318	-122.921
B [MeV]	165.3	219.21	249.5	55.343
g ₀ [MeV fm ²]	—	24.569	21.86	18.286
g _τ [MeV]	—	9.70	5.9357	6.439
C _s [MeV]	25	32	32	32
κ _s [fm ²]	—	0.08	0.08	0.08
γ	4/3	7/6	7/6	2
η ^a	—	5/3	5/3	5/3
ρ ₀ [fm ⁻³] ^a	0.168	0.160	0.165	0.1452
BE [MeV] ^a	-16.00	-15.97	-15.77	-15.83
K ₀ [MeV] ^a	237.8	230.2	216.8	355.9

2. α-cluster structure



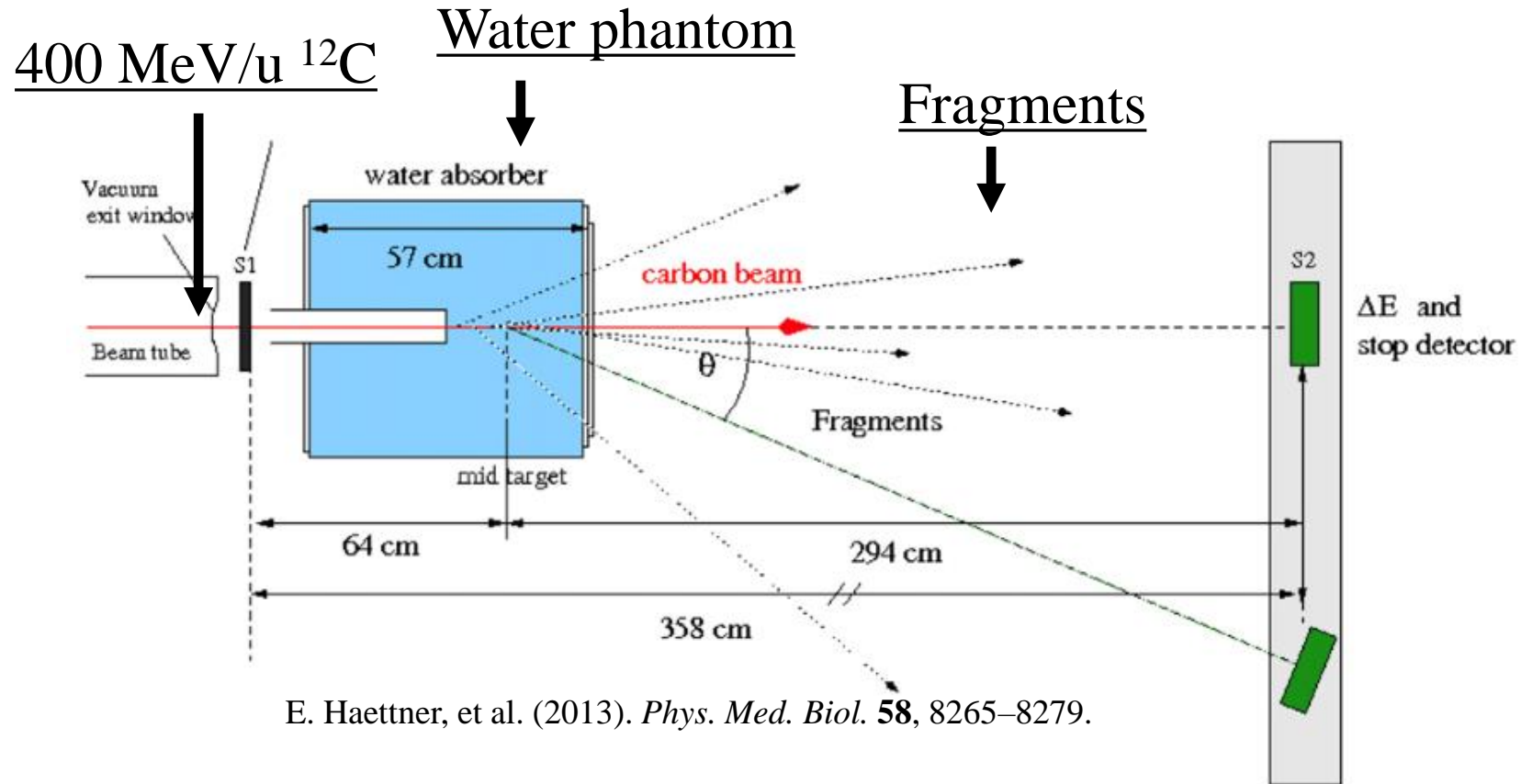
3. QMD model Parameter optimization

$$\varphi_i(\mathbf{r}) \equiv \frac{1}{(2\pi L)^{3/4}} \exp\left(-\frac{(\mathbf{r} - \mathbf{r}_i)^2}{4L} + \frac{i}{\hbar} \mathbf{r} \cdot \mathbf{p}_i\right)$$

$|\mathbf{r}_{ij}^2| \leq R^2$

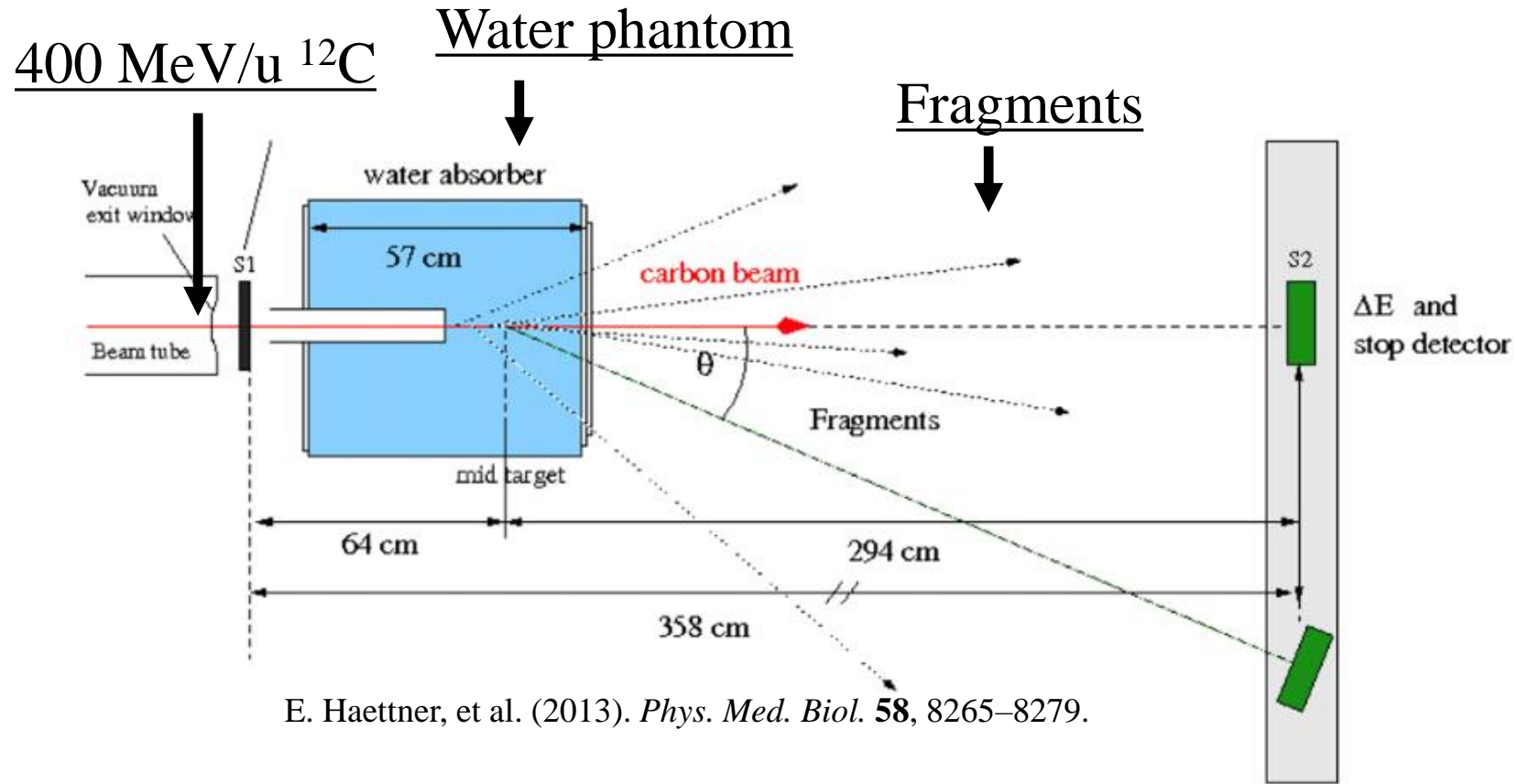
Water Phantom

Materials and Methods



Water Phantom

Materials and Methods



Light Ion QMD (LIQMD)

Fragment

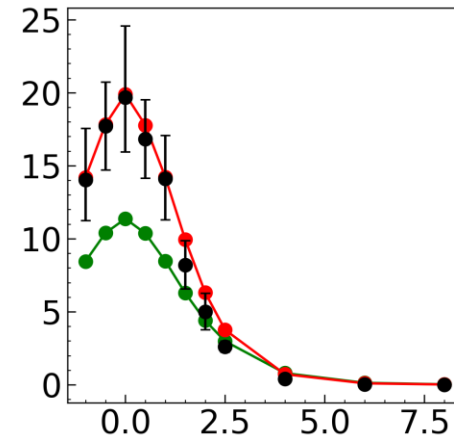
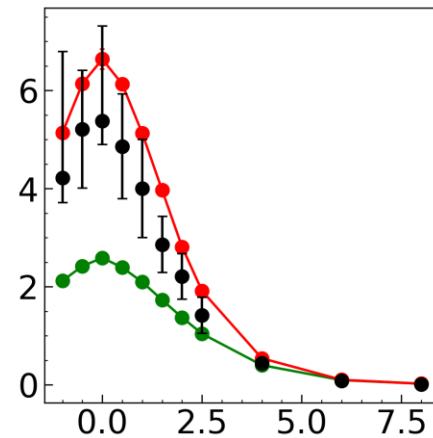
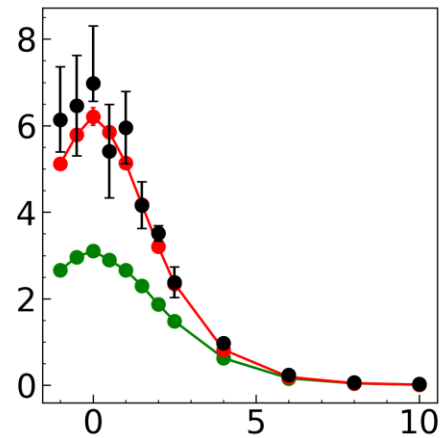
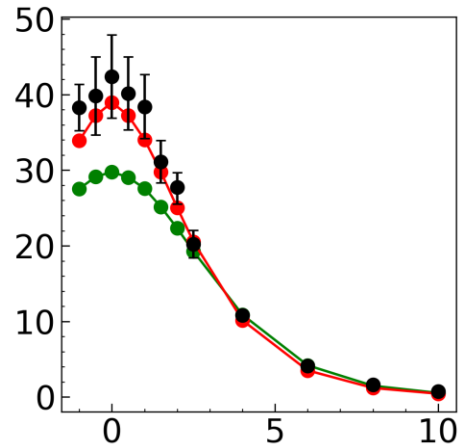
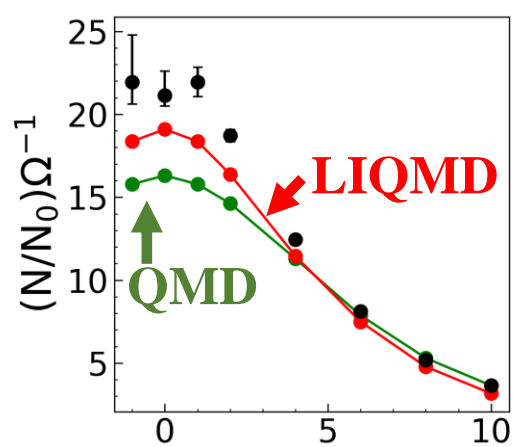
H

He

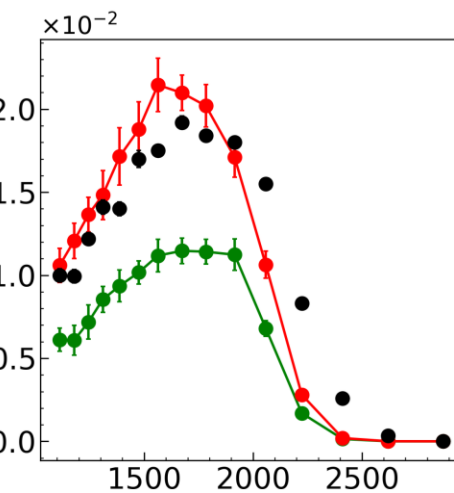
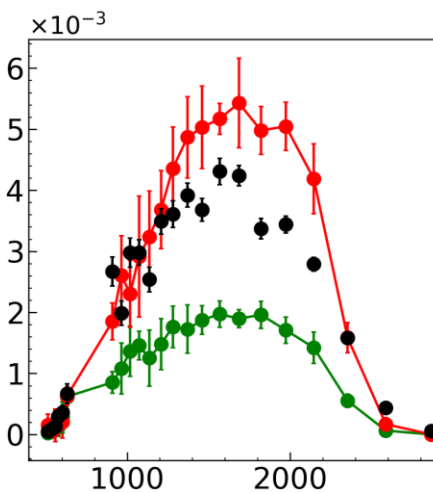
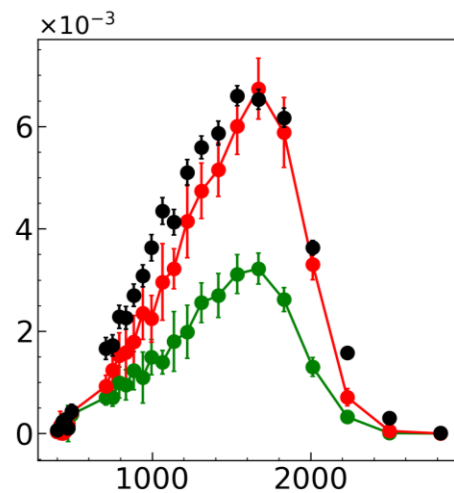
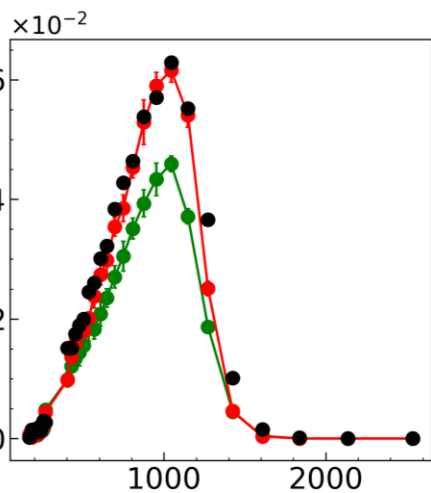
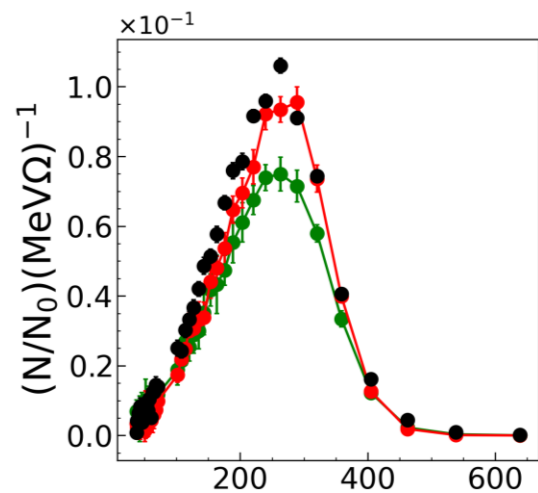
Li

Be

B



Angle (deg)



Energy (MeV)

Validation

- Water phantom with 400 MeV/u ^{12}C beam [1]
- Thin target with 95 MeV/u ^{12}C beam [2]
- Dose-Depth curve of water phantom [3]
- Positron-emitting nucleus yield [3]

[1] E. Haettner, et al. (2013). *Phys. Med. Biol.* **58**, 8265–8279.

[2] Dudouet, J. et al. (2013). *Physical Review C*, 88(2), 024606.

[3] A. Chacon, et. al.(2019), *Phys. Med. Biol.* **64**, 155014.

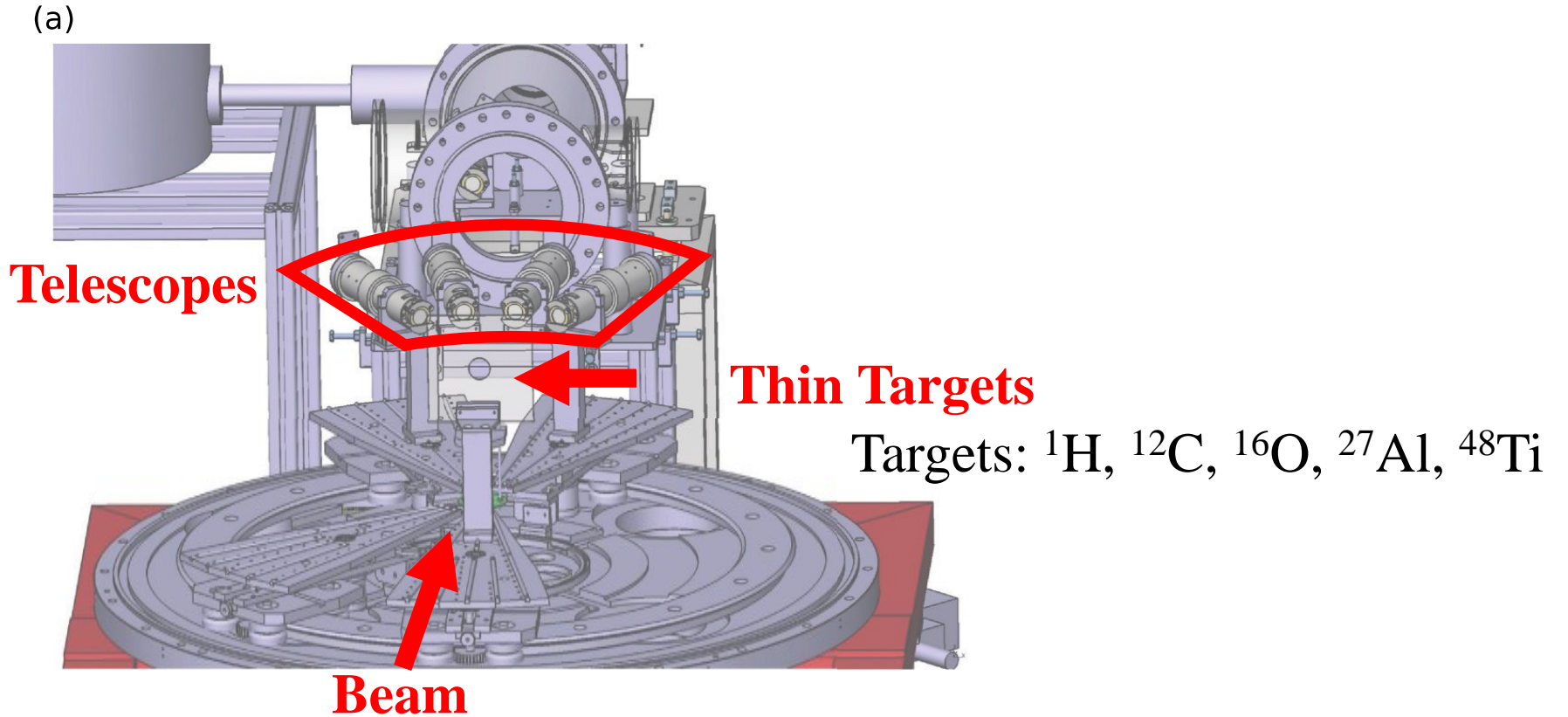
Validation

- Water phantom with 400 MeV/u ^{12}C beam
- **Thin target with 95 MeV/u ^{12}C beam**
- Dose-Depth curve of water phantom
- Positron-emitting nucleus yield

Thin Target

Materials and Methods

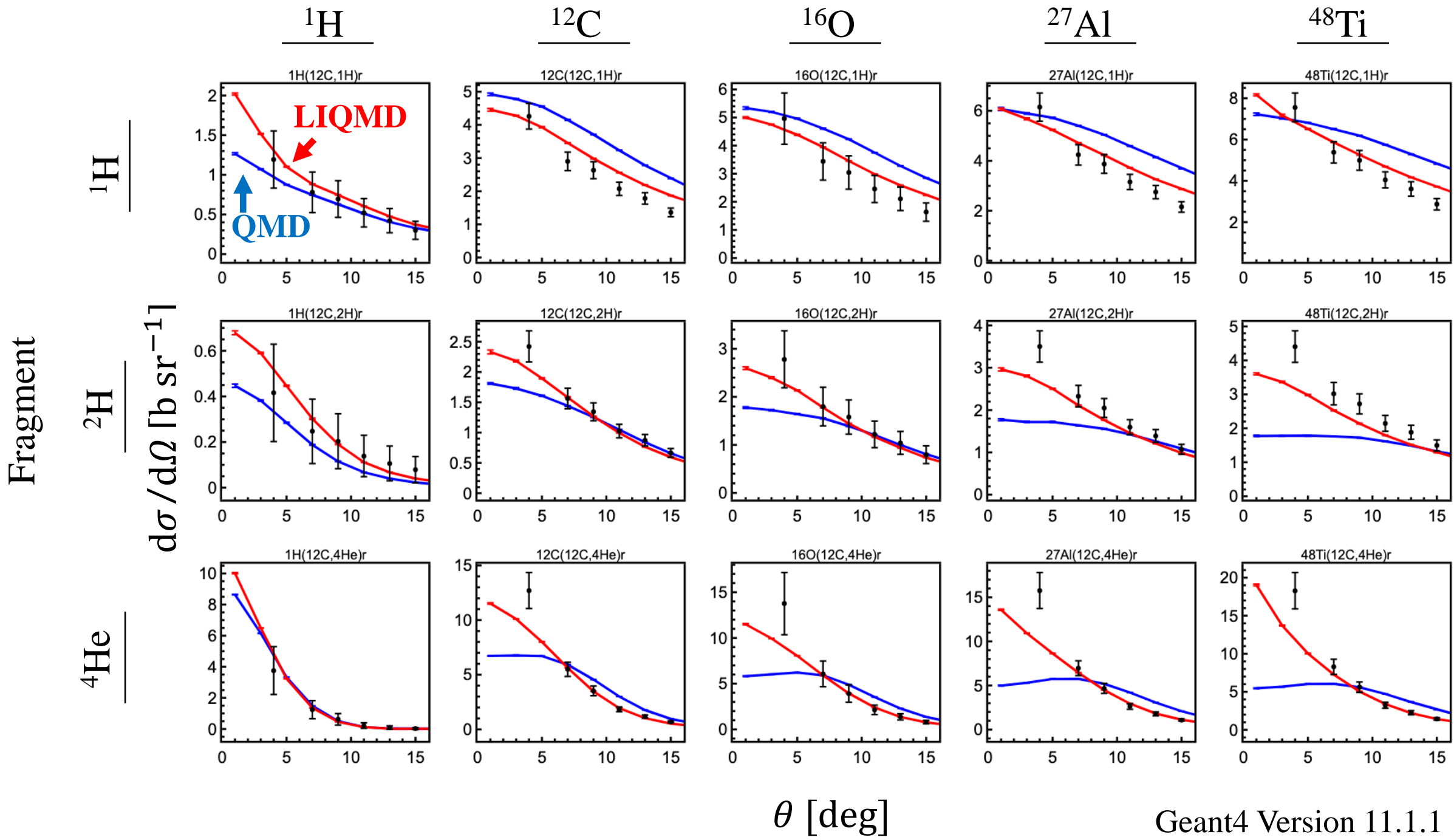
Fragments: $1,2,3\text{H}$, $3,4,6\text{He}$, $6,7\text{Li}$, $7,9,10\text{Be}$, $8,10,11\text{B}$, $10,11,12\text{C}$

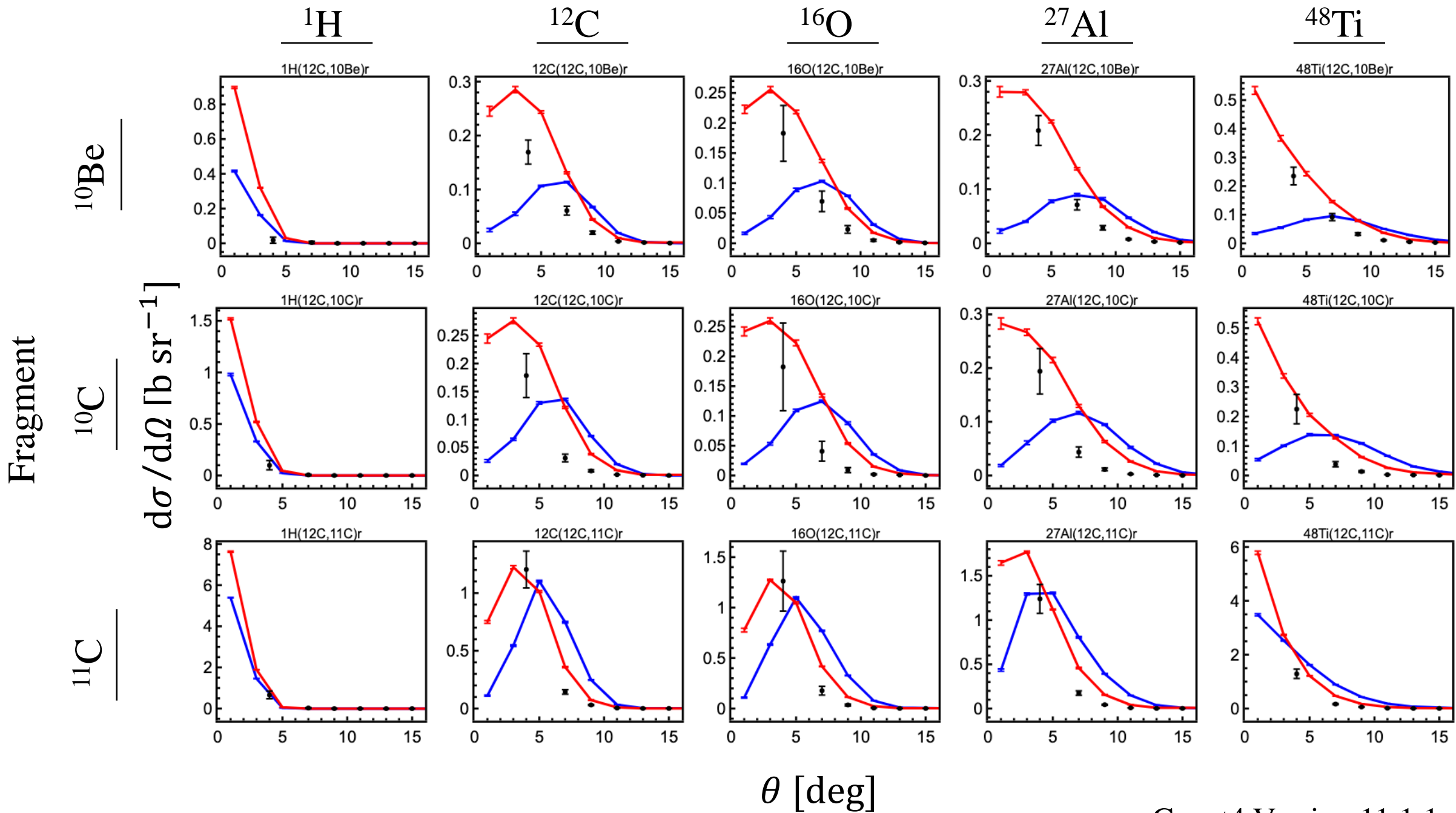


Beam: 12C , $94.6 \pm 0.09 \text{ MeV/A}$

Dudouet, J. et al.. (2013). *Physical Review C*, 88(2), 024606.

E600 experiment at the Grand Accelérateur National d'Ions Lourds (GANIL) facility.



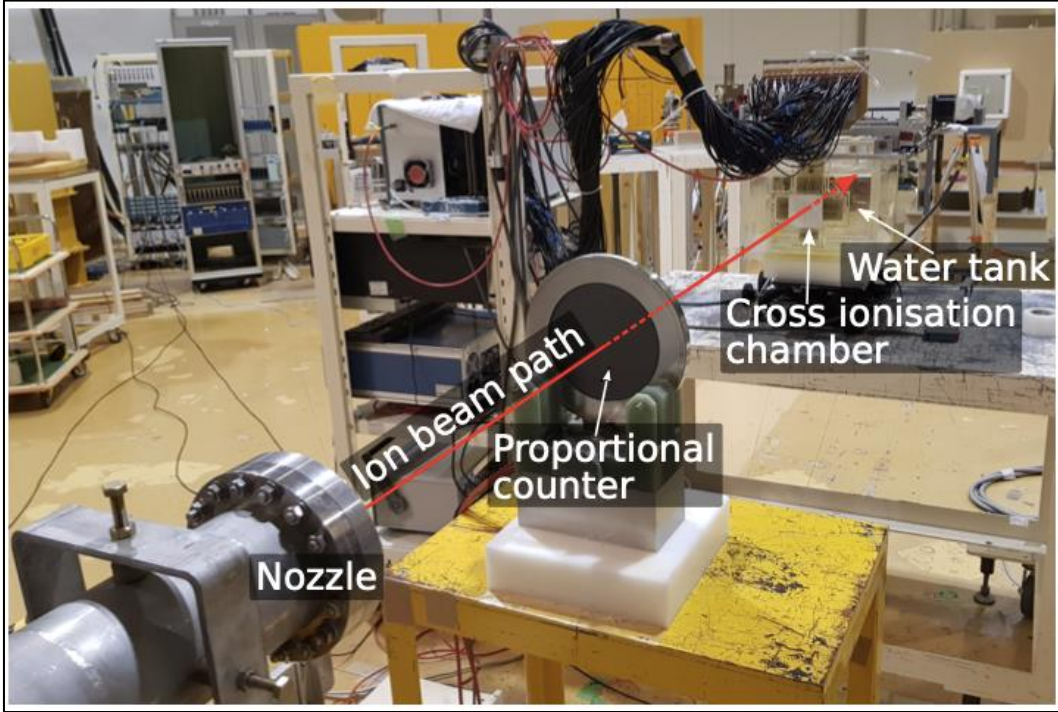


Validation

- Water phantom with 400 MeV/u ^{12}C beam
- Thin target with 95 MeV/u ^{12}C beam
- Dose-Depth curve of water phantom
- Positron-emitting nucleus yield

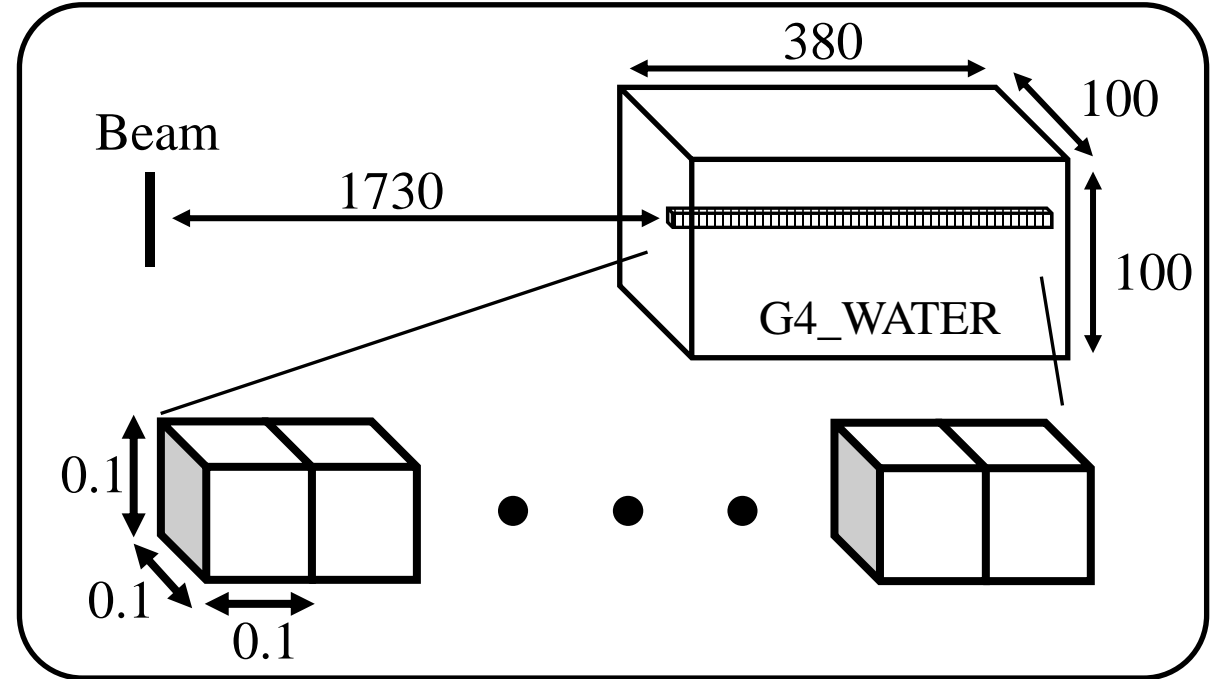
Depth-Dose Curve

Materials and Methods



[1] A. Chacon, et. al., *Phys. Med. Biol.* **64**, 155014 (2019).

Simulation geometry (Unit : mm)

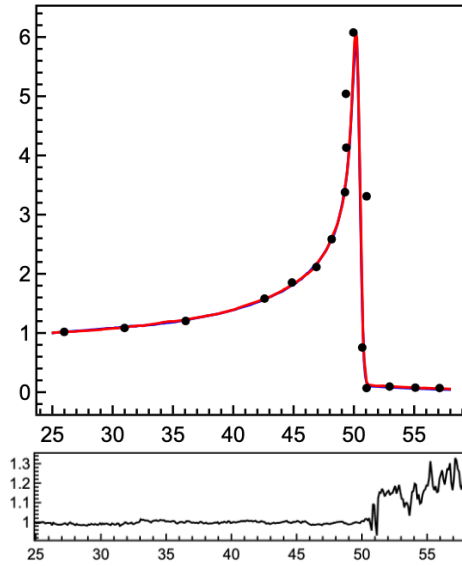


- ❑ Recording the **energy deposition** to each voxel on the beam axis by Command-based scoring.

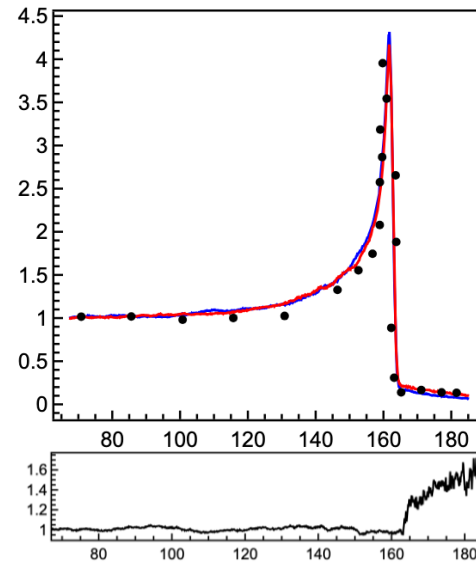
$$N_0 = 10^8$$

Normalized Dose
&
Dose Ratio (LIQMD/QMD)

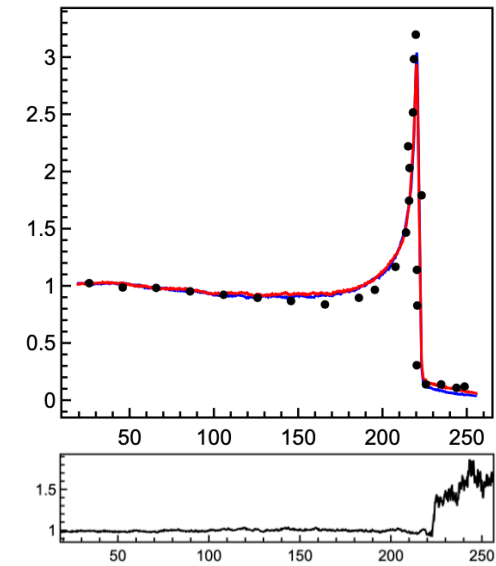
^{12}C , 148.5 MeV/u



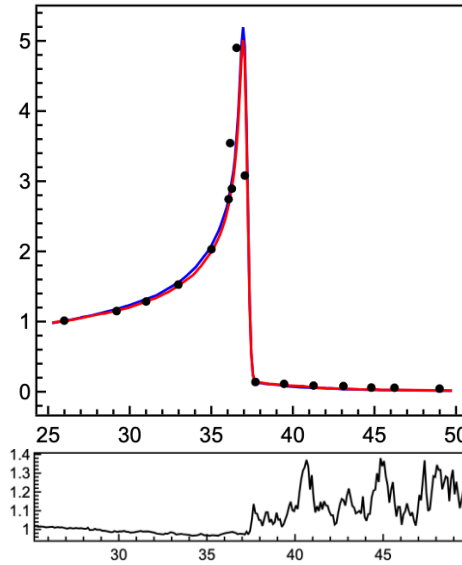
^{12}C , 290.5 MeV/u



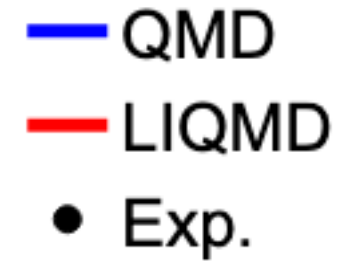
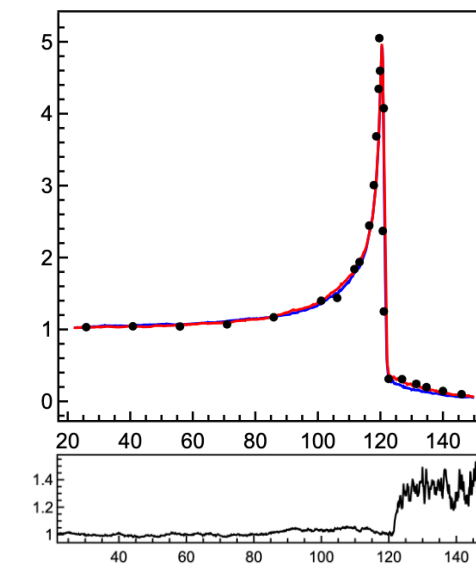
^{12}C , 350 MeV/u



^{16}O , 148 MeV/u



^{16}O , 290 MeV/u



Geant4 Version 11.1.1

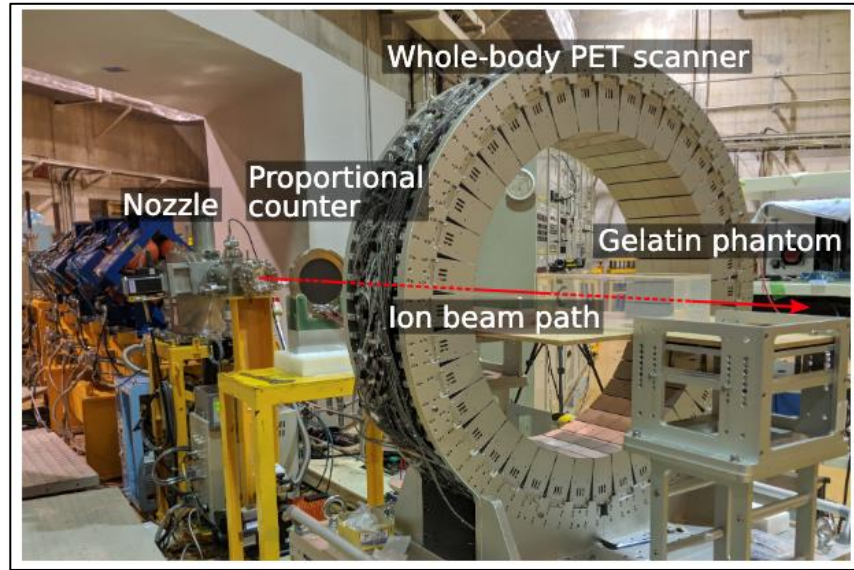
Depth in water [mm]

Validation

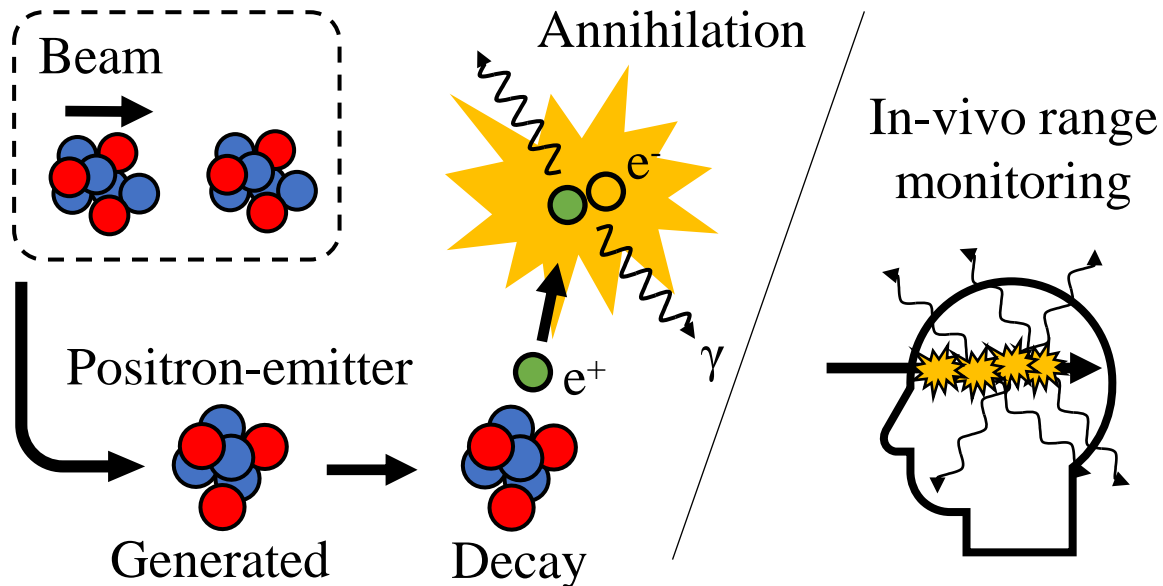
- Water phantom with 400 MeV/u ^{12}C beam
- Thin target with 95 MeV/u ^{12}C beam
- Dose-Depth curve of water phantom
- Positron-emitting nucleus yield

Positron-Emitting Nuclide Yield

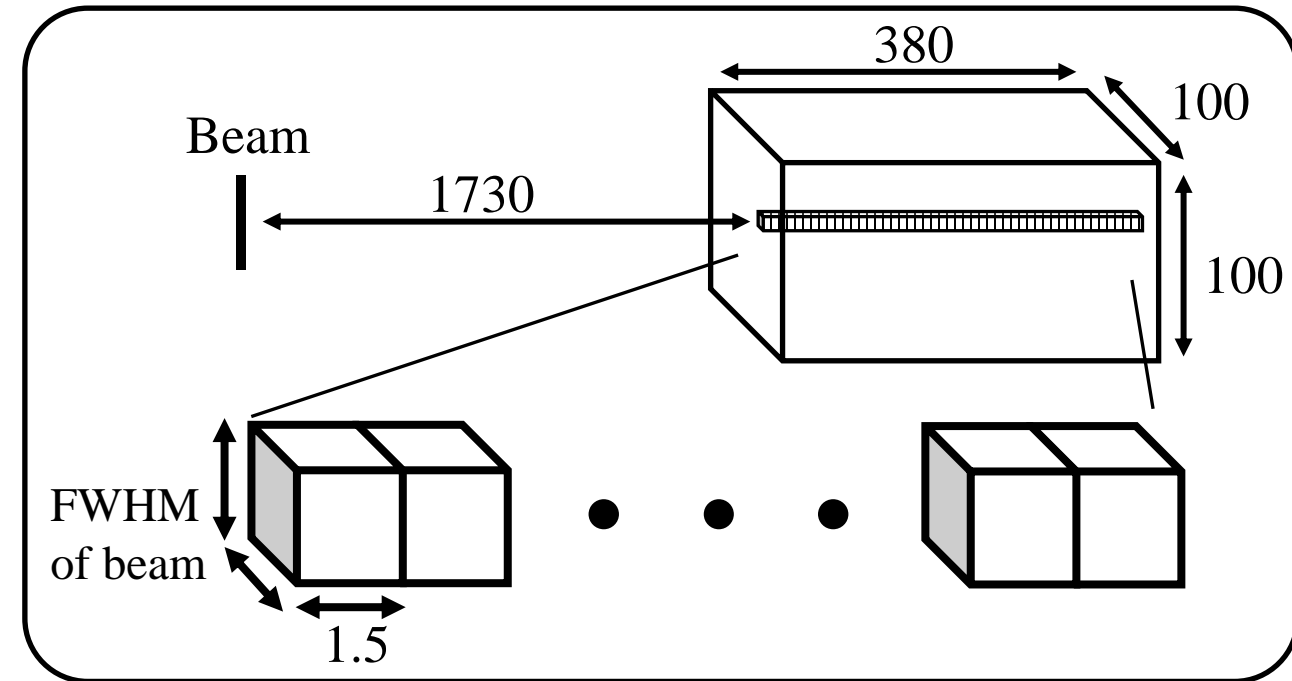
Materials and Methods



A. Chacon, et. al., *Phys. Med. Biol.* **64**, 155014 (2019).



Simulation and analysis geometry (Unit : mm)



- Calculating the **number of positron-emitting nuclide** in each voxel on the beam axis at the initial.

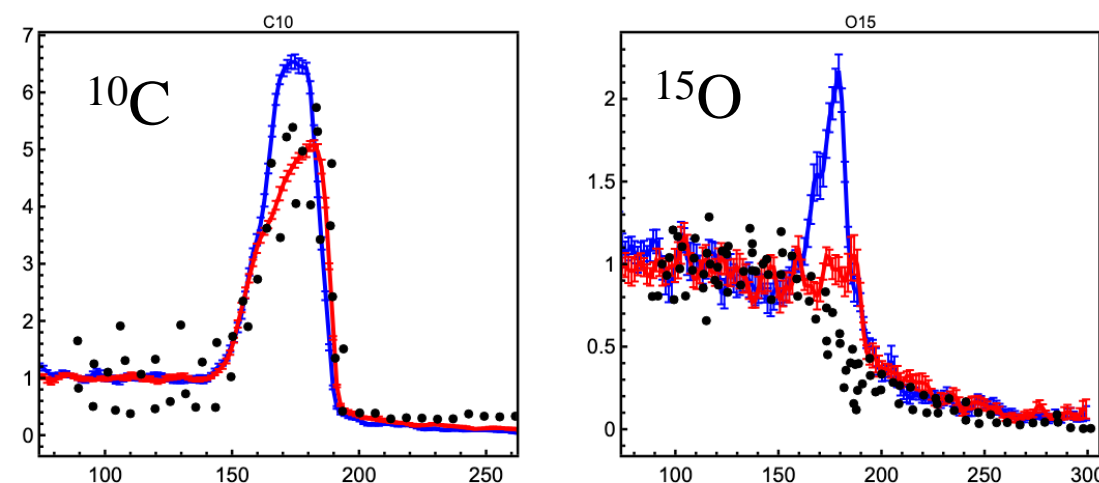
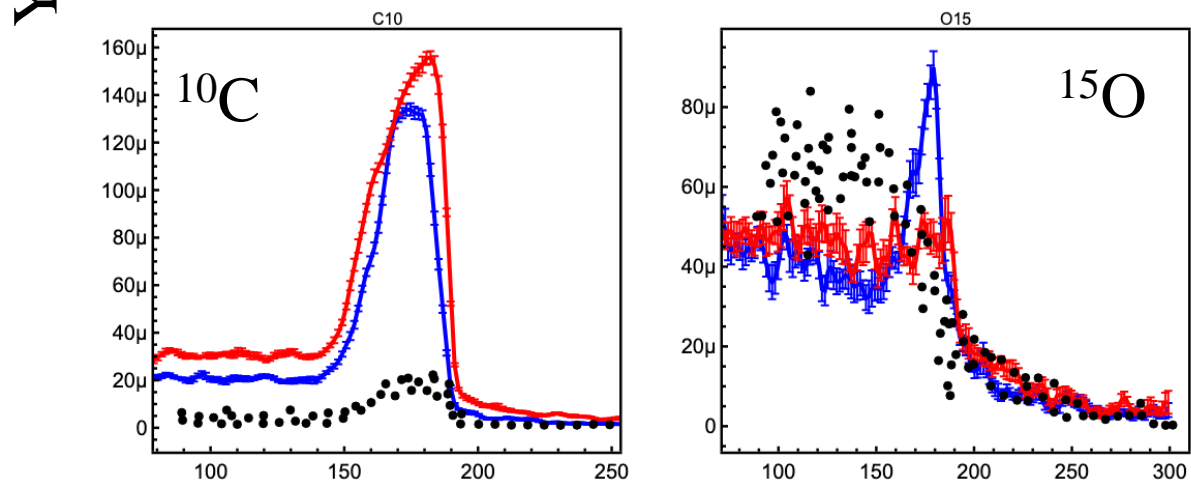
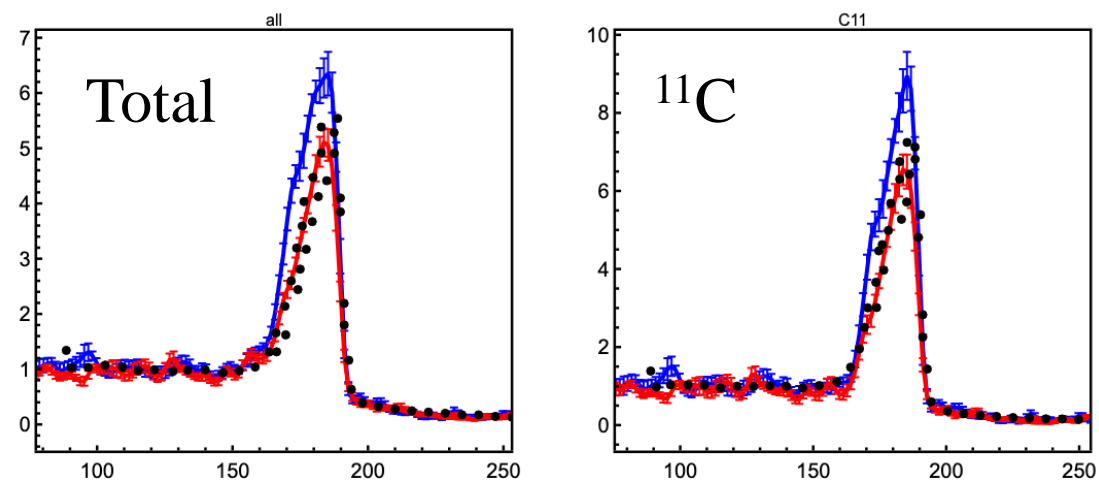
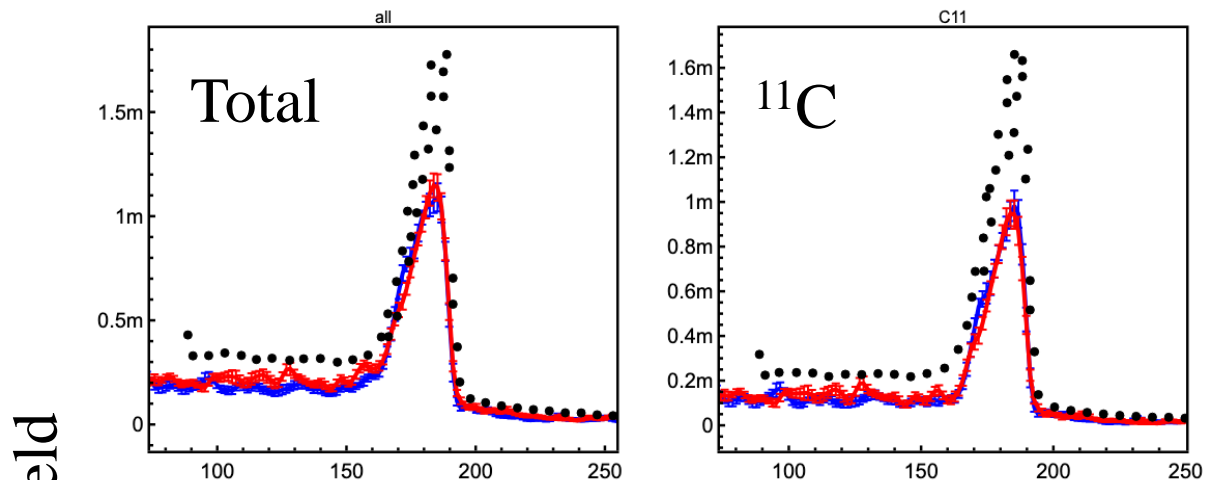
— QMD — LIQMD • Exp.

^{12}C , 350MeV/u, PMMA

$N_0 = 10^8$

10 repetitions

Normalized (Focus on shape)



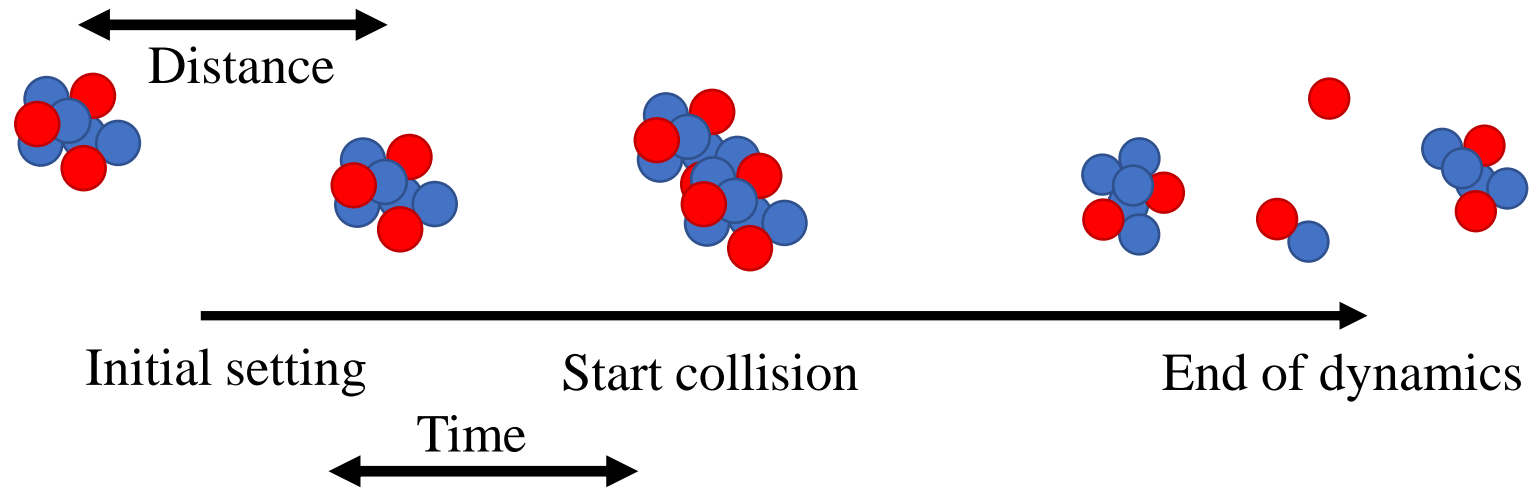
Limitation in LIQMD

- Energy range
- Mass range

Energy Range Recommended in LIQMD

Materials and Methods

Projectile and target should be separated by at least about b -value.

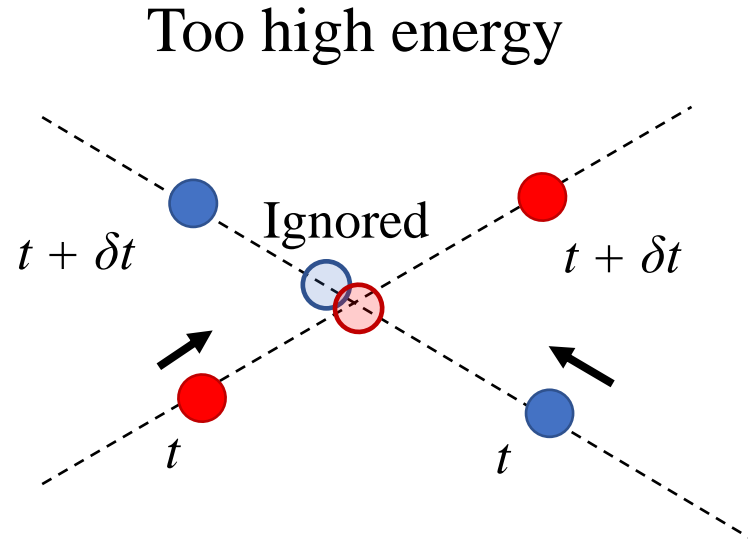


If the speed is slow (Low kinetic energy per nucleon), it will take a long time to collide, leaving not enough time for simulation after the collision.

It is better to have at least about 30 MeV/u.

Energy Range Recommended in LIQMD

Materials and Methods



500 MeV/u gives 1.5 fm gap in time step $\delta t = 2 \text{ fm}/c$.

Due to this large gap, collisions that should occur may be ignored.

Energies below about 500 MeV/u are recommended.

Mass Range Recommended in LIQMD

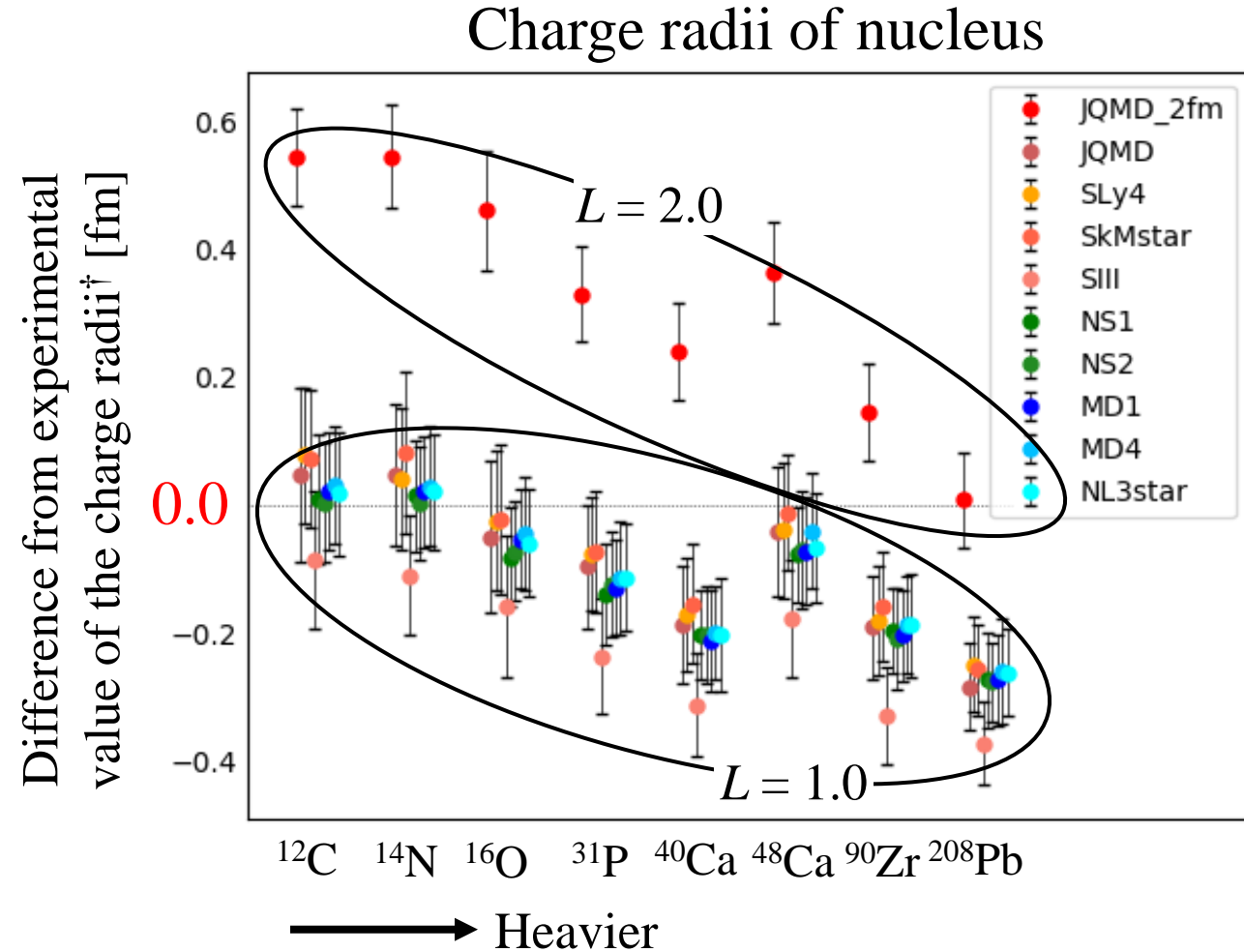
Materials and Methods

Wave function of nucleon

$$\varphi_i(\mathbf{r}) \equiv \frac{1}{(2\pi L)^{3/4}} \exp\left(-\frac{(\mathbf{r} - \mathbf{r}_i)^2}{4L} + \frac{i}{\hbar} \mathbf{r} \cdot \mathbf{p}_i\right)$$

$L = 2.0$ [fm] for QMD

$L = 1.26$ [fm] for LIQMD



†: I. Angeli, K.P. Marinova, Atomic Data and Nuclear Data Tables, 99, 2013, 69-95.

Mass Range Recommended in LIQMD

Materials and Methods

Wave function of nucleon

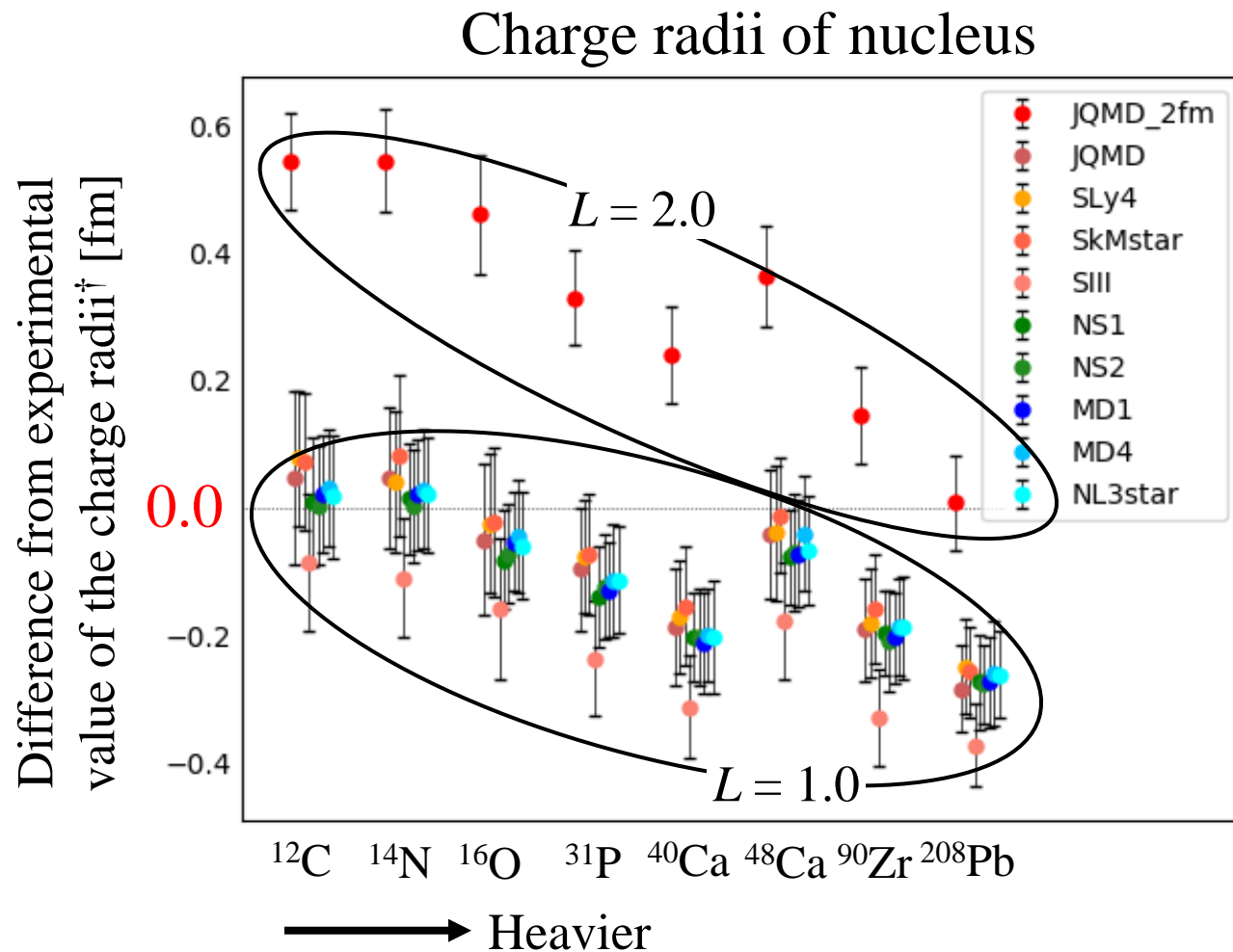
$$\varphi_i(\mathbf{r}) \equiv \frac{1}{(2\pi L)^{3/4}} \exp\left(-\frac{(\mathbf{r} - \mathbf{r}_i)^2}{4L} + \frac{i}{\hbar} \mathbf{r} \cdot \mathbf{p}_i\right)$$

$L = 2.0$ [fm] for QMD

$L = 1.26$ [fm] for LIQMD



For heavier nuclei, the difference from the experimental values is larger with small L . For this reason, at best, mass number $A \leq 64$ is recommended.



†: I. Angeli, K.P. Marinova, Atomic Data and Nuclear Data Tables, 99, 2013, 69-95.

Implementation

Implementation of LIQMD

Materials and Methods

qmd

Source code

light_ion_qmd

source/processes/hadronic/models/

G4QMDReaction

G4QMDMeanField

G4QMDCollision

G4QMDParameters

G4QMDNucleus

G4QMDGroundStateNucleus

G4QMDSystem

G4QMDParticipant

G4LightIonQMDReaction

G4LightIonQMDMeanField

G4LightIonQMDCollision

G4LightIonQMDParameters

G4LightIonQMDNucleus

G4LightIonQMDGroundStateNucleus

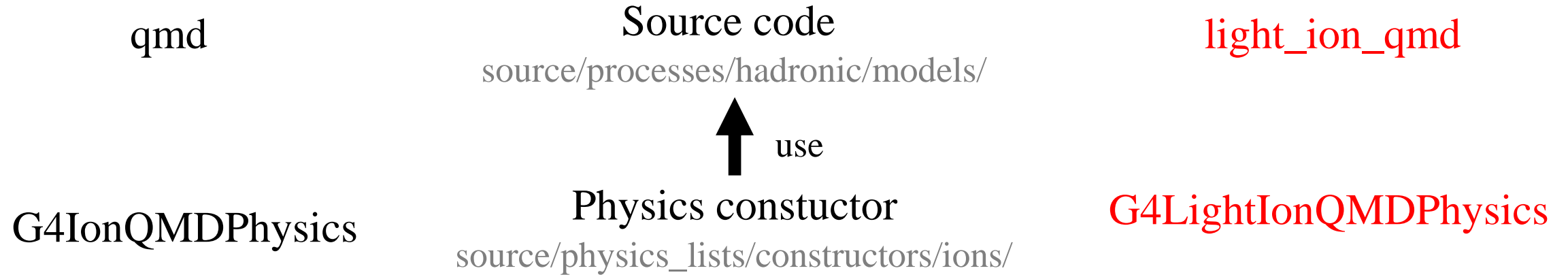
G4LightIonQMDSystem

G4LightIonQMDParticipant

I might put these files in the qmd folder.

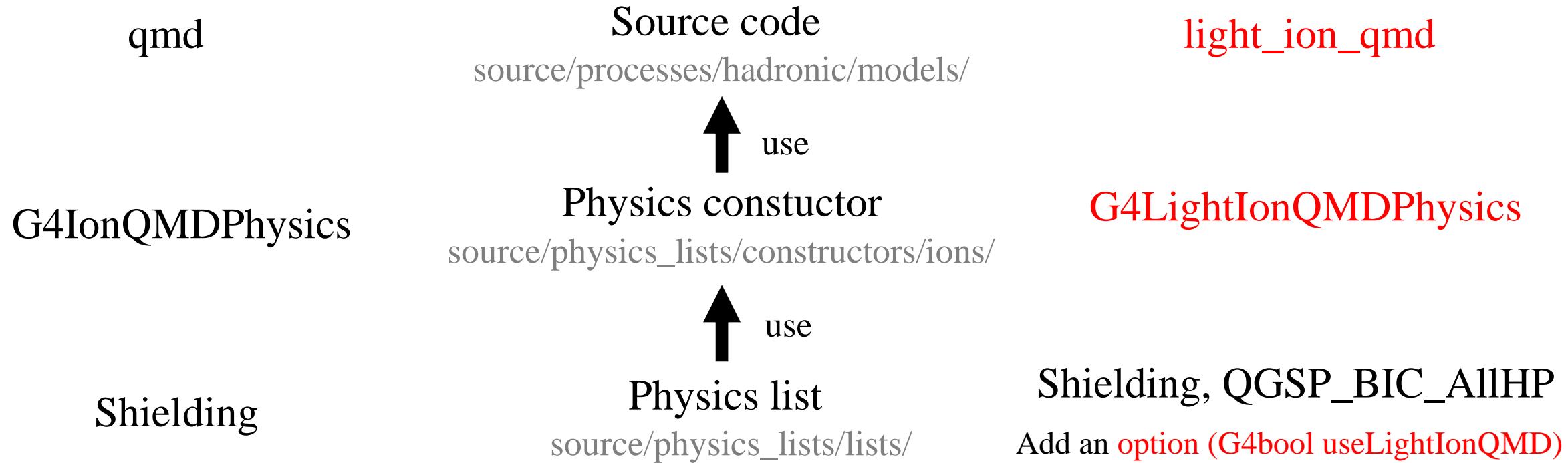
Implementation of LIQMD

Materials and Methods



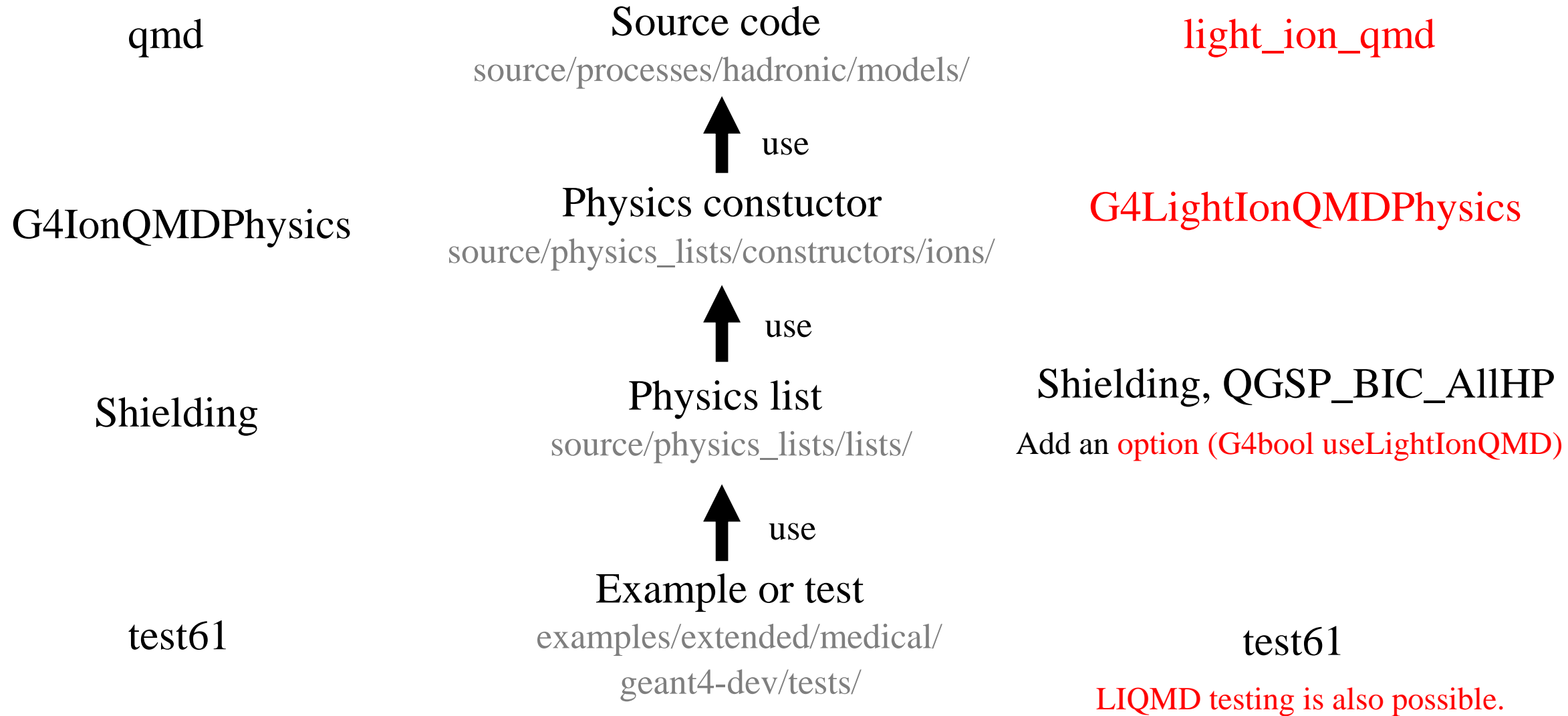
Implementation of LIQMD

Materials and Methods



Implementation of LIQMD

Materials and Methods



Summary and Future plan

- ❑ We have improved the QMD especially for hadron therapy using light ions and are considering to implement it as “Light Ion QMD” in Geant4.
- ❑ We have verified the performance of Light Ion QMD with various experimental data and confirmed that it is as good as or better than the conventional QMD model, at least in light ion beams and targets.
- ❑ Currently, the merge request is underway.

Thank you for listening