

## Activities at INFN-LNS in Catania, Italy

GAP Cirrone, F Farrokhi, S Fattori, L Pandola, G Petringa, A Sciuto

# Group presentation



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INFN-LNS group is active since 2002

**G.A.Pablo Cirrone:** responsible for "hadrontherapy" and "radiobiology" examples pablo.cirrone@lns.infn.it

**Serena Fattori:** contact person for UHDR simulations in Geant4-DNA; Geant4-DNA tutorials (chem6) serena.fattori@lns.infn.it

**Alberto Sciuto**: machine learning development in Geant4 / Geant4-DNA alberto.sciuto@lns.infn.it

**Giada Petringa**: LET/RBE simulations vs. microdosimetric data giada.petringa@lns.infn.it

Luciano Pandola: Coordination of the EM Working Group; Maintenance and bug fix of Penelope; STT shifts; Evolution Task Force luciano.pandola@lns.infn.it

A new post-doc will join the group on November 2023

100 % for Geant4 work

100 % for Geant4 work

Geant4 activities

## Geant4 activities

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Hadrontherapy progress, maintenance and G4Med connections

LET/RBE simulations vs microdosimetry data

Contribution into the EM Working group

Developing and maintenance of the Hadrontherapy example



Hadrontherapy was published in 2002 and it has been evolved during the years;

**Details** in the dedicated talk on the "Examples" Session;

Part of the example output inserted in the **G4-MED** framework.

GAP Cirrone, PhD - pablo.cirrone@Ins.infn.it

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# Coordination / service tasks within LNS

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Deputy coordination of the EM Working Group (Low Energy)

# Maintenance and bug fix of the Penelope (and other EMLowEnergy) models

Bug #2465 on fluorescence after Livermore photoelectric still open

Could not reproduce with TestEm5 (a lot of support by Zhuxin!)

Contribution to the **STT shifts** (3 weeks/year)

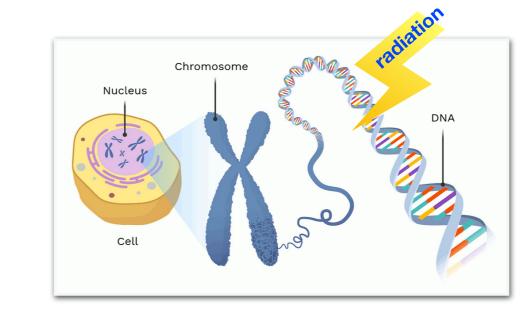
Contribution to the Collaboration Evolution Task Force

Technical support to the **Ferrara group** for the implementation of the Baier-Katkov model --> See talk by A. Sytov

G4DNA activities (from macro to micro and nano)

## G4DNA activities

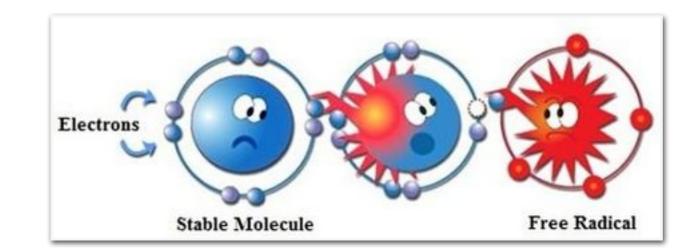


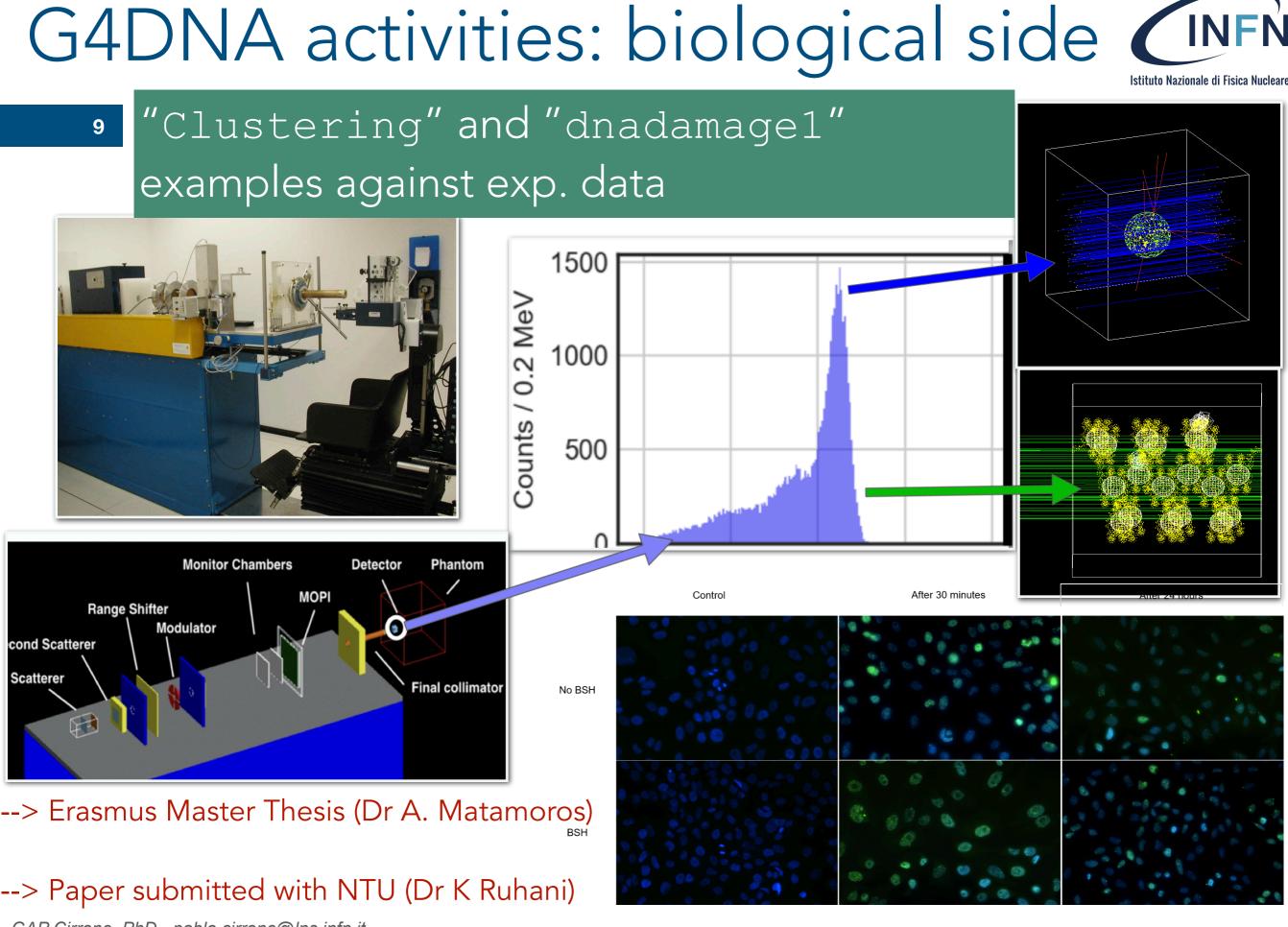


## Biological side

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## Chemical side





# G4DNA activities: chemical side

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Oxygen depletion studies with Geant4-DNA: Geant4-DNA (scavenger) compared to TRAX-CHEM

\$759	
E.C.M.	
ELSEVIER	

Radiation Physics and Chemistry

journal homepage: www.elsevier.com/locate/radphyschem

Contents lists available at ScienceDirec

Effects of the Oxygen depletion in FLASH irradiation investigated through Geant4-DNA toolkit

Fateme Farokhi <sup>a,b</sup>, Babak Shirani <sup>a,\*</sup>, Serena Fattori <sup>b,\*\*</sup>, Mohammad Ali Asgarian <sup>a</sup>, Giacomo Cuttone <sup>b</sup>, Sayyed Bijan Jia <sup>c</sup>, Giada Petringa <sup>b</sup>, Alberto Sciuto <sup>b</sup>, G.A. Pablo Cirrone <sup>b,d,e</sup> <sup>a</sup> Faculty of Physics, University of Isfahan, Isfahan, Iran <sup>b</sup> Istituto Naxionale di Fisica Nucleare (INFN), Laboratori Nazionali del Sud, Catania, Italy

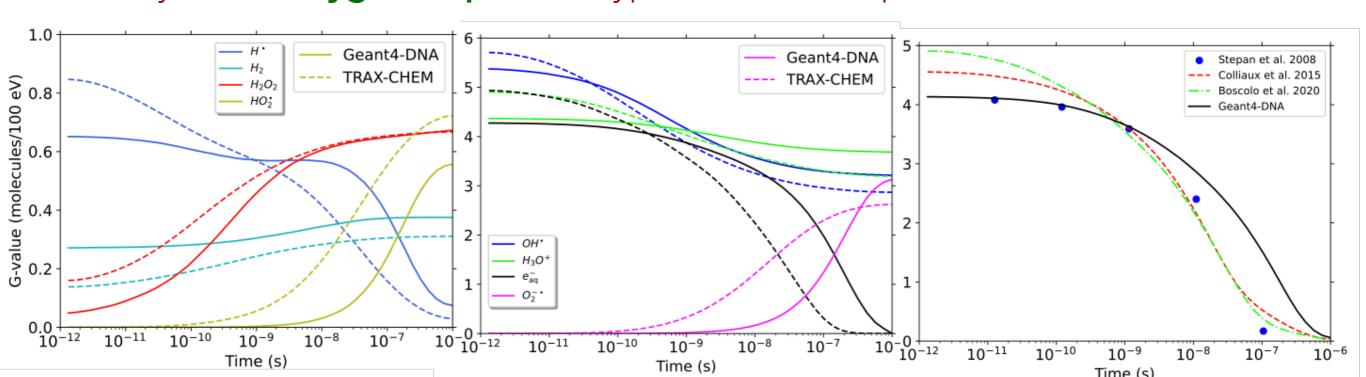
<sup>c</sup> Department of Physics, University of Bojnord, Bojnord, Iran <sup>d</sup> Centro Scillano di Fisica Nucleare e Struttura della Materia, Catania, Italy

<sup>d</sup> Centro Siciliano di Fisica Nucleare e Struttura della Materia, Catania, Italy <sup>e</sup> Dipartimento di FISICA ED ASTRONOMIA "Ettore Majorana" - Università degli Studi di Catania, Catania, Italy

#### (2023) doi.org/10.1016/j.radphyschem.2023.111184



#### **G-values** of different radiolitic species after **proton and electron irradiation**. --> Study of the **oxygen depletion** hypothesis in comparison with TRAX-CHEM



# Side activities: the BIORAD III project

## The BIORAD-ESA activity

# Istituto Nazionale di Fisica Nucleare





Experimental biological damage quantification and simulation

Institutes

INFN (Italy), IBFM (Italy), IRSN (France), CENBG (France), VINCA (Serbia), Univ. Sevilla (Spain)

Experimental radiobiological data database for comparison against Monte Carlo simulation (i.e. the output from the "Hadrontherapy" example (RBE module from Survival Fraction)











Christelle ADAM-GUILLERMIN, Pierre BEAUDIER, Marco CALVARUSO, Giovanni CANTONE, Konstantinos CHATZIPAPAS, Davide CHIAPPARA, Giuseppe Antonio Pablo CIRRONE\*, Miguel CORTES-GIRALDO, Milos DORDEVIC, Serena FATTORI, Orsola GIAMPICCOLO, Franck GOBET, Sébastien INCERTI, Giada PETRINGA, Ivan PETROVIC, Aleksandra RISTIC-FIRA, Alberto SCIUTO, Hervé SEZNEC, Ngoc Hoang TRAN

## The BIORAD-ESA activity





### Two kinds of biological endpoints:

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5 1, 0.756, 0.126 6 2, 0.516, 0.066 7 3, 0.409, 0.069

8 4, 0.257, 0.050

10 21, 0.056, 0.015

10, 0.109, 0.022

FOCI via the H2AX protein Phosphorylation and

Survival Fraction (or Clonogenic)

Clone	ogenicSurvival_Data_example $ angle$ No Selection	
1		on; SOBP; vertical; 24.2; 60.3; 1.0; 1.0; 02; 0.036; 0.010; 0.003; 28.6; -1;
2		
3	DOSE, SF, SEM_SF	
4	0, 1.000, 0.185	

Example of **Clonogenic** data file

B < > FOCL\_Data\_example\_normal
FOCL\_Data\_example\_normal > No Selection
1 92-1; 5., 55., 25., 15.; 60Co-gamma; SOBP; vertical; 24.0; 1.0; 1.0; immuno-yH2AX; 10.3109/09553002.2010.481322
2
3 TIME, FOCI, SEM\_F, DSB, SEM\_D, LET, ENERGY, DURATION
4 -0, 5.82, 0.87, 0.97, 0.14, 36.0 20. 5.
5 0.5, 22.13, 0.62, 3.69, 0.10, 36.0 20. 5.
6 24.2, 11.63, 1.07, 1.94, 0.18, 36.0 20. 5.

Example of **FOCI-H2AX** data file

## The BIORAD-ESA activity



**D**eesa



#### Readme for the different part

Structure of the Clonogenic information

Structure of the FOCI information

Structure of the Cell-lines information

CellName; CellCycle; BeamParticle; BeamType; BeamOrientation; Depth; Dose; DoseRate; Assay; DOI

TIME, FOCI, SEM\_F, DSB, SEM\_D, LET, ENERGY, DURATION
Time\_1, Foci\_1, Err\_F\_1, DSB\_1, Err\_D\_1, Let1, Energy1, Duration1
Time\_2, Foci\_2, Err\_F\_2, DSB\_2, Err\_D\_2, Let2, Energy2, Duration2

========	
CellName: String of one word containing the "code-name" of the	he irradiated cell Database/
line, according to the cell line file;	IBFM/
CellCycle: Group of 4 float values, comma separated with each	h other, indicating FOCI 92-1 Proton CNR-IB
the percentage of cells in each cycle phase during the irrad	
the following order: sub-G1, G1, S, G2/M;	FOCT MCF10A Proton CNR-
BeamParticle: String of one word containing the type of beam	(use: Proton, 4He,
12C, 16O, 60Co-gamma, etc.);	ClonogenicSurvival_U87_
BeamType: String indicating the beam type with the acronym:	IRSN/
"FE": for Full Energy peak beam	FOCI_Data_Layout_example
"SOBP": for Spread Out Bragg Peak	FOCI_Data_Layout_example
"uBF": for micro Beam Focused (using particle by part	cicle in a specific FOCI Data Layout example
subcellular compartment)	FOCI Data Lavout example
"uBD": for micro Beam Defocused (using several partic	cles as a defocused US/
beam irradiating a large cell surface)	
BeamOrientation: String indicating the beam orientation with	
"horizontal" or "vertical";	U VINCA/
Depth: Float value indicating the depth in water in mm;	92.1 cell line gamma ir
Energy: Float value indicating the kinetic energy at the cell	l-layer point, in 92.1 cell line proton i
AMeV;	ARPE19 cell line gamma
Dose: Float value indicating the dose in Gy;	ARPE19 cell line proton
DoseRate: Float value indicating dose-rate in Gy/min;	UTR177 coll line gamma
Assay: String of one word indicating the method used to quant	
DOI: doi of publication related to the data contained in the	file. HTB177 cell line proton

### Format of the data file and data list

Database/				
IBFM/				
FOCI 92-1 Proton CNR-IBFM				
FOCI ARPE19 Proton CNR-IBFM				
- FOCI MCF10A Proton CNR-IBFM				
ClonogenicSurvival U87 Proton CNR-IBFM				
IRSN/				
FOCI_Data_Layout_example92.1_NEW-updated_MICRO_GononG1_Proton_gH2AX				
FOCI_Data_Layout_example92.1_NEW-updated_MICRO_GononG2_Proton_53BP1				
FOCI_Data_Layout_example92.1_NEW-updated_MICRO_GononG3_Alpha_gH2AX				
FOCI_Data_Layout_example92.1_NEW-updated_MICRO_GononG4_Alpha_53BP1				
US/				
L USE U2OS 2019				
VINCA/				
- 92.1 cell line gamma irradiation				
ARPE19 cell line gamma irradiation				
- ARPE19 cell line proton irradiation				
- HTB177 cell line gamma irradiation				
HTB177 cell line proton irradiation				

Machine Learning approaches

# Machine Learning and Geant4

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Current activities

We are currently developing a **post-processing tool** to bridge results across Geant4 and Geant4DNA in medical physics applications;

Tree based **regression models** are used to learn damage behaviours at the DNA scale in the form of **SSBs and DSBs** 

Trained models are used at the end of macroscopic simulations to give **an estimate of DNA damage** 

## Machine Learning



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### Future activities

Improve computational efficiency in Geant4 based simulations for medical physics trough the **integration of a ML super-resolution models** in voxelized detectors

Post-processing tool based on a CNN (Convolutional Neural network) model trained with high density voxelized detector data.

Application of the model to **retrieve spatial distribution of secondary particles** on low resolution voxelized detectors.

Roberto Catalano, Davide Rassarello, Pablo Cirrone, Emilio Zappalà, Nino Amato, Luciano Pandola, Giuliana Milluzzo, Michele Costa, Mariacristina Guarrera, Serena Fattori, Antonio Russo, Beatrice Cagni, Alma Kurmanova, Carmen Altana, Andrea Matamoros, Giuliana Navarra, Salvo Tudisco, Giacomo Cuttone, Giada Retringa, Gustavo Messina INFN-LNS Medical Physics Group - Catania, April 30, 2021

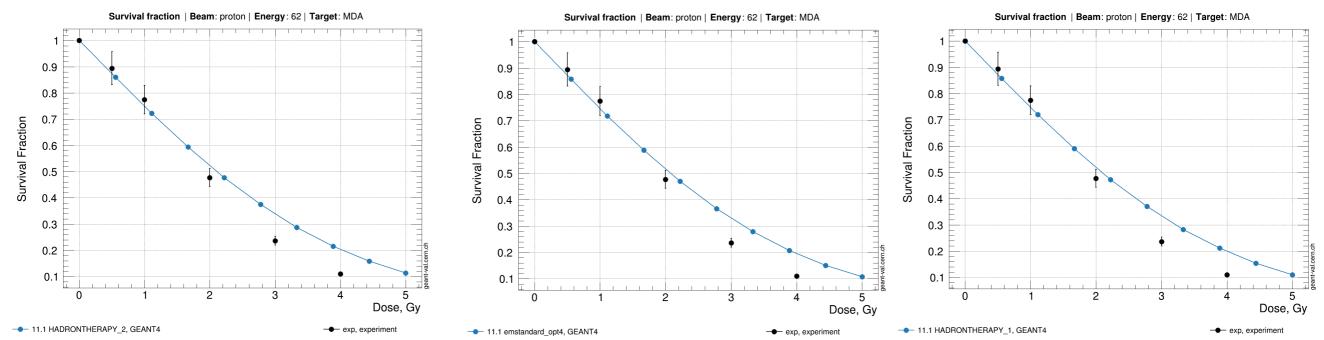
Thanks or listening

IN N LABORATORI NAZIONALI DEL SUD

# Hadrontherapy testing in G4-MED

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# Hadrontherapy is one of the examples present in the **Geant-Val platform**



Current version monitories the cell **Survival Fraction and RBE calculation** for a 62 MeV Proton Beam with the three em options of the example: emstandard\_opt4, HADRONTHERAPY\_1 and HADRONTHERAPY\_2;

LET calculation and comparison with data for proton, Carbon, Helium and Oxygen beams of clinical interest, are being inserted GAP Cirrone, PhD - pablo.cirrone@Ins.infn.it

### LET simulations vs microdosimetry data: protons



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$$\bar{L}_d = \frac{\sum_{i=1}^N L_i \varepsilon_i}{\sum_{i=1}^N \varepsilon_i}$$

$$\bar{L}_d^{Total} = \frac{\sum_{j=1}^n \left[\sum_{i=1}^N L_i \varepsilon_i\right]_j}{\sum_{j=1}^n \left[\sum_{i=1}^N \varepsilon_i\right]_j}$$

 $\bar{L}_T = \frac{\sum_{i=1}^N L_i l_i}{\sum_{i=1}^N l_i}$ 

 $\bar{L}_{T}^{Total} = \frac{\sum_{j=1}^{n} [\sum_{i=1}^{N} L_{i} l_{i}]_{j}}{\sum_{j=1}^{n} [\sum_{i=1}^{N} l_{i}]_{j}}.$ 

L: total electronic stopping power  $\varepsilon$ : energy loss

t: track length

GAP Cirrone, PhD - pablo.cirrone@Ins.infn.it

Physics in Medicine & Biology

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

PAPER

Monte Carlo implementation of new algorithms for the evaluation of averaged-dose and -track linear energy transfers in 62 MeV clinical proton beams

G Petringa<sup>1</sup>, L Pandola<sup>1</sup>, S Agosteo<sup>2,3</sup>, R Catalano<sup>1</sup>, P Colautti<sup>4</sup>, V Conte<sup>4</sup>, G Cuttone<sup>1</sup>, K Fan<sup>5</sup>, Z Mei<sup>5</sup>, A Rosenfeld<sup>6</sup>, A Selva<sup>4</sup> and GAP Cirrone<sup>1,\*</sup>

(2020) doi.org/10.1088/1361-6560/abaeb9

**New formulation** for LET-dose and LET-track

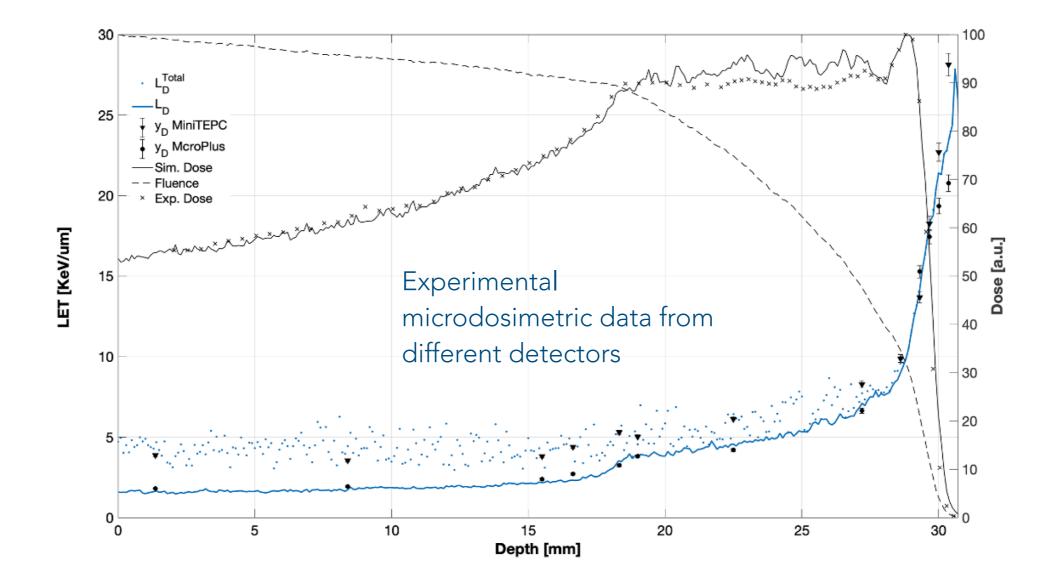
Independence from the production cut

Extensible to higer-Z ions

Taking into account the **primary** beam and the **secondaries produced** in hadronic interactions

## LET simulations vs microdosimetry data: protons

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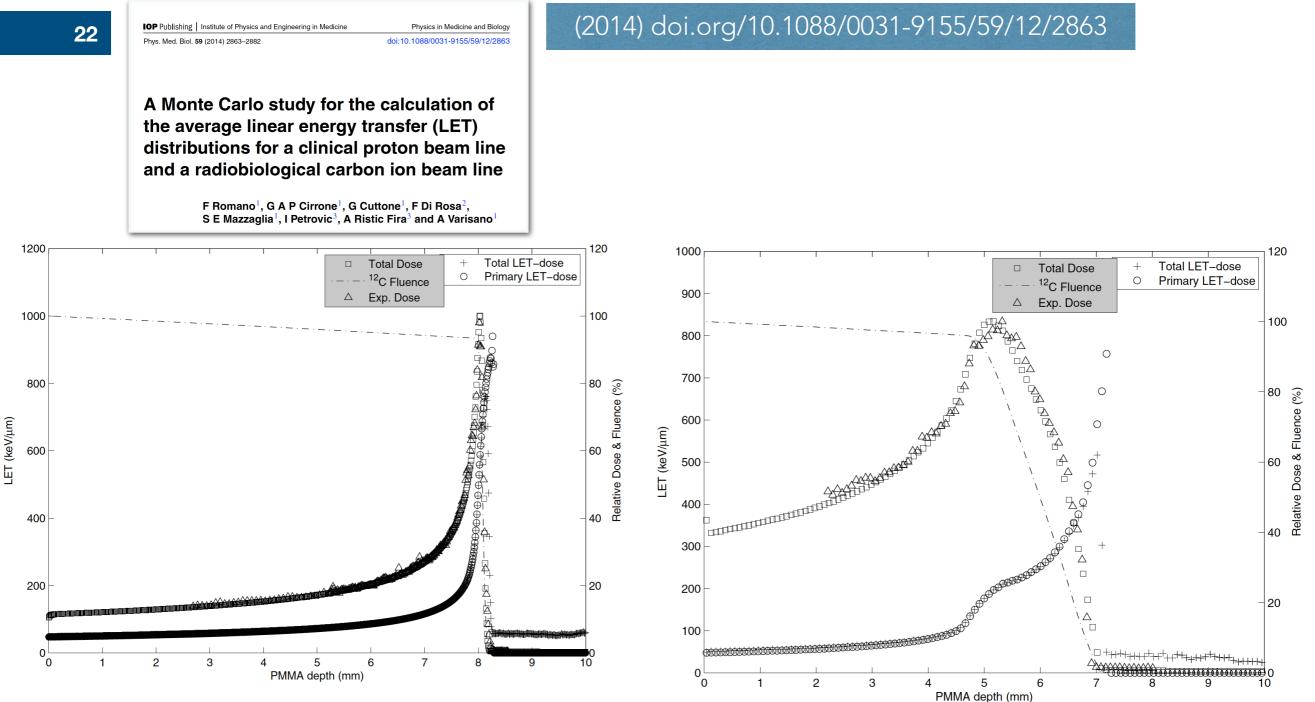
Istituto Nazionale di Fisica Nucleare

#### Contribution of **secondaries**;

Agreement with detectors able to also evaluate the **fragments**; **Projectile fragmentation** already predicted in 2014 by our group

### LET simulations vs microdosimetry data: Carbon

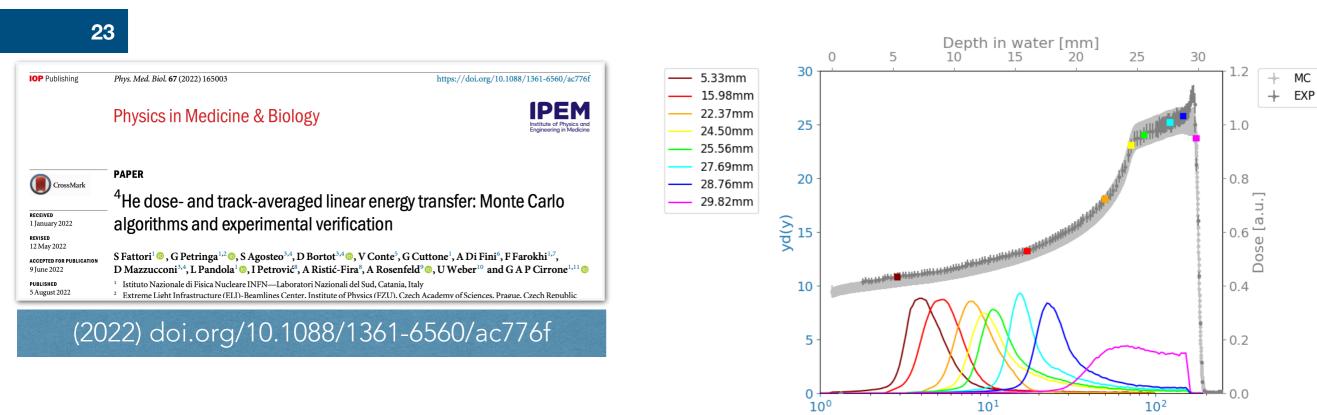


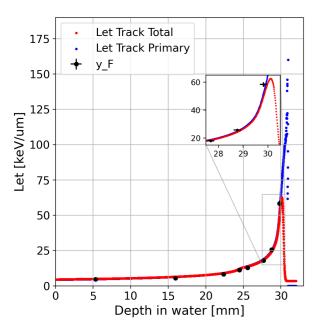


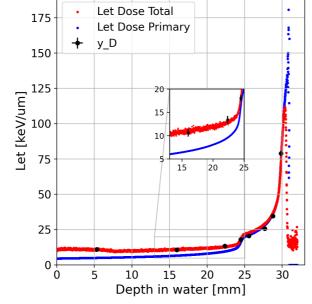
Mono- and poly-chromatic 62 AMeV Carbon beam Also validated against microdosimetric data

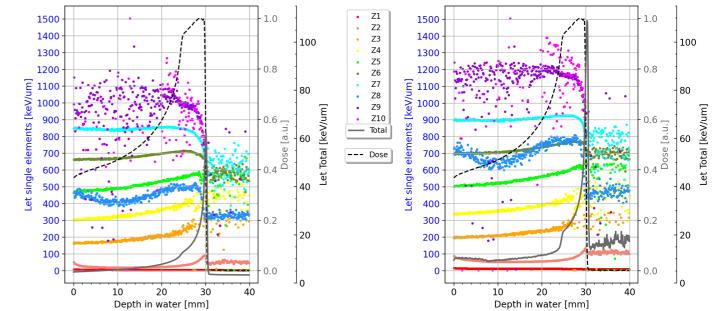
### LET simulations vs microdosimetry data: Helium, 62 AMeV











y [keV/um]

#### Oxygen study is ongoing