



# Summary of Hadronic Parallel Session 3A

Alberto Ribon  
CERN EP-SFT

# Parallel Session 3A : Hadronic Physics 1

- Yoshihide Sato “*Development and implementation of a new Geant4 QMD model and its validation*”
- Lorenzo Arsini “*Graph Neural Networks for fast emulation of nuclear interaction models*”
- Julia Yarba “*Update on tuning of hadronic model parameters*”

Yoshihide Sato

*“Development and implementation of a new Geant4  
QMD model and its validation”*

# Development and implementation of new Geant4 QMD model and its validation

**Yoshihide Sato,**  
Graduate School of Health Science,  
Tokushima University

## 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

**IPEM**  
Institute of Physics and  
Engineering in Medicine

**PAPER**

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

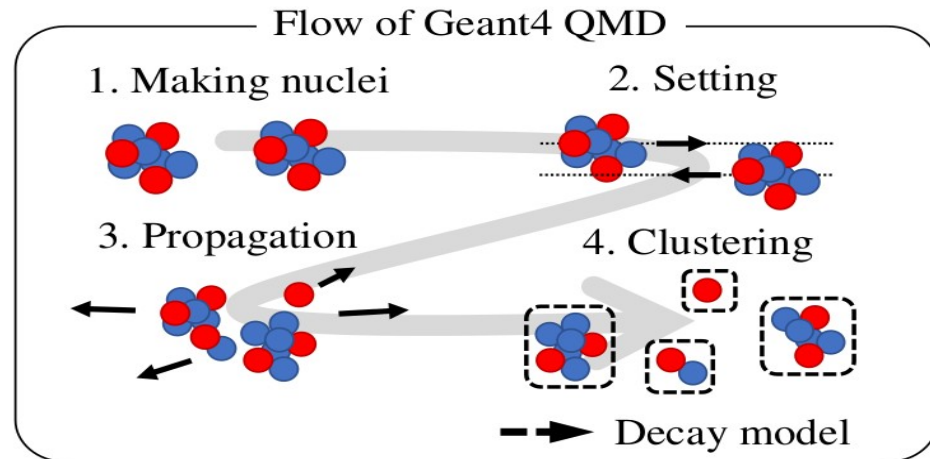
Yoshihide Sato<sup>1</sup>, Dousatsu Sakata<sup>2,3,\*</sup>, David Bolst<sup>4</sup>, Edward C Simpson<sup>5</sup>, Susanna Guatelli<sup>6</sup> and Akihiro Haga<sup>1,6</sup>

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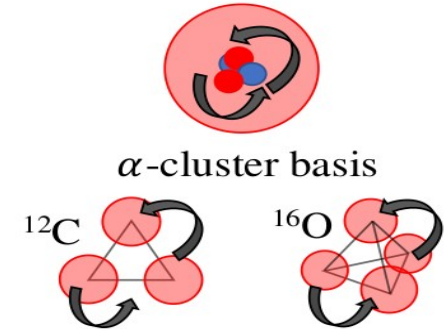


### Three improvements

#### 1. Modern interaction

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

#### 2. α-cluster structure



#### 3. QMD model Parameter optimization

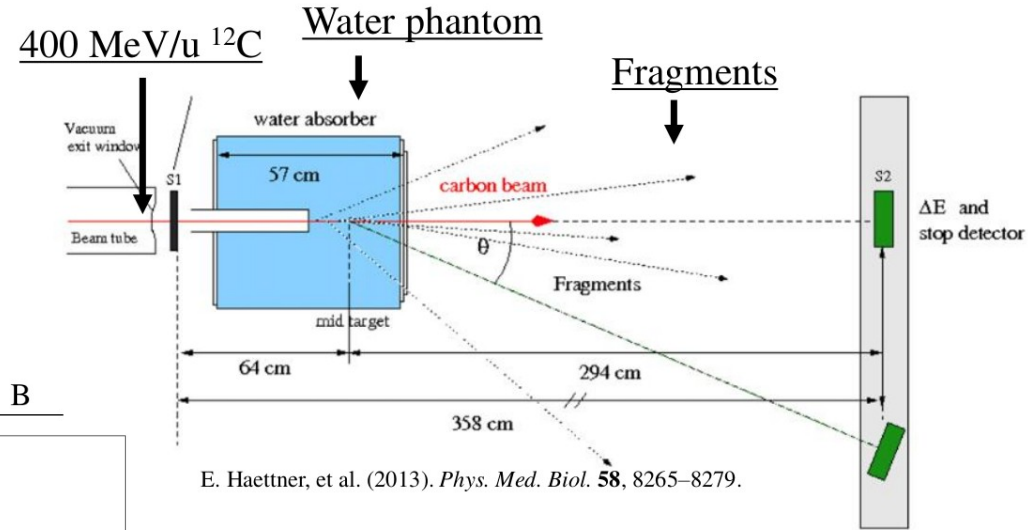
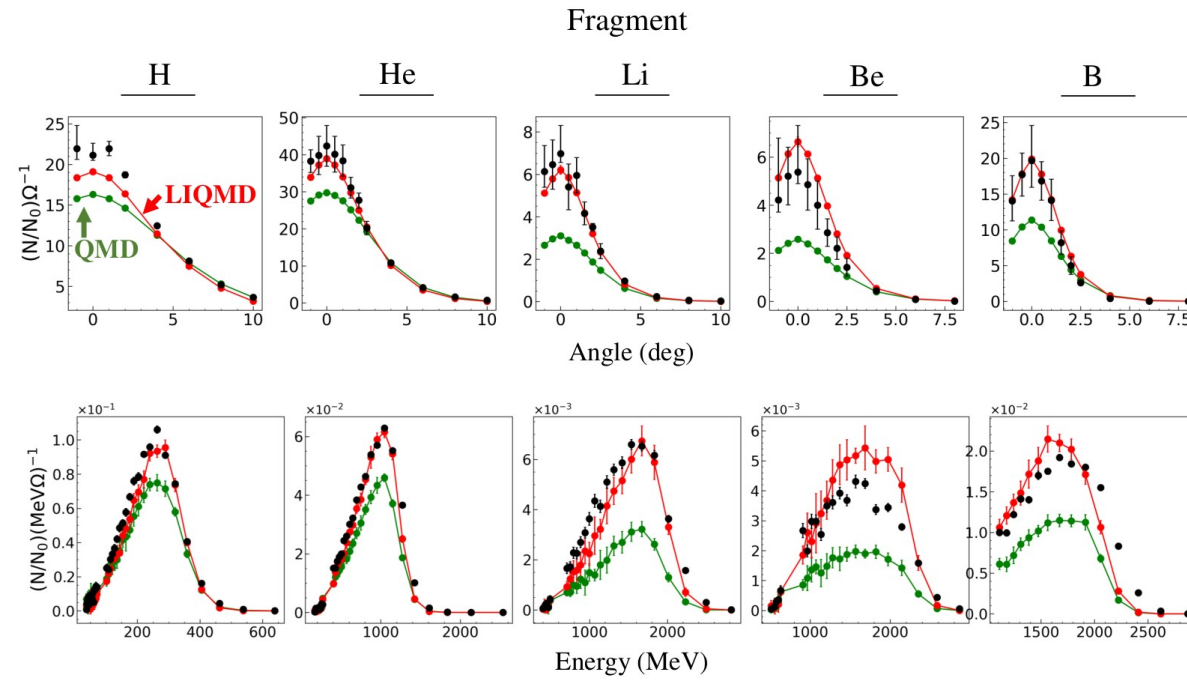
$$\varphi_i(\mathbf{r}) \equiv \frac{1}{(2\pi L)^{3/4}} \exp \left( -\frac{(\mathbf{r} - \mathbf{r}_i)^2}{4L} + \frac{i}{\hbar} \mathbf{r} \cdot \mathbf{p}_i \right)$$

$r_{ij}^2 \leq R^2$

# Development and implementation of new Geant4 QMD model and its validation

**Yoshihide Sato,**  
Graduate School of Health Science,  
Tokushima University

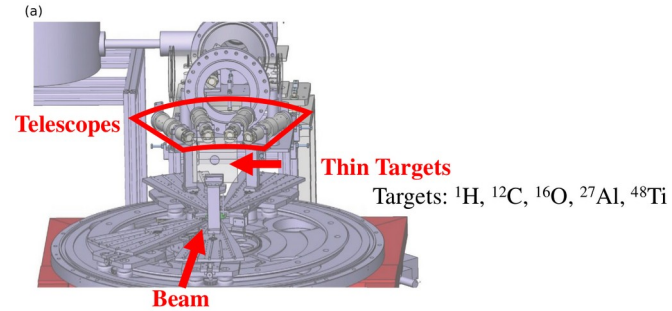
Recommended energy range: 30 – 500 MeV/u



# Development and implementation of new Geant4 QMD model and its validation

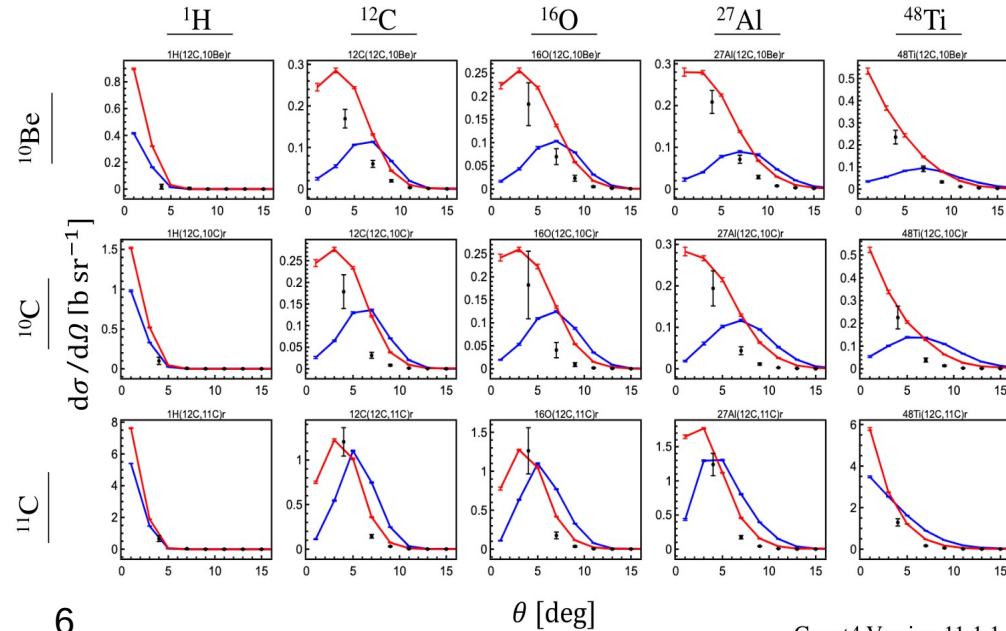
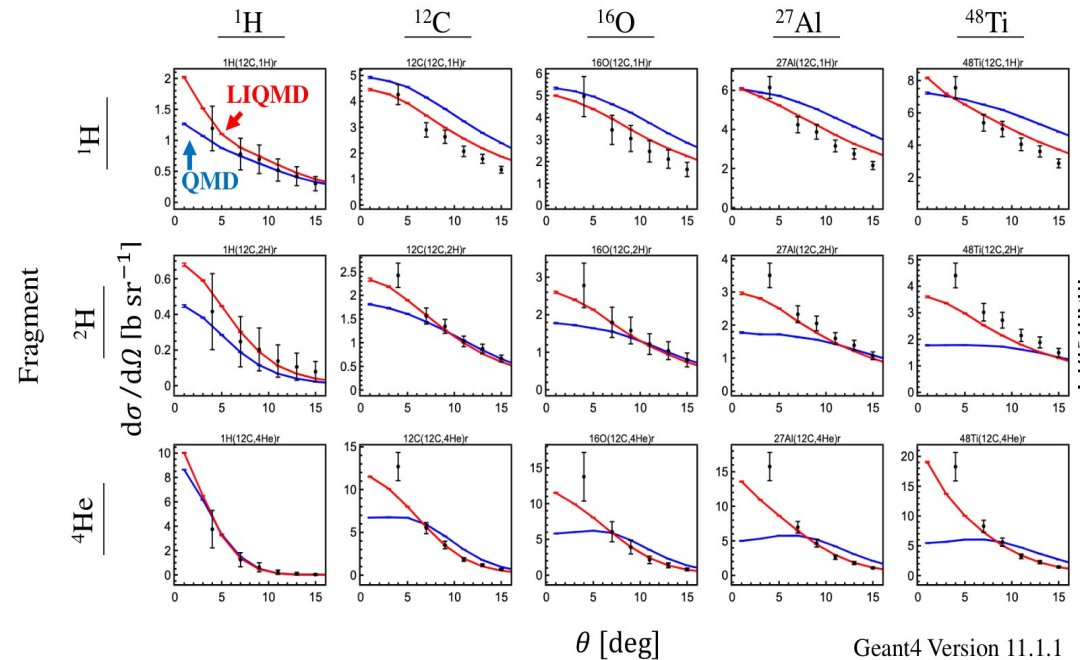
**Yoshihide Sato,**  
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Tokushima University

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

Recommended energy range: 30 – 500 MeV/u



Lorenzo Arsini

*“Graph Neural Networks for fast emulation of  
nuclear interaction models ”*

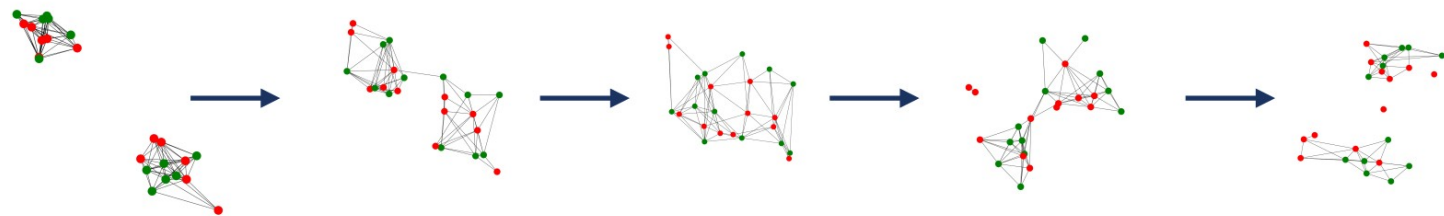


# Graph Neural Networks for fast emulation of nuclear interaction models

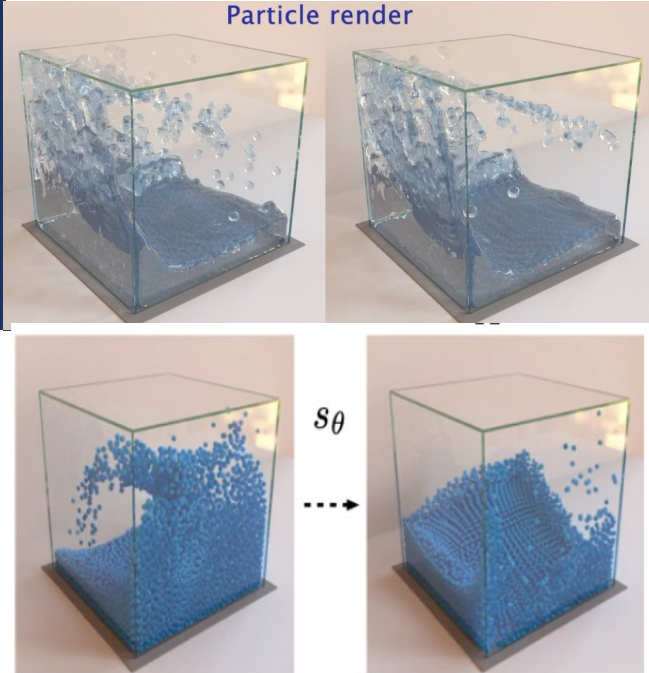
L. Arsini<sup>1,2</sup>, B. Caccia<sup>3</sup>, A. Giardiello<sup>1</sup>, M. Colonna<sup>4</sup>, S. Giagu<sup>1,2</sup>, C. Mancini Terracciano<sup>1,2</sup>

## GNN approach

Emulating the dynamics of **QMD** in  $^{12}\text{C}$  on  $^{12}\text{C}$  reaction at 12 MeV/u



Each nucleon is a node of the graph

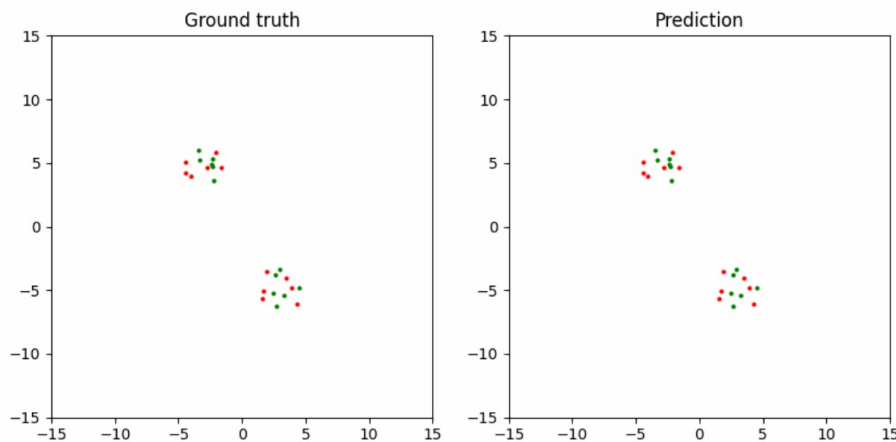




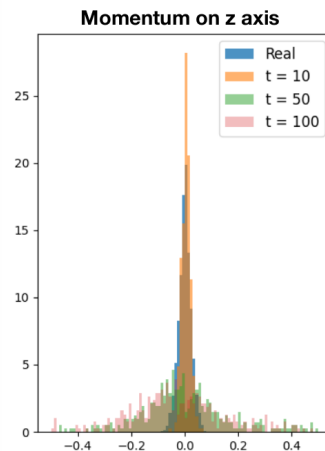
# Graph Neural Networks for fast emulation of nuclear interaction models

L. Arsini<sup>1,2</sup>, B. Caccia<sup>3</sup>, A. Ciardiello<sup>1</sup>, M. Colonna<sup>4</sup>, S. Giagu<sup>1,2</sup>, C. Mancini Terracciano<sup>1,2</sup>

Visually satisfying results...



... which are not Physical



In the **best case scenario**:

Quantities are conserved on average

Variance explodes increasing time steps

Cannot be used to infer physical quantities  
at the end of the reaction

Differences in the Physics

**QMD, BLOB**

Long range interactions  
Collisions

**Liquid simulations**

Short range interactions  
Gravity



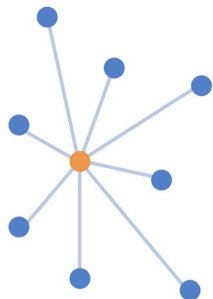
Building **fully connected** graphs

# Graph Neural Networks for fast emulation of nuclear interaction models

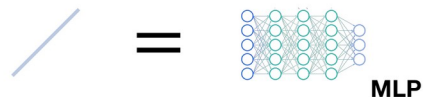
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## Learning the Potential: DL model

Particle-wise MLP for Potential Prediction



$$V_i = \sum V_{ij} = \sum f(q_i, q_j, p_i, p_j, c_i, c_j)$$



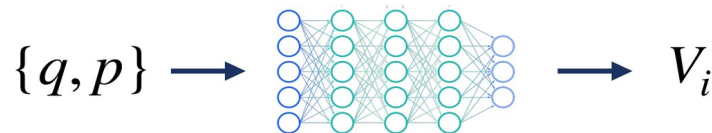
Embed particle exchange symmetry

## Why the Potential

Get more control on the Physics

No **blackbox** AI solution

DL computes a **known**,  
but complex function



Enforce physical **conservation laws** in the model

## Once you've learned the Potential

Mixing Deep Learning with  
standard methods

**Hybrid Models**

Get  $F_i$  **automatically  
differentiating**  $V_i$

**Integrate** the equation of  
motion with standard methods

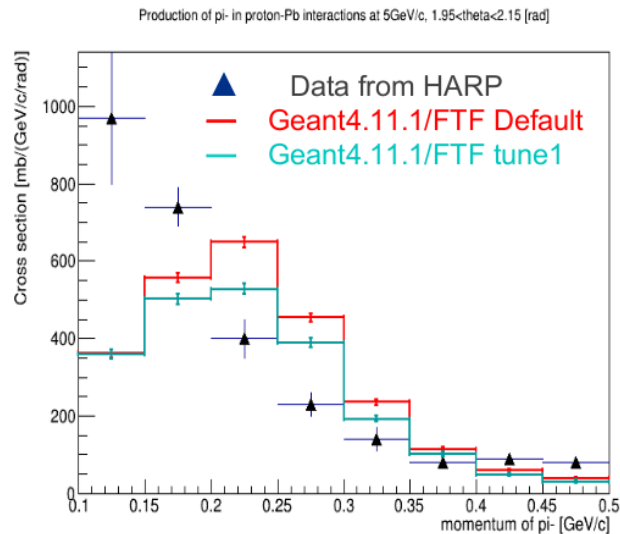
**Export** the DL model  
as a function

Integrate it into existing code,  
**replacing** potential computation  
GetPotential() method in QMD

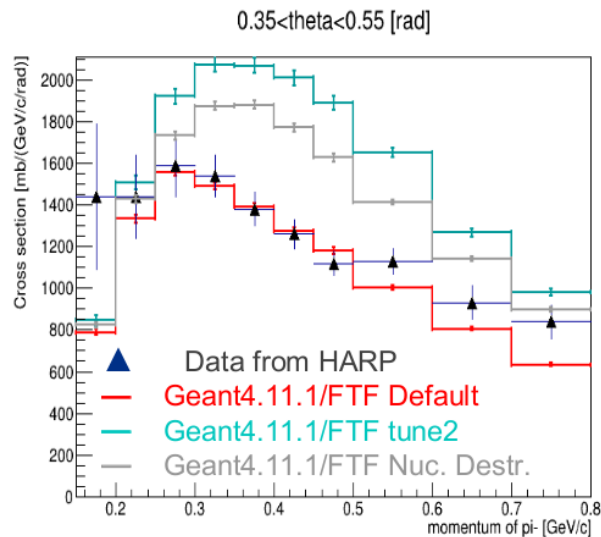
Julia Yarba

*“Update on tuning of hadronic model parameters”*

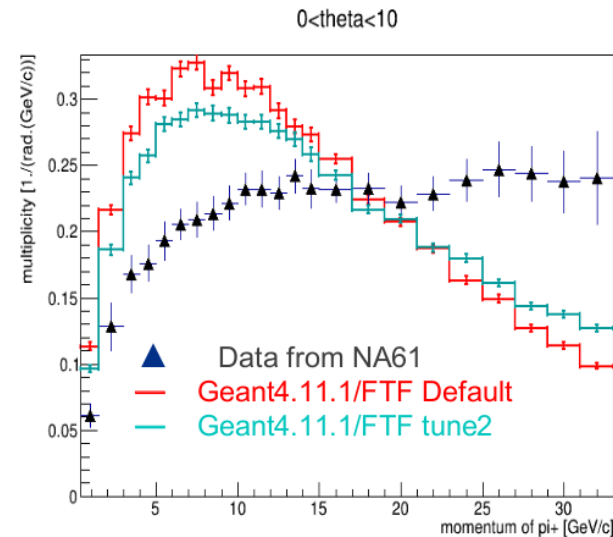
## Some examples of concerns with tune1 and tune2



5GeV/c proton on Pb  $\rightarrow \pi^-$   
For  $\pi^-$  production in the backward hemisphere, can we get the shape of the simulated spectrum right ?



5GeV/c  $\pi^+$  on Pb  $\rightarrow \pi^-$   
Increase in  $\pi^-$  production in the forward hemisphere is an artefact of using best fit parameters for the nuclear destruction (grey curve).  
Can we compensate ?



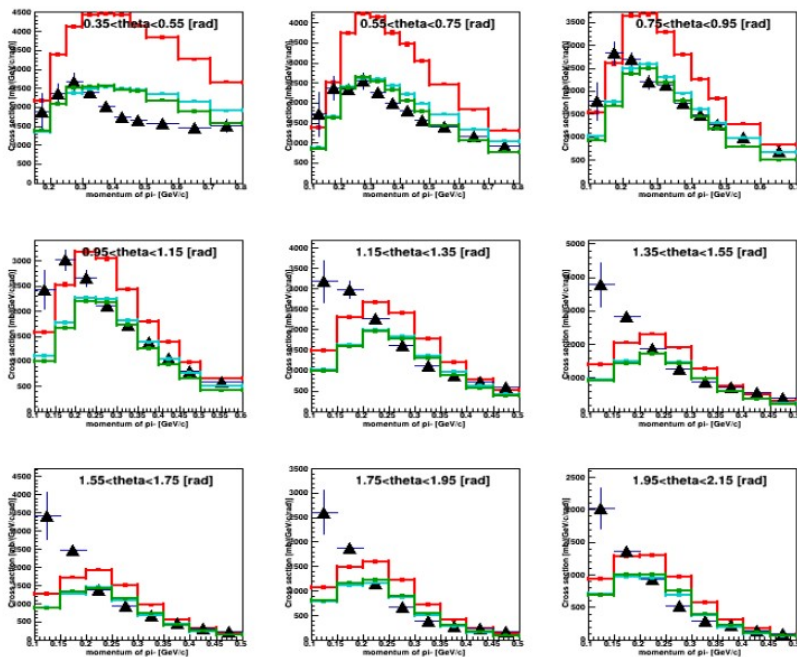
60GeV/c  $\pi^+$  on C  $\rightarrow \pi^+$   
Can we get the shape of the simulated spectrum right ?

# Update on Tuning Geant4 FTF Parameters Julia Yarba, Fermilab

Trying to change the thresholds in the **string mass sampling** to compensate for some discrepancies in the backward hemisphere, for the proton projectile tuning (tune1) → does not bring major benefits !

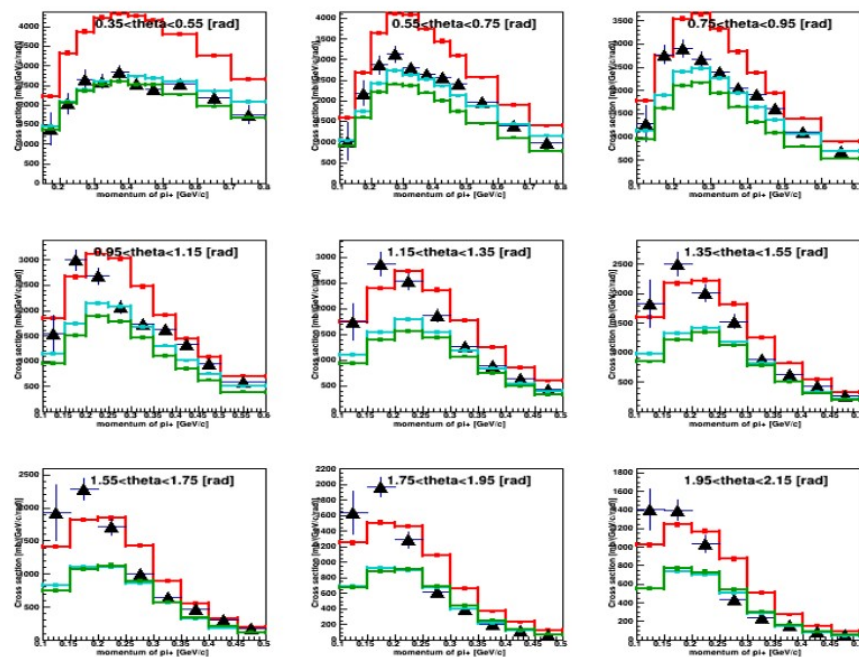
## 12 GeV/c proton on Pb (LA production of pions)

G4/FTF: 12.0GeV proton on Pb → piminus + X; data by HARP



G4/FTF Default  $\chi^2/NDF = 67.84$   
G4/FTF tune1  $\chi^2/NDF = 10.48$   
G4/FTF BestFit  $\chi^2/NDF = 11.0$

G4/FTF: 12.0GeV proton on Pb → piplus + X; data by HARP



G4/FTF Default  $\chi^2/NDF = 22.59$   
G4/FTF tune1  $\chi^2/NDF = 12.37$   
G4/FTF BestFit  $\chi^2/NDF = 7.71$

## Summary

- In general, using tuned (best fit) FTF parameters improves agreement between MC and thin target data, for both baryon (proton) or pion projectiles
- For the baryon projectile it includes FTF parameters involved in modeling nuclear target destruction and those that define contribution from quark exchange
  - Fitting/changing thresholds of the string mass sampling in the string formation does not bring sizeable benefits (although these parameters can produce non-negligible impact if varied individually)
  - Discrepancies between FTF results and thin target data for pions produced in the backward hemisphere by proton projectile on heavy nuclei are due to the underlying algorithms rather than currently available configurable parameters
- For the pion projectile, using FTF best fit parameters related to nuclear target destruction and quark exchange generally improves agreement between FTF results and thin target data but changing parameters of nuclear destruction may induce undesired increase in pion production on heavy targets
  - Perhaps the effect needs to be revisited/explored some more
  - The effect might be somewhat compensated by adjusting thresholds in string mass sampling
  - Some discrepancies between FTF results and experimental data, e.g. on pion production in the very forward direction by higher momentum beam (e.g. 60 GeV/c) are likely to be due to underlying mechanisms and not only the combination of FTF parameters defining contribution(s) from quark exchange (vs non-diffractive interactions)