

# **Computational and experimental methods applied for treatment planning, quality assurance, and research at the clinically operating proton therapy facility**

**Antoni Rucinski, IGFAE Workshop, 10.05.2023**

[antoni.rucinski@ifj.edu.pl](mailto:antoni.rucinski@ifj.edu.pl)

# Witnessing PT development in Europe

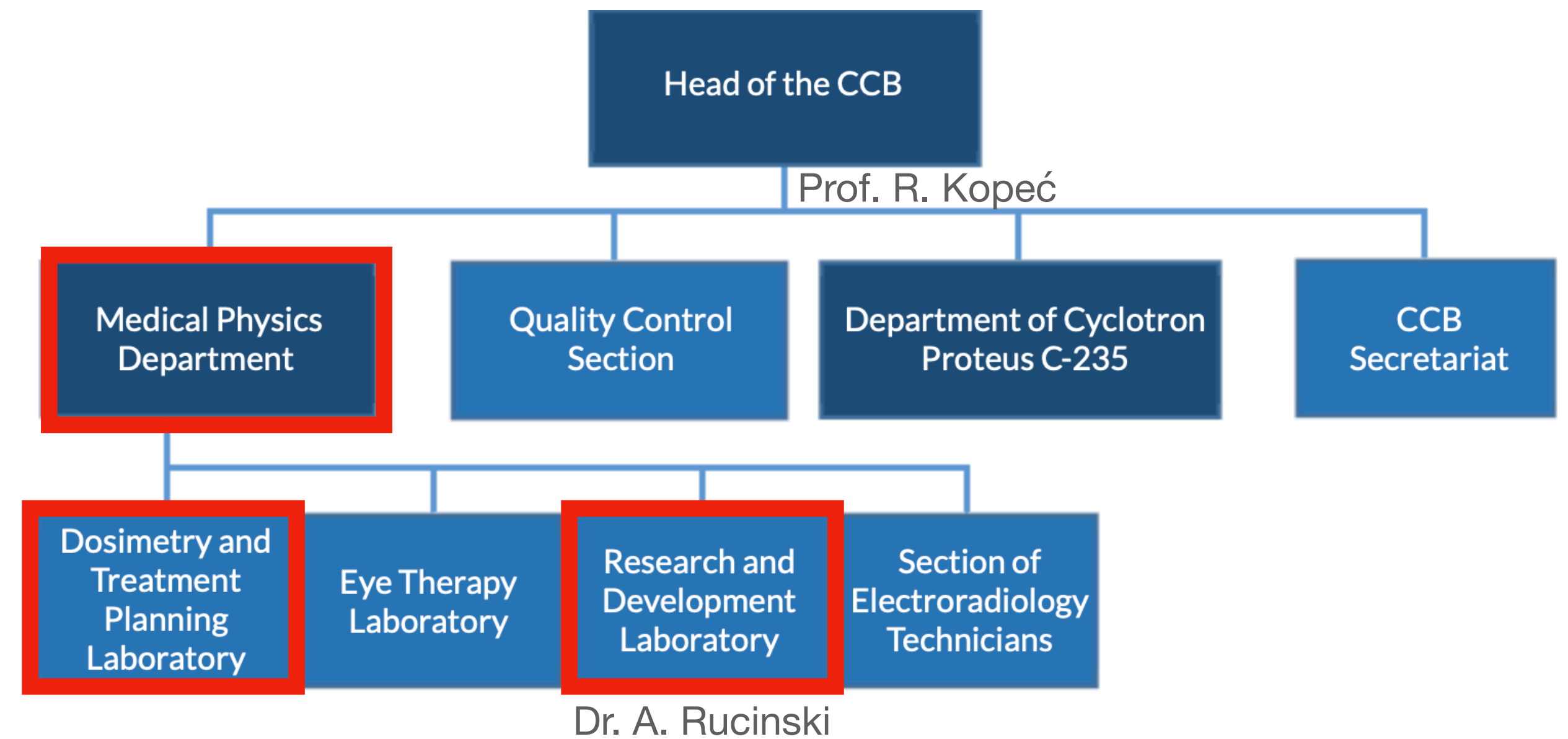
## My story



1	2004 -2009	Warsaw Technical University Universität Stuttgart (Erasmus Fellow)	MSc
2	2010 -2013	University of Heidelberg	PhD
3	2013 -2014	University Clinic Heidelberg SLK-Kliniken Heilbronn	MPE
4	2015 -2016	INFN & Sapienza University of Rome	Postdoc
5	2017 -now	Institute of Nuclear Physics Polish Academy of Sciences CCB Kraków Proton Beam Therapy Centre	Assistant Professor

# CCB Kraków proton center

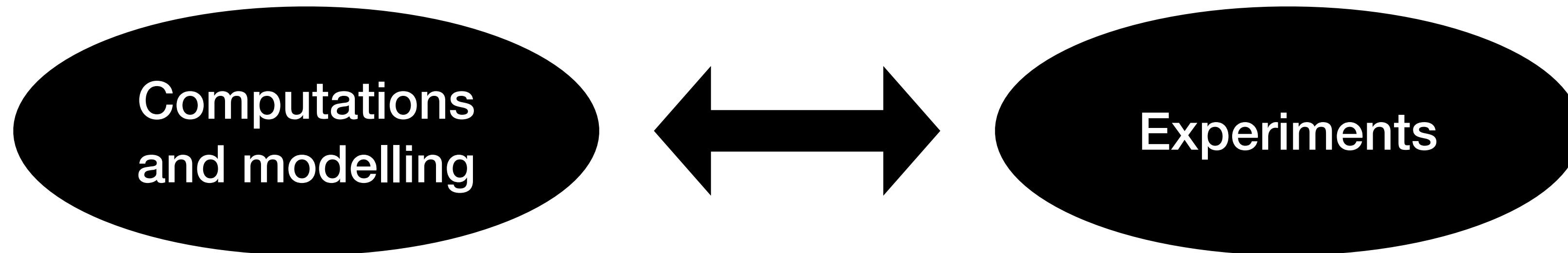
## Structure and relevant data/equipment



- Access to proton beams in Gantry rooms for experiments
- Anonymised patient data for TP studies and protocols development
- Radiobiology labs equipped to perform in vitro experiments (from 2023)

# Translational research at CCB proton center

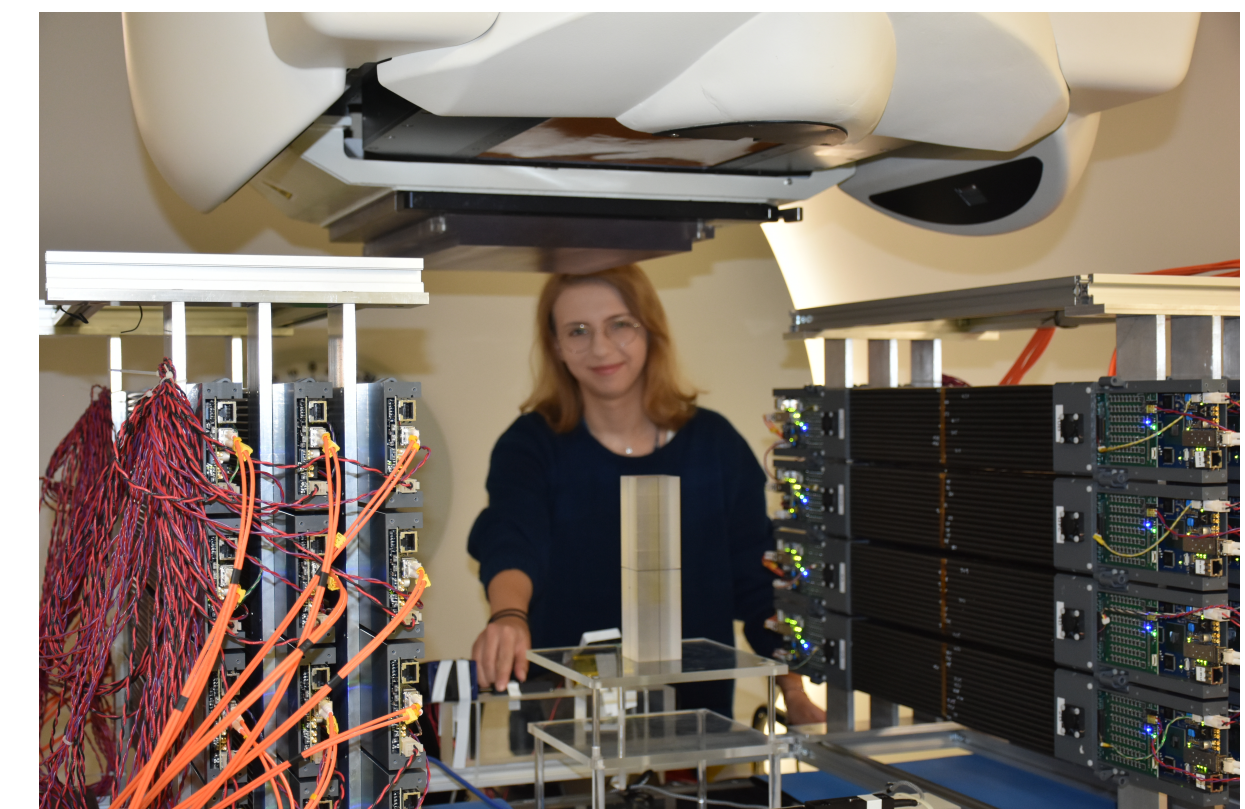
Physics, radiation biology, and oncology to improve clinical protocols



Biologically and LET weighted treatment planning



Quality assurance

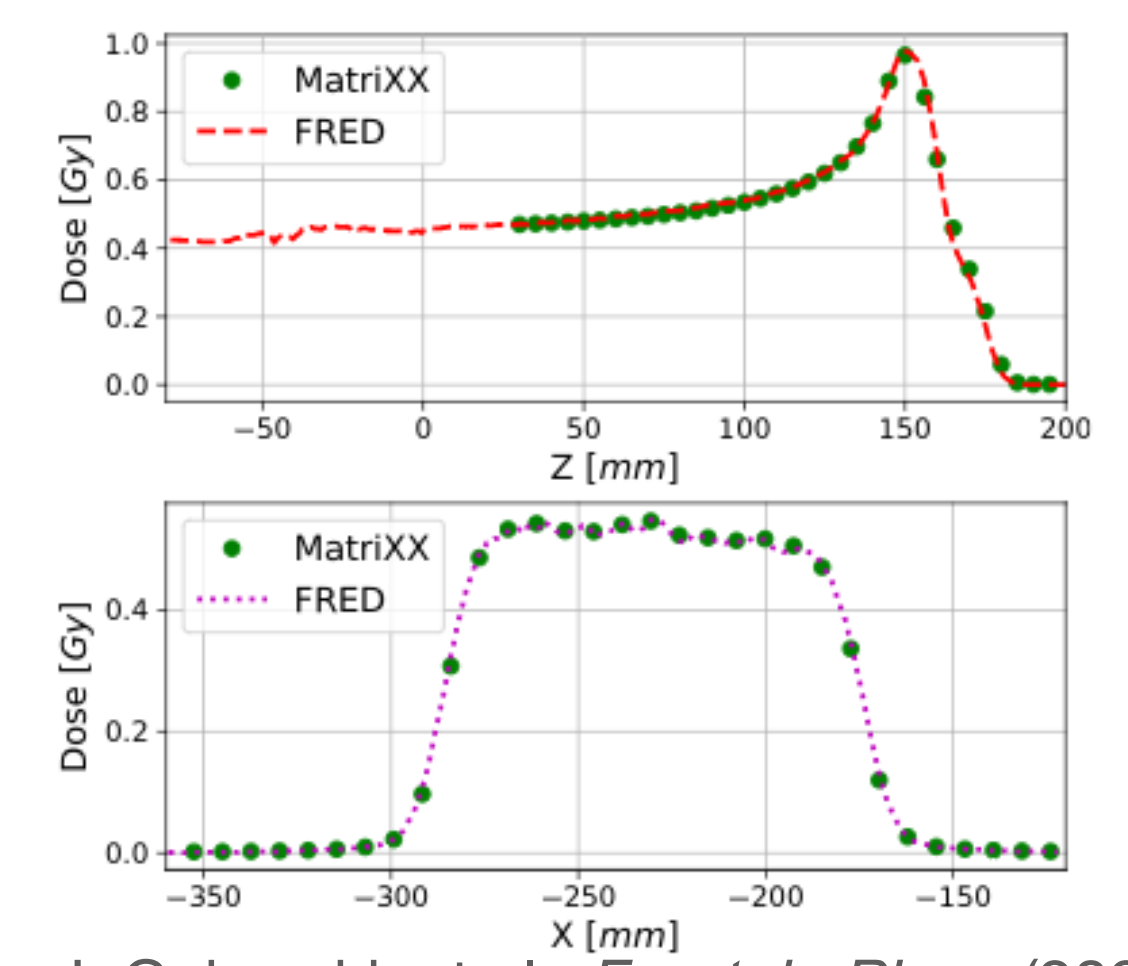
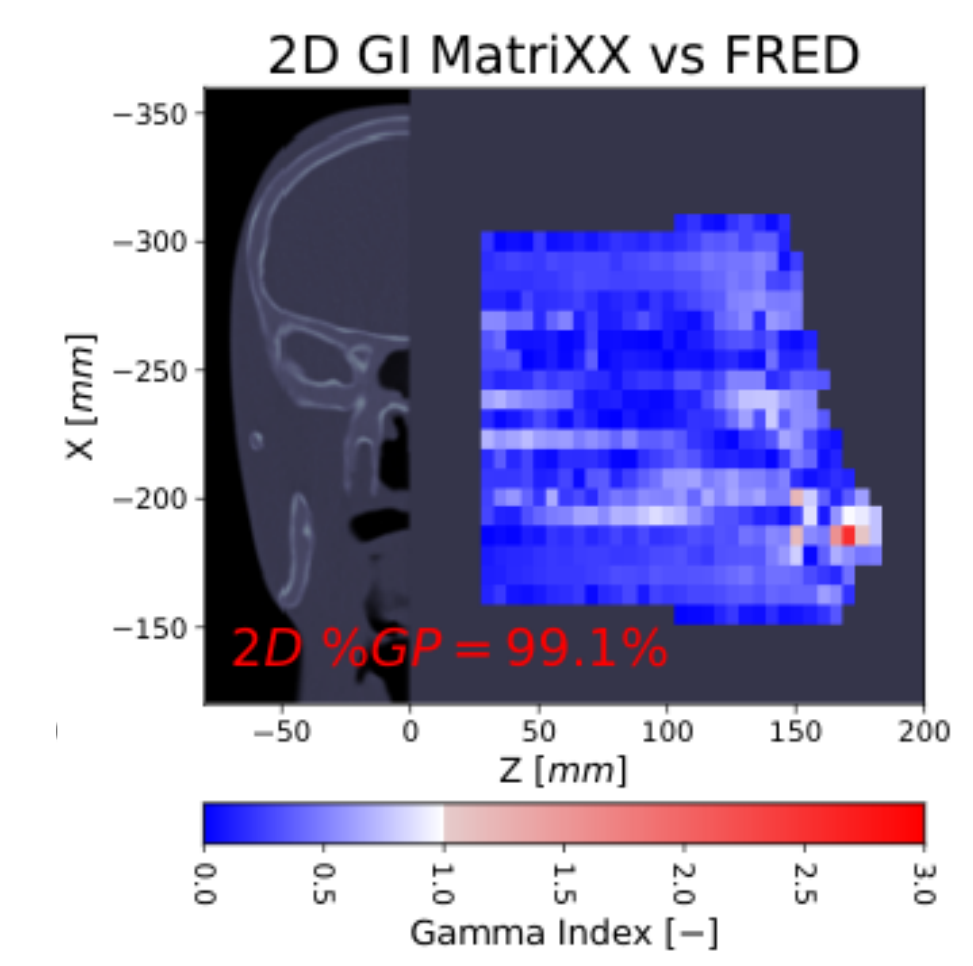
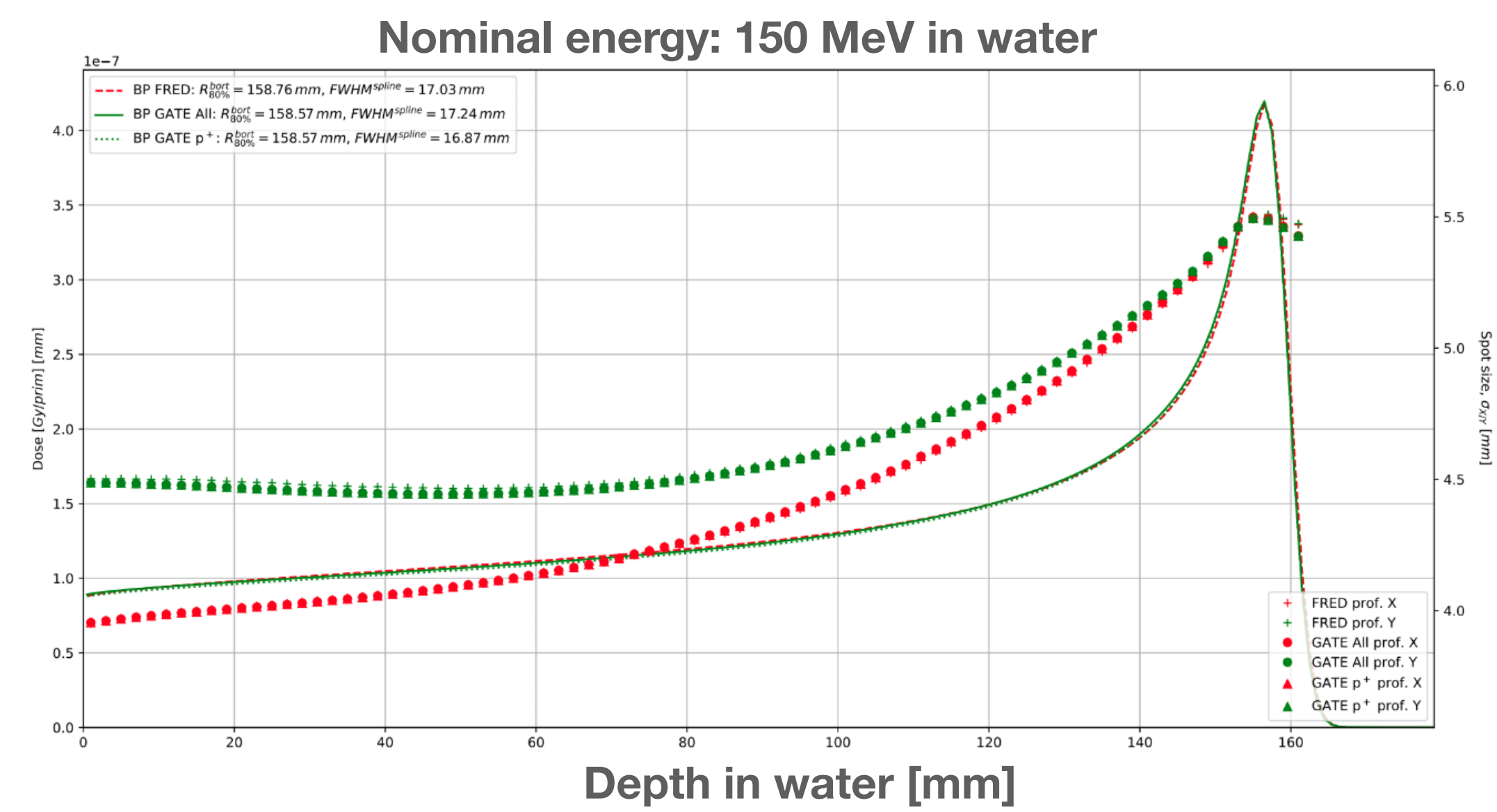
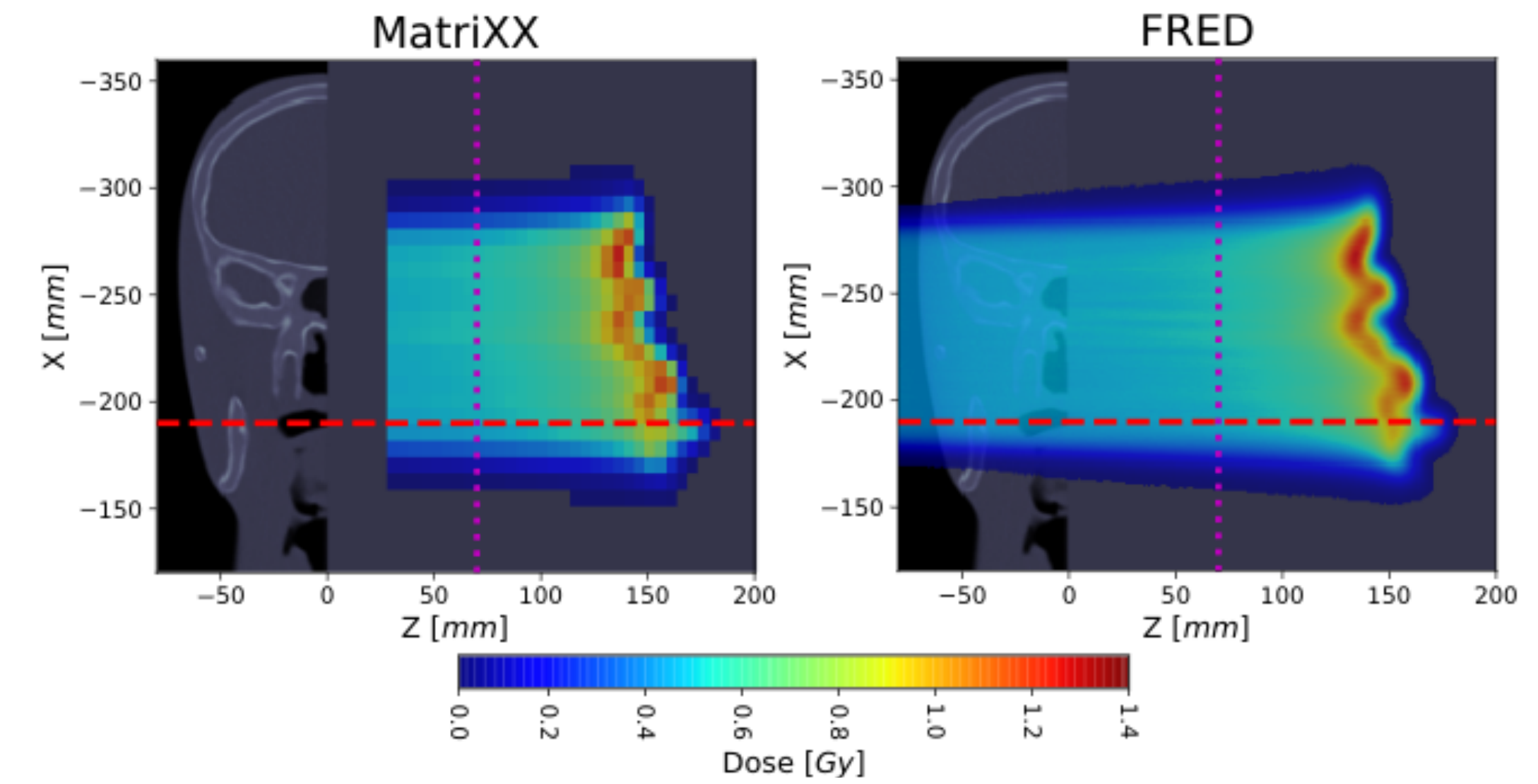
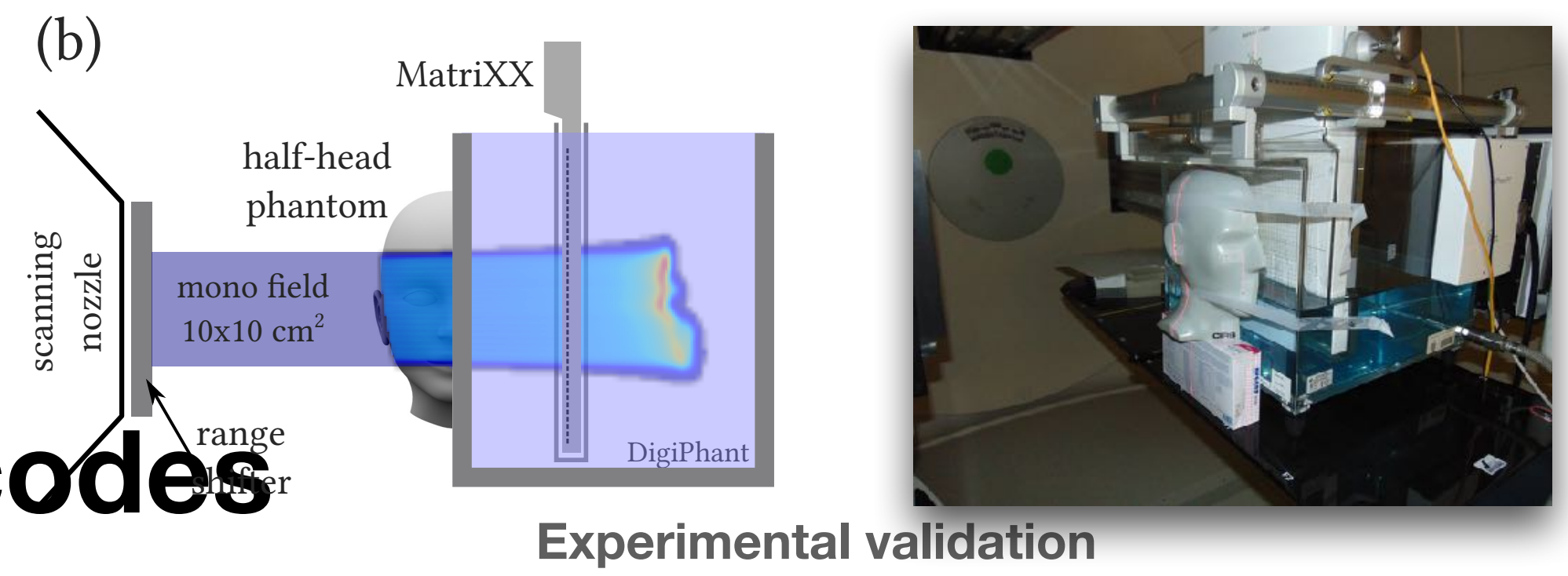


Detectors for range monitoring for therapy adaptation

# Beam modeling

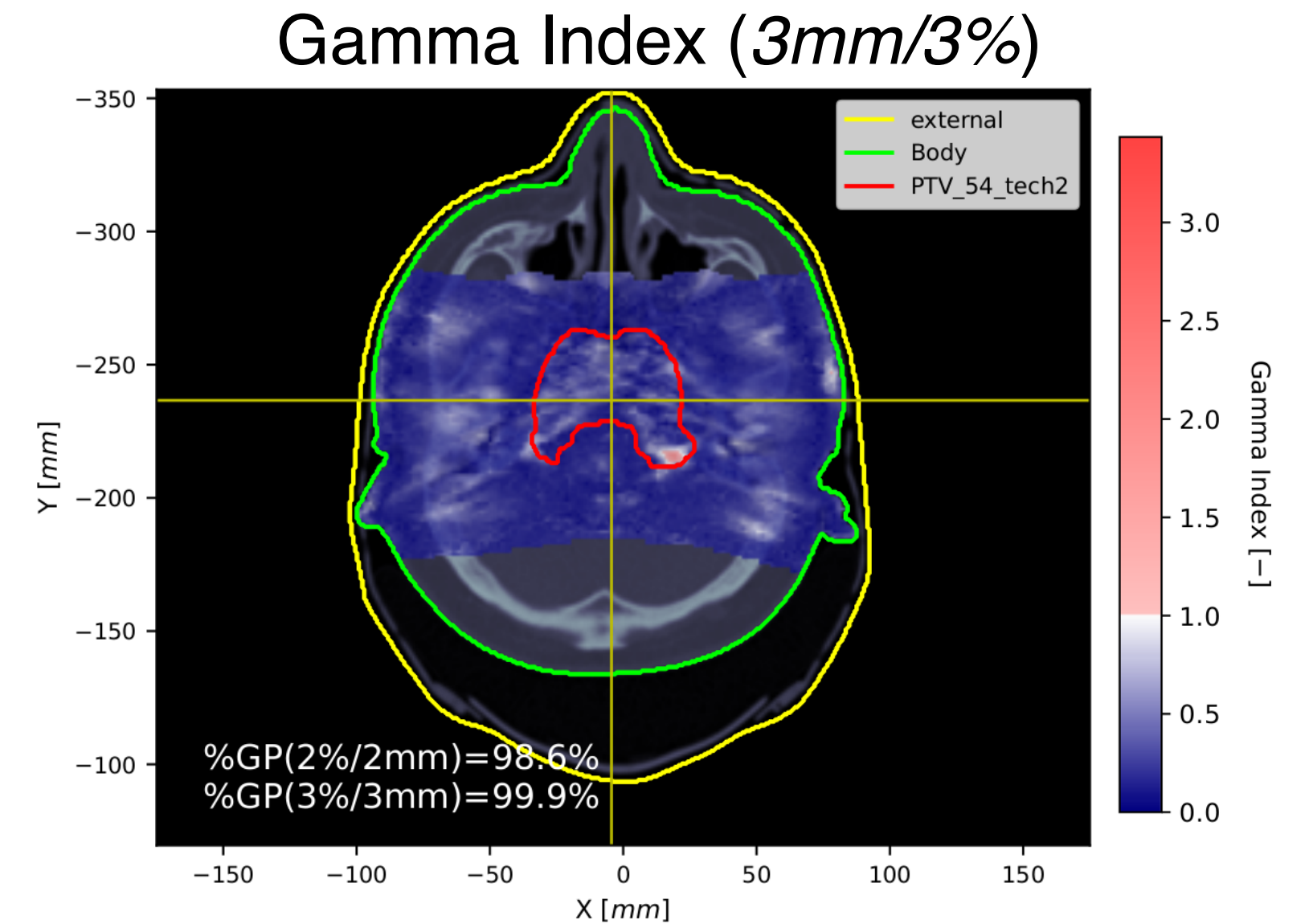
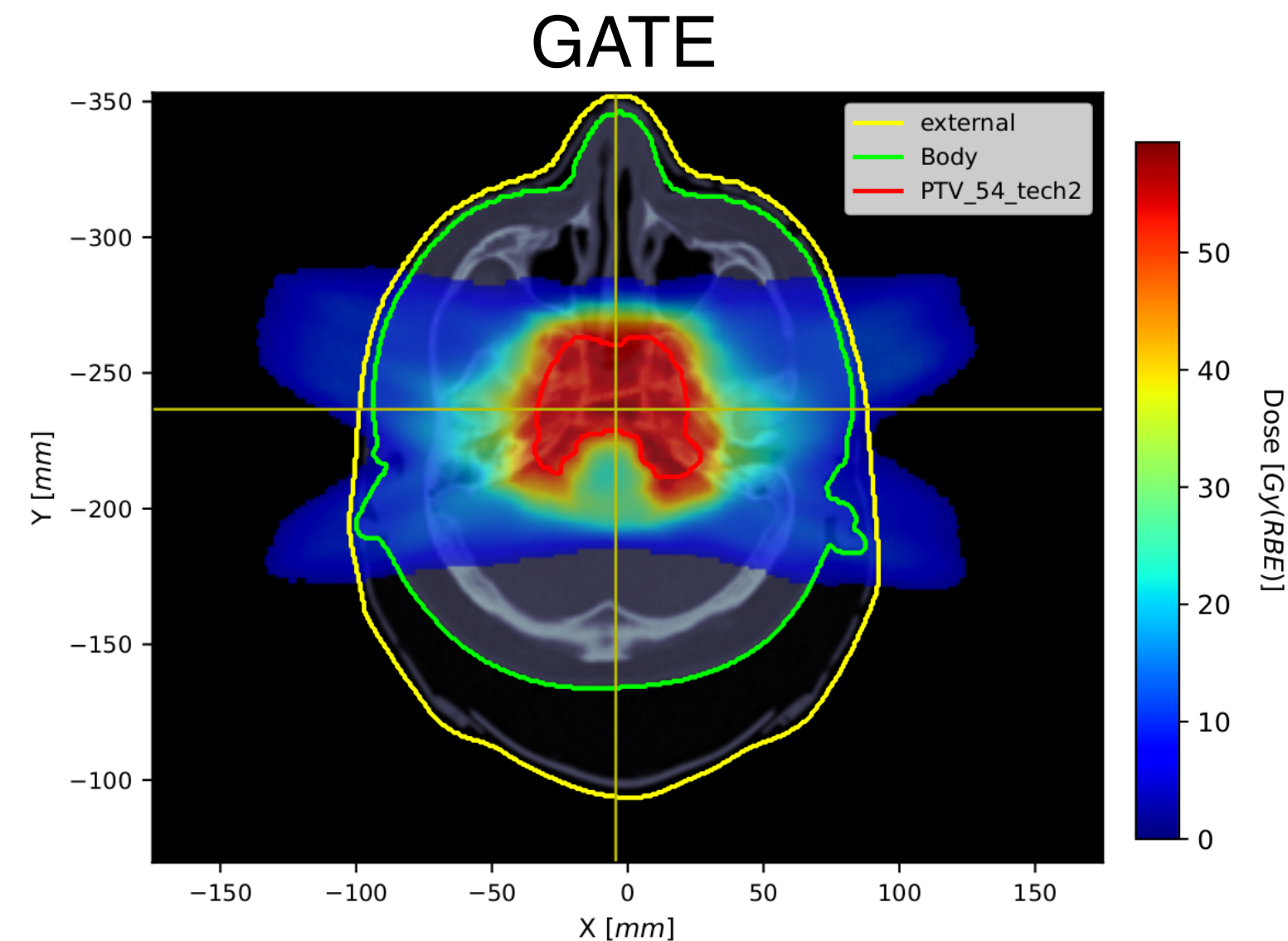
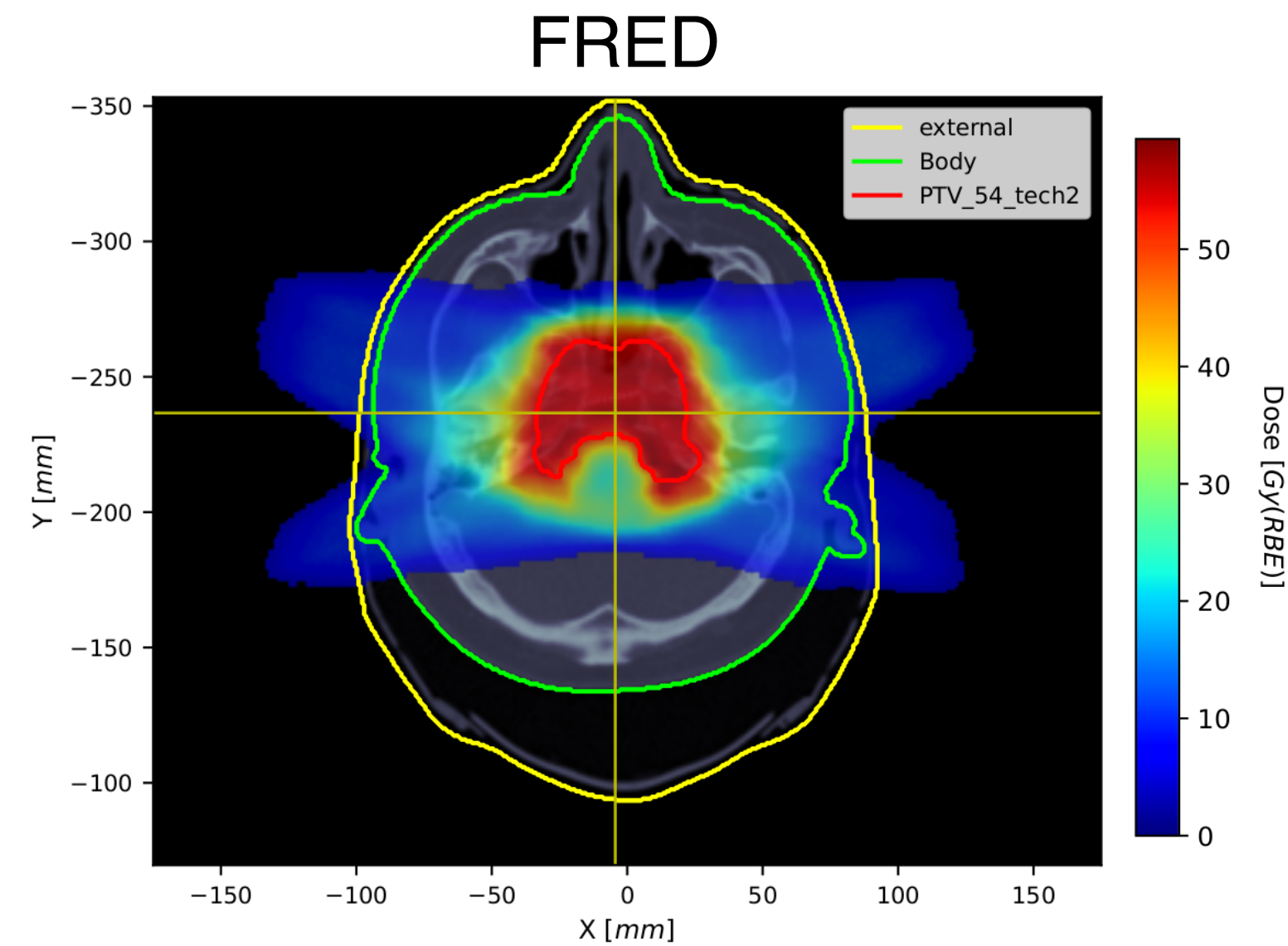
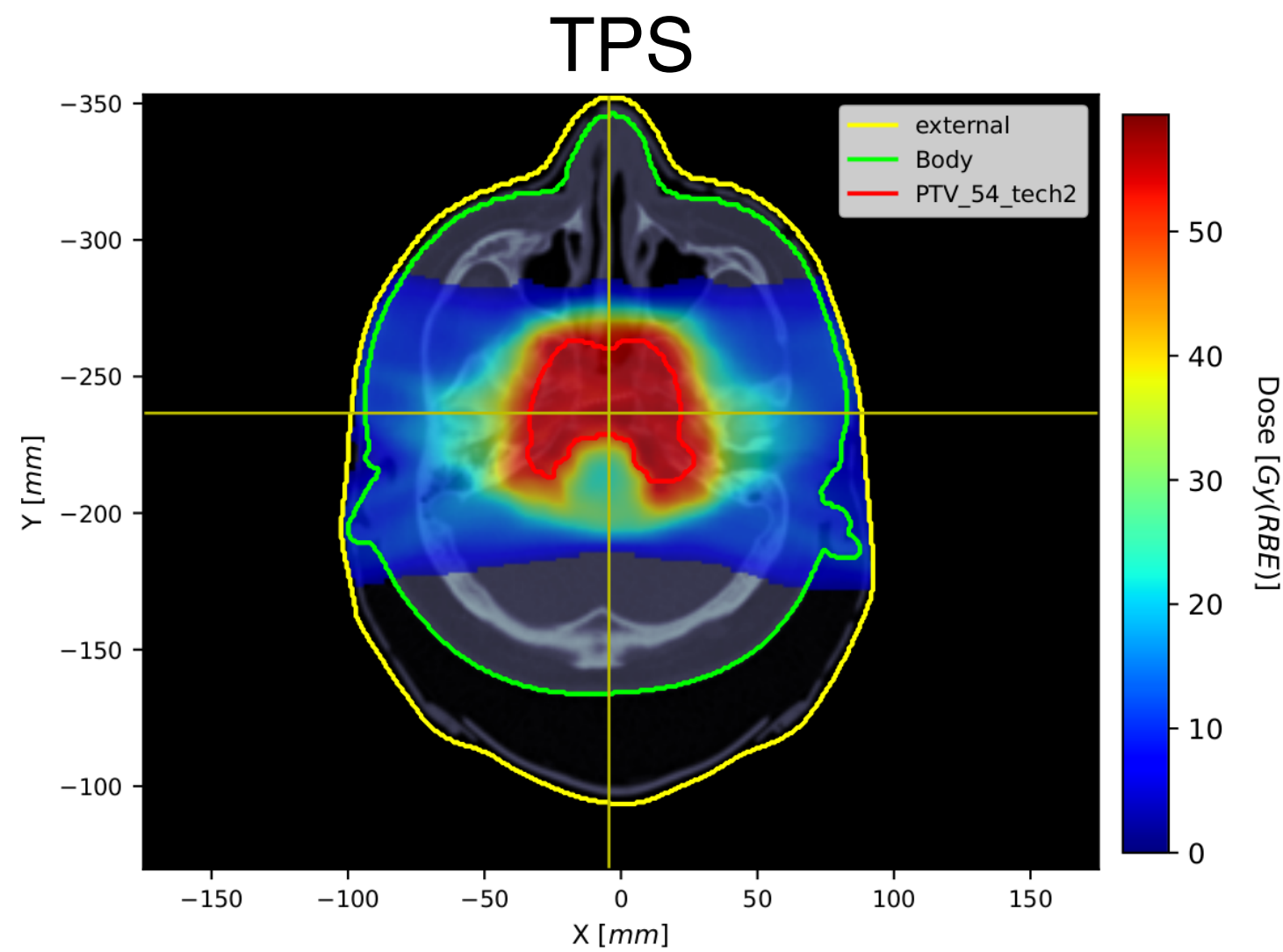
## Eclipse TPS, **Fred** and **GATE** MC codes

- Beam model based on 9 parameters for 17 beam energies:
- Initial energy, energy spread
- Lateral propagation (6 emittance parameters)
- Dosimetric calibration
- Range shifters
- Gantry/patient rotations + isocenter shifts



# Dosimetric cross-validation TPS-GATE-FRED

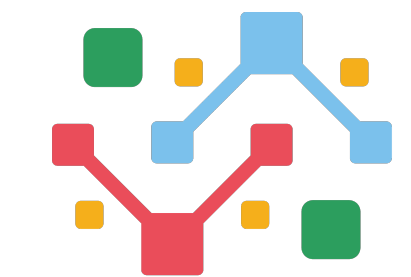
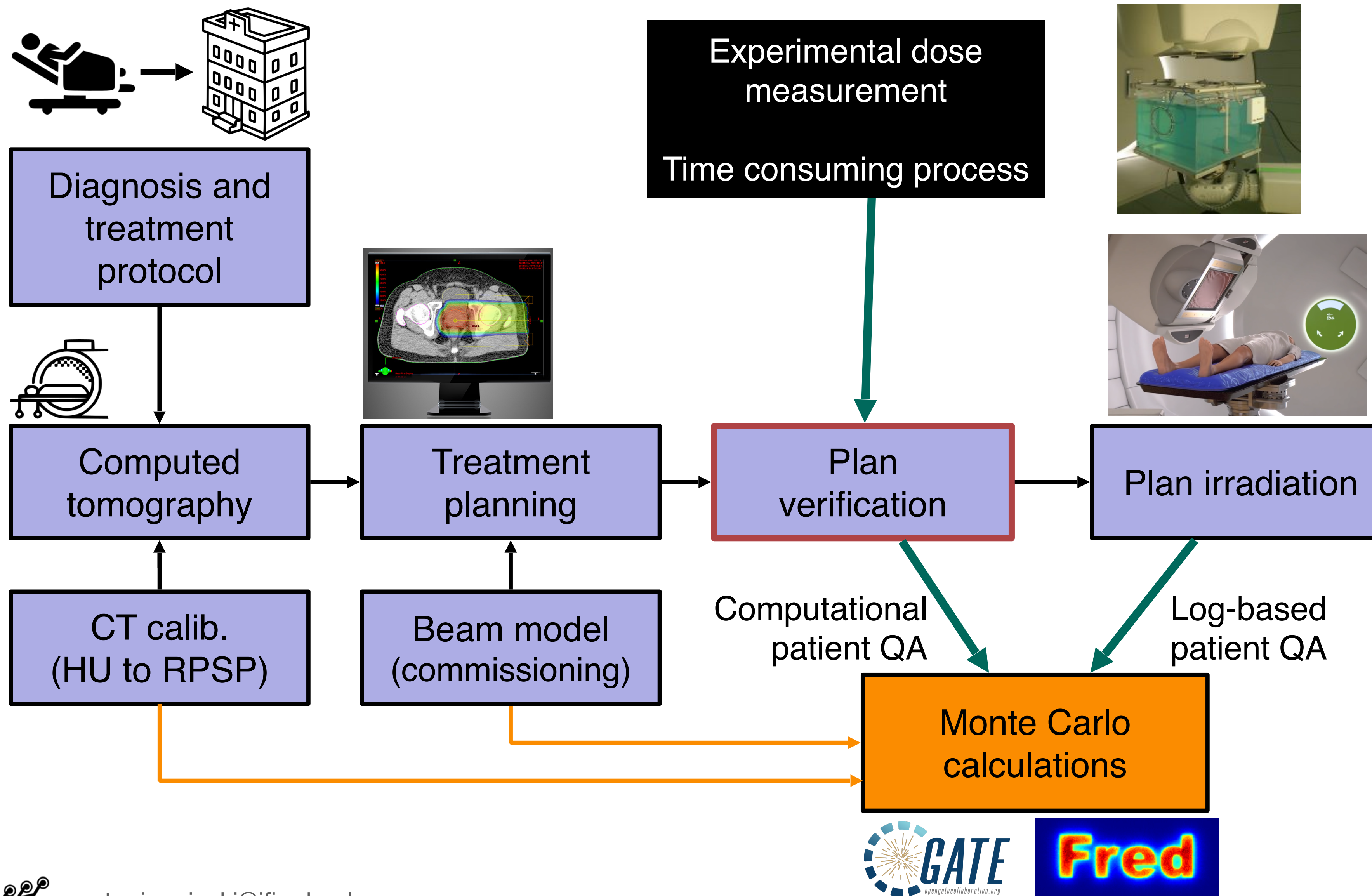
- 35 patient treatment plans
- precise physics
  - QGSP\_BIC\_HP\_EMZ
  - 0.1 mm prod. cuts
- GATE on ZIEMOWIT cluster



HPC Ziemowit

SAPIENZA  
UNIVERSITÀ DI ROMA

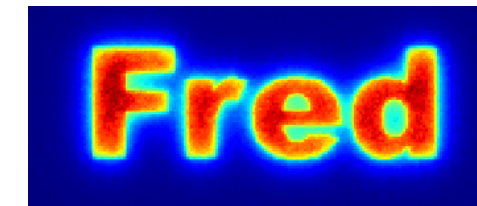
# Computational Patient QA with FRED and GATE



SCIENCE FOR THE SOCIETY

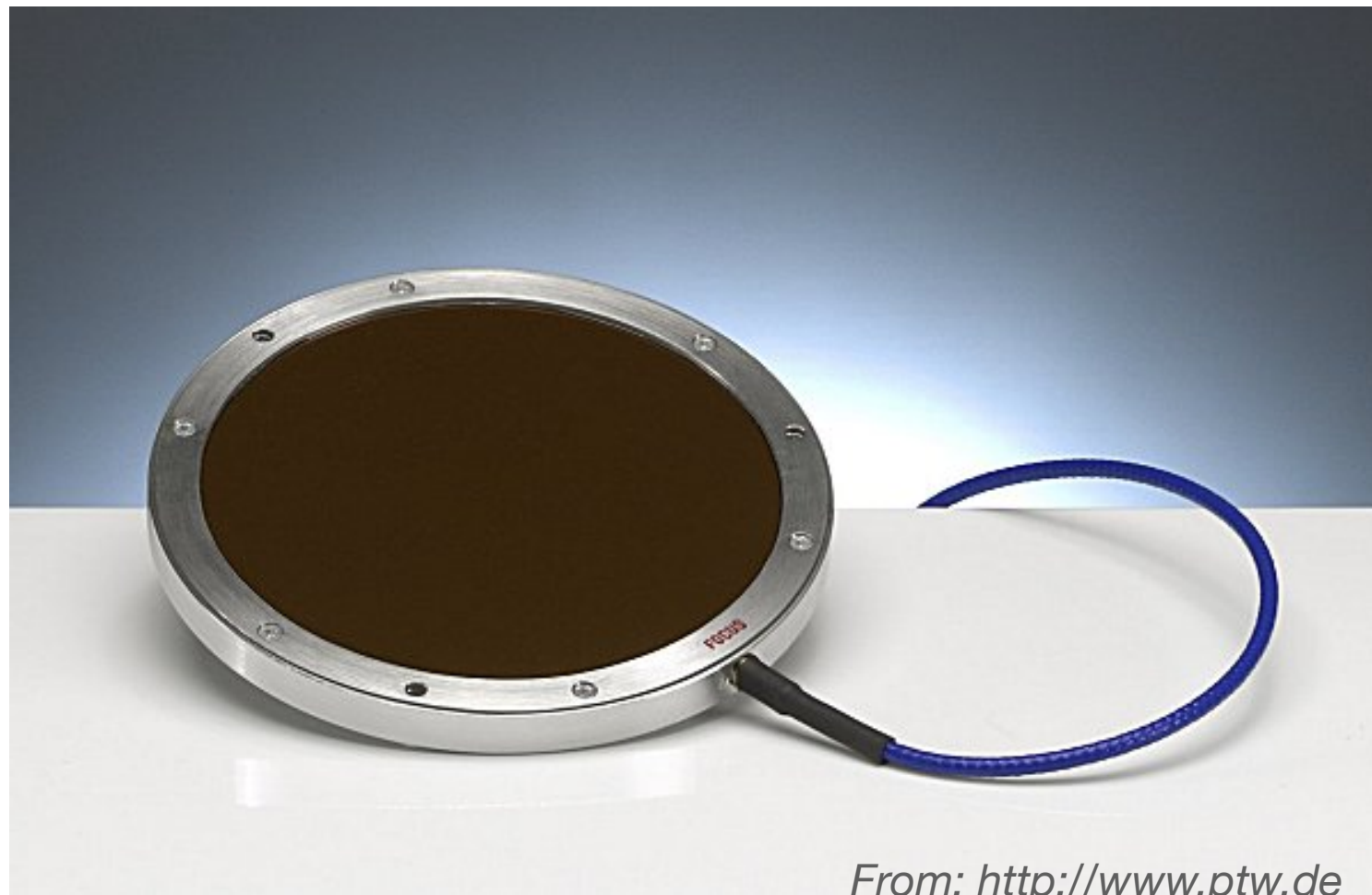
Grant for computational patient QA workflow implementation + simultaneous LET and dose optimisation

myQA® iON

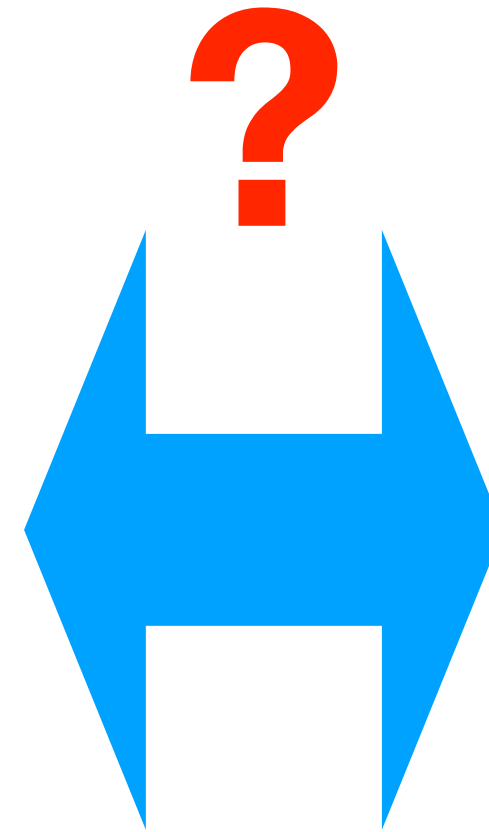


# Advances in radiobiological modeling

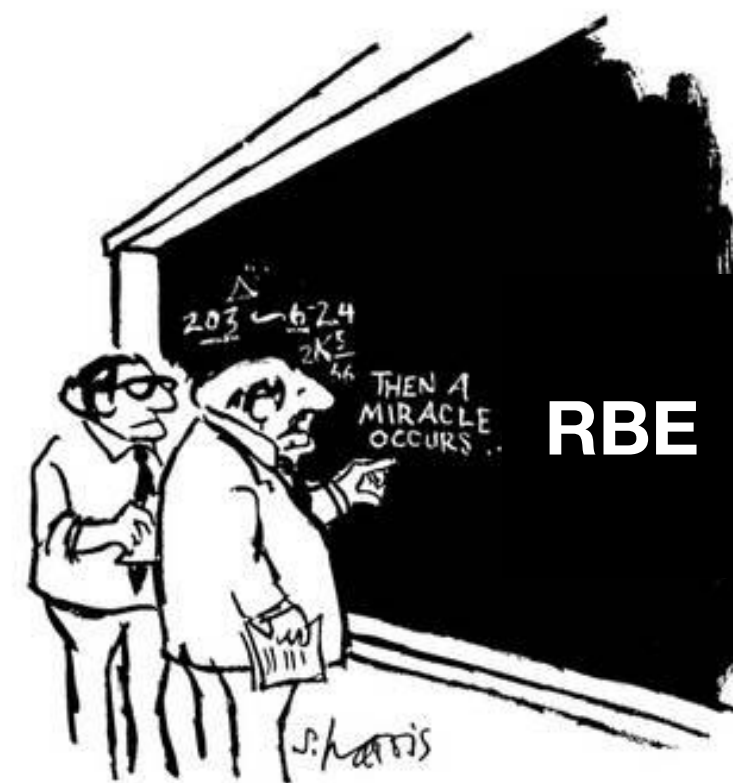
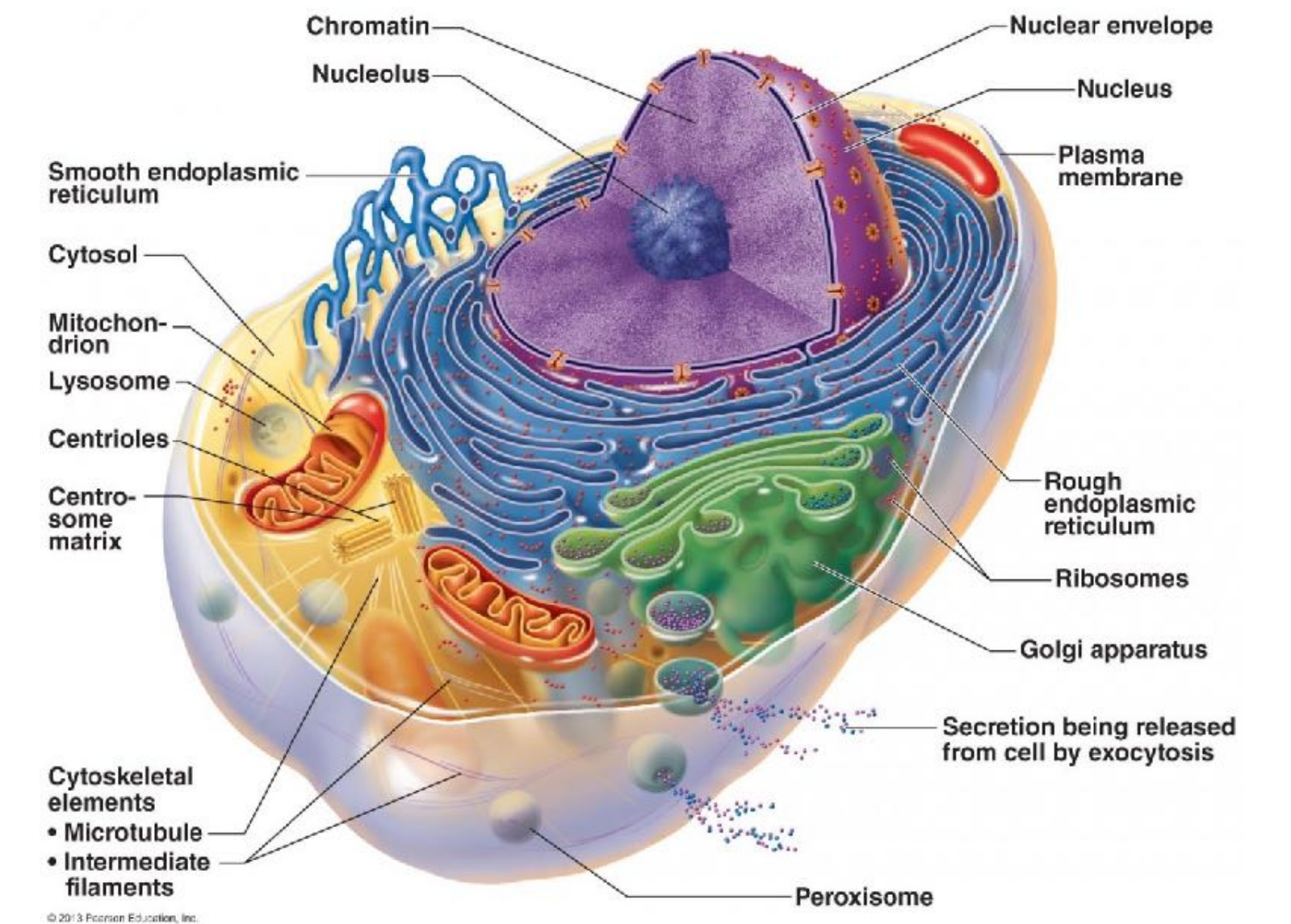
## Dose: Macroscopic concept



RBE modeling



## Cell inactivation: Microscopic concept



"I think you should be more explicit here in step two."

**RBE modeling:**  
 - Phenomenological  
 - Microdosimetric

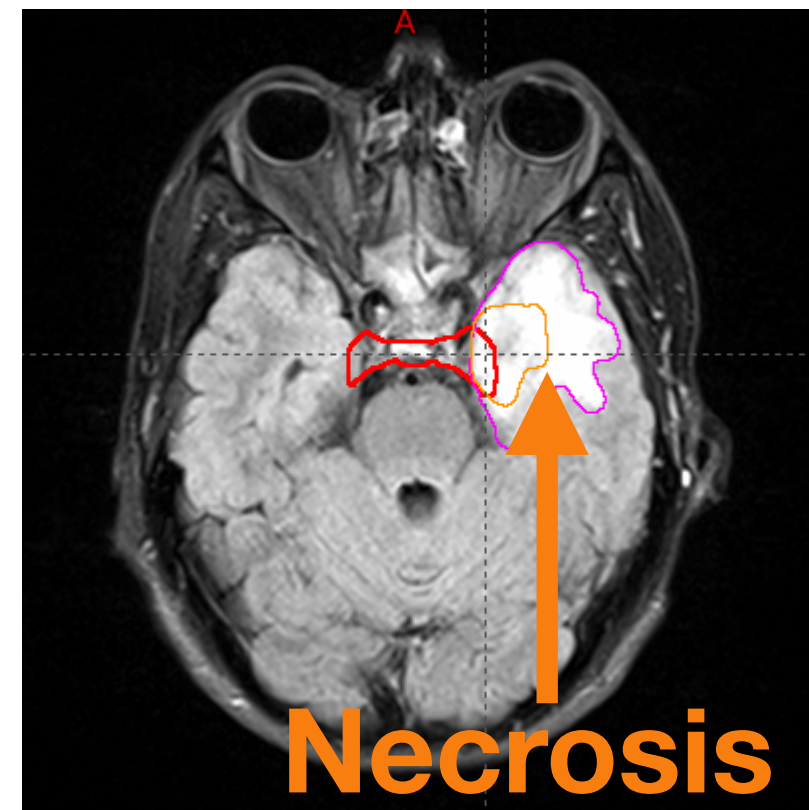
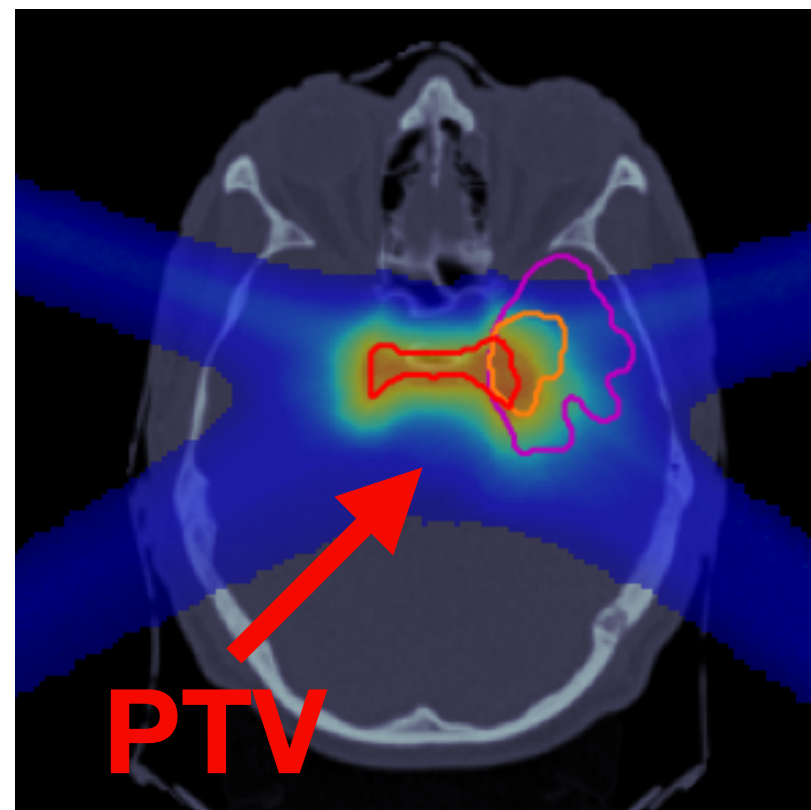


# MC to support LET and biologically weighted TP

Chordoma, 74 Gy(RBE), 2Gy(RBE)/Fx, 4 radiation fields

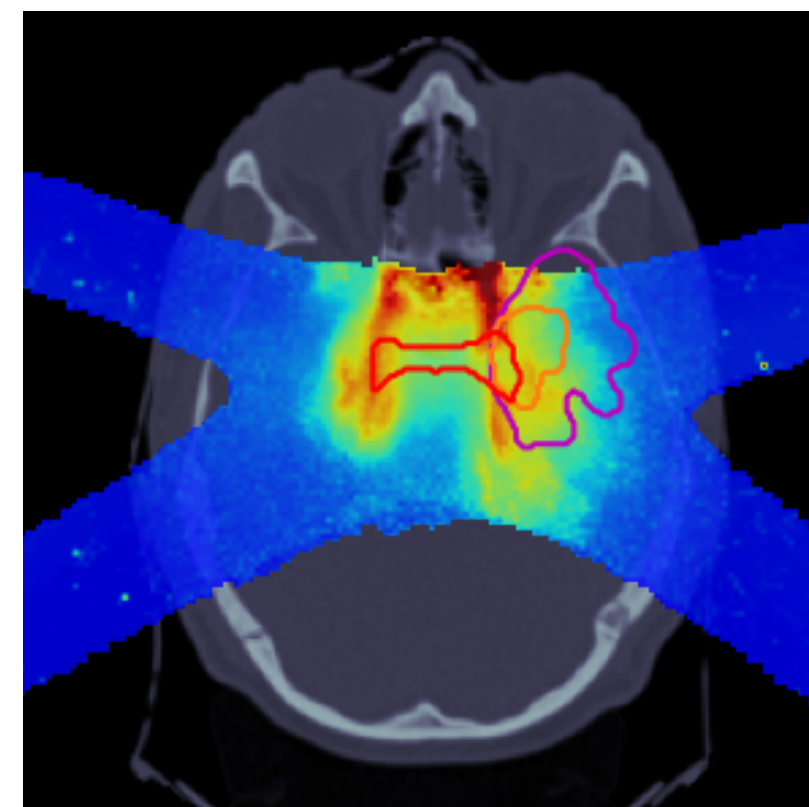
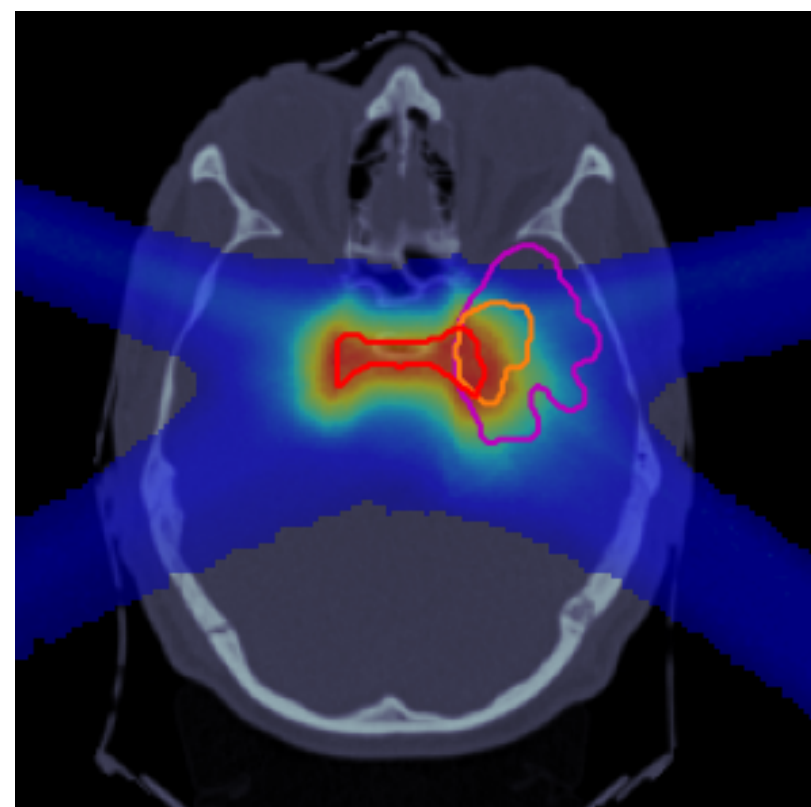
Dose RBE=1.1

Post treatment MRI



RBE-weighted dose\*

LET<sub>d</sub>



## • Hypothesis

- LET spectrum (not LET<sub>d</sub>) in a voxel is essential for biological effect
- High-LET particles cause complex DNA damage

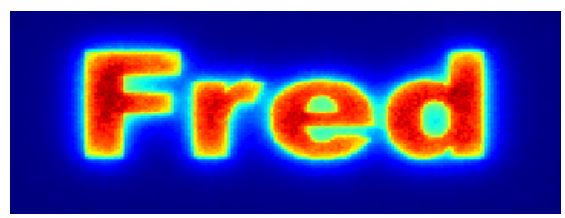
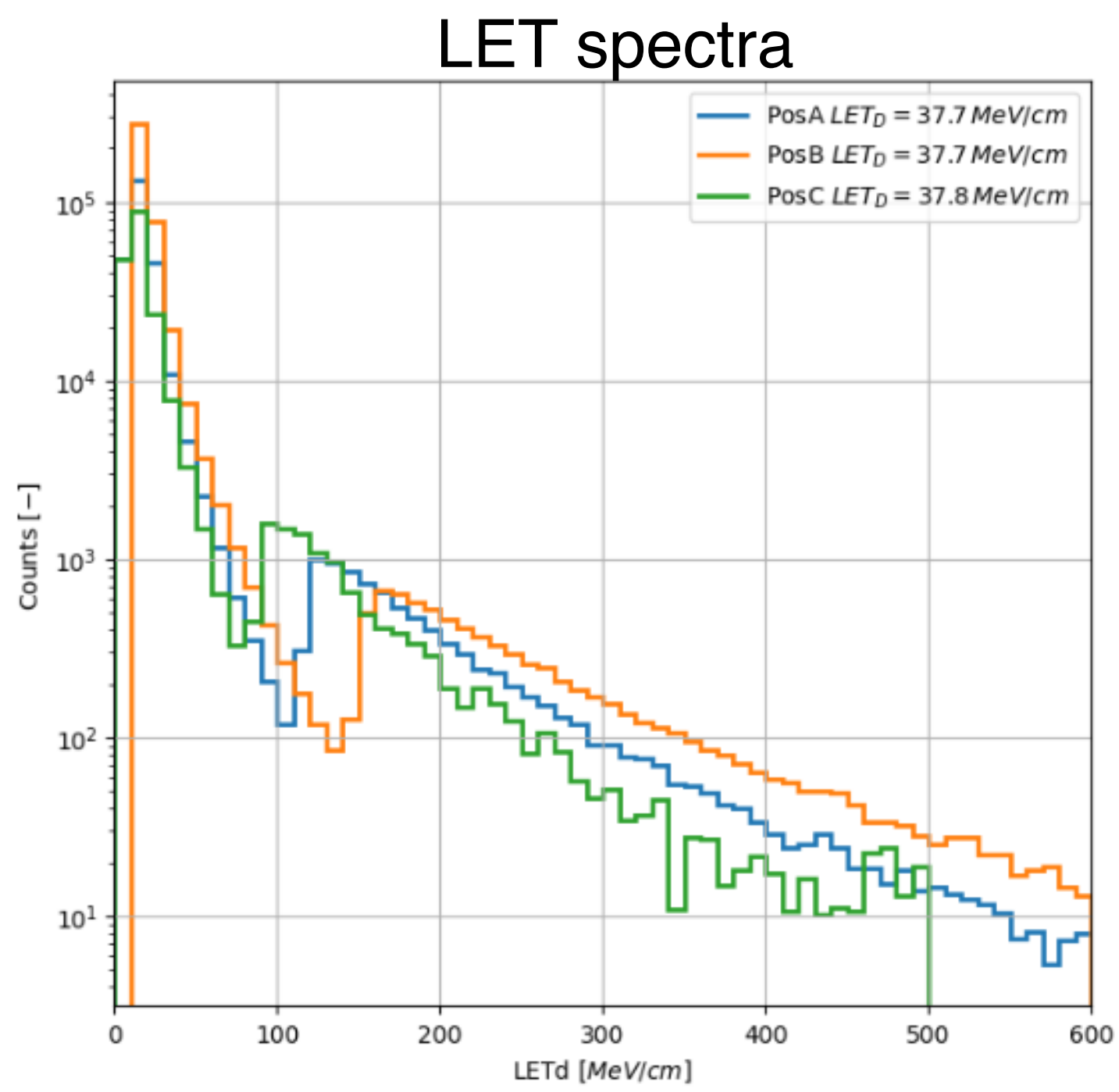
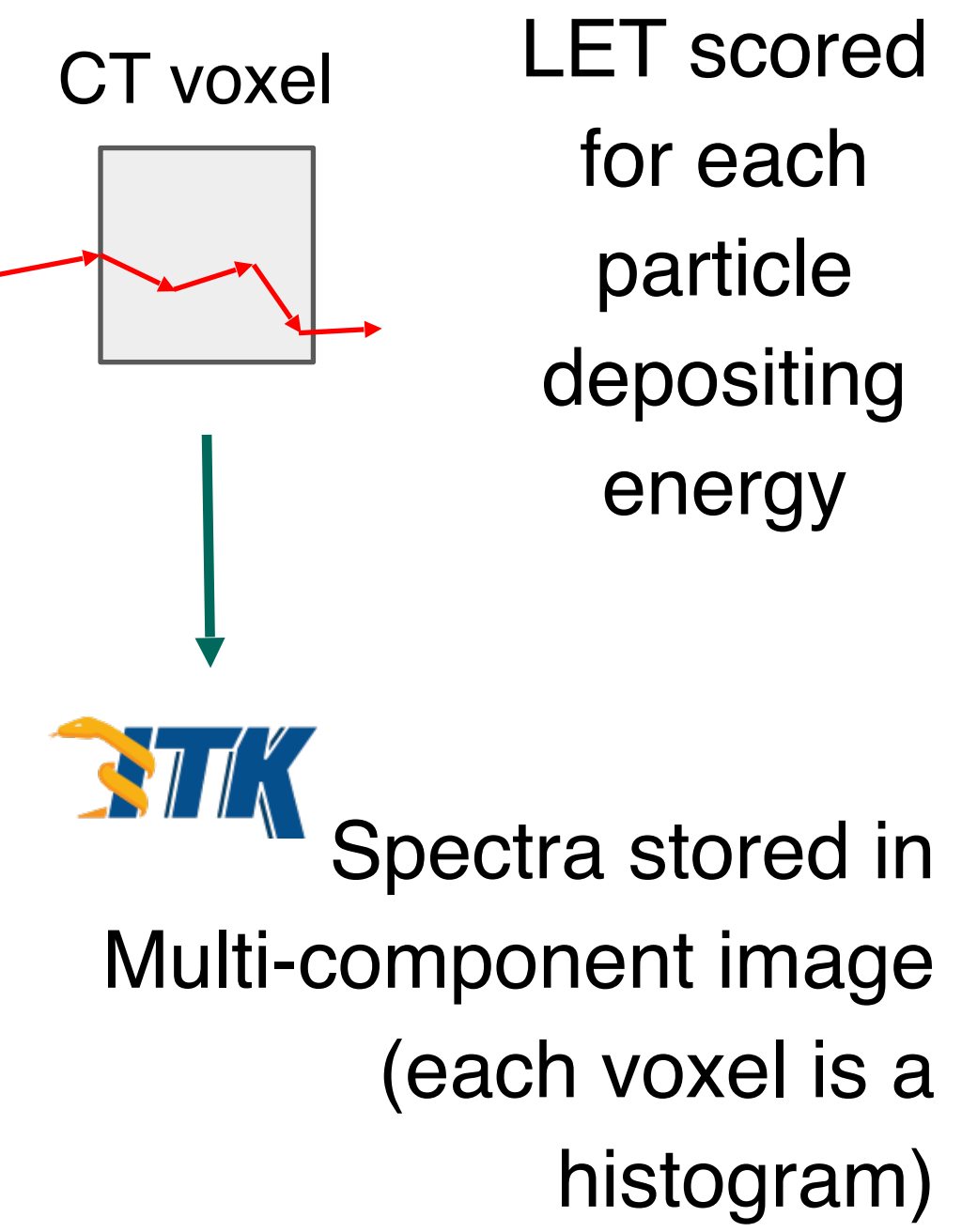
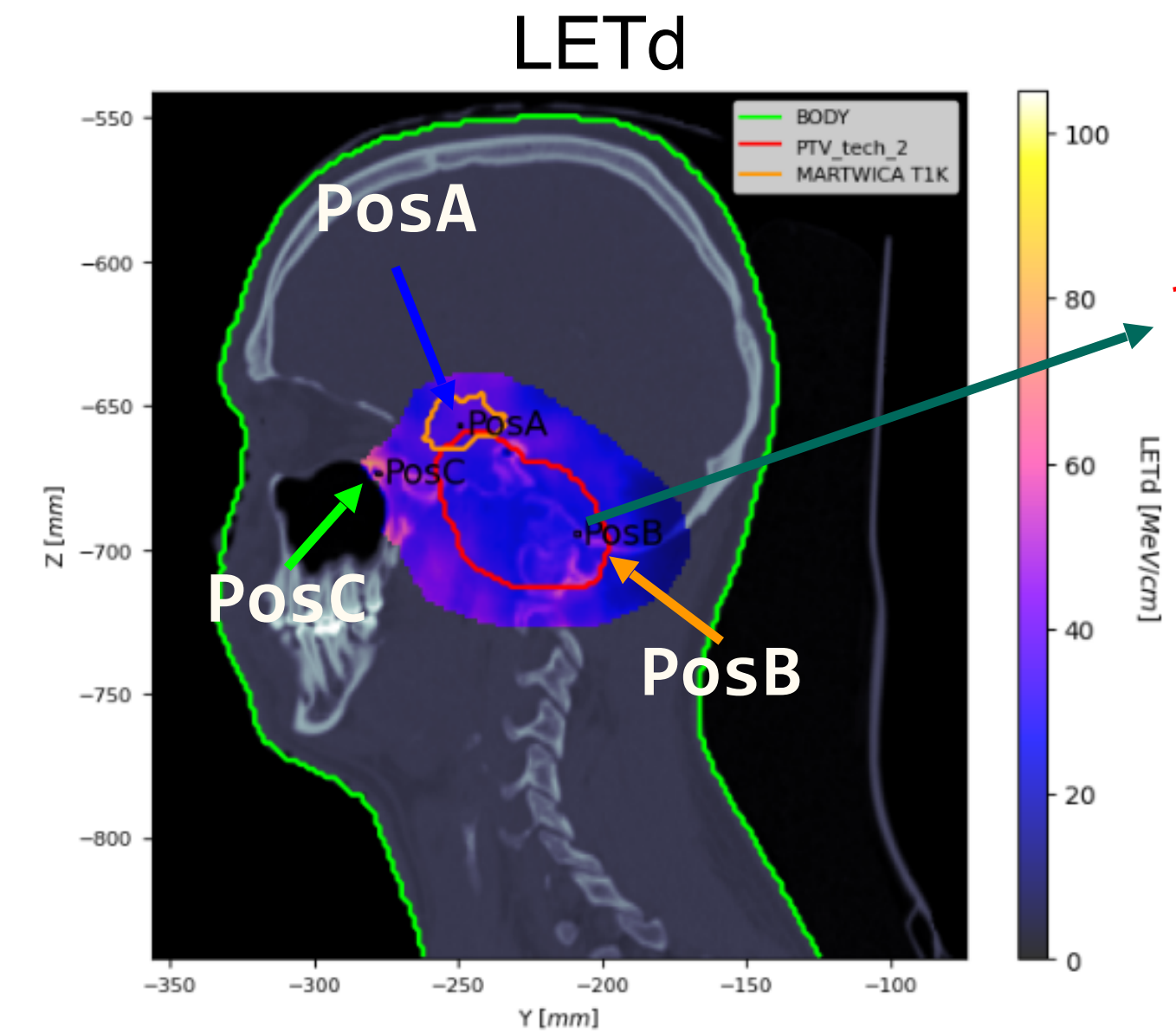
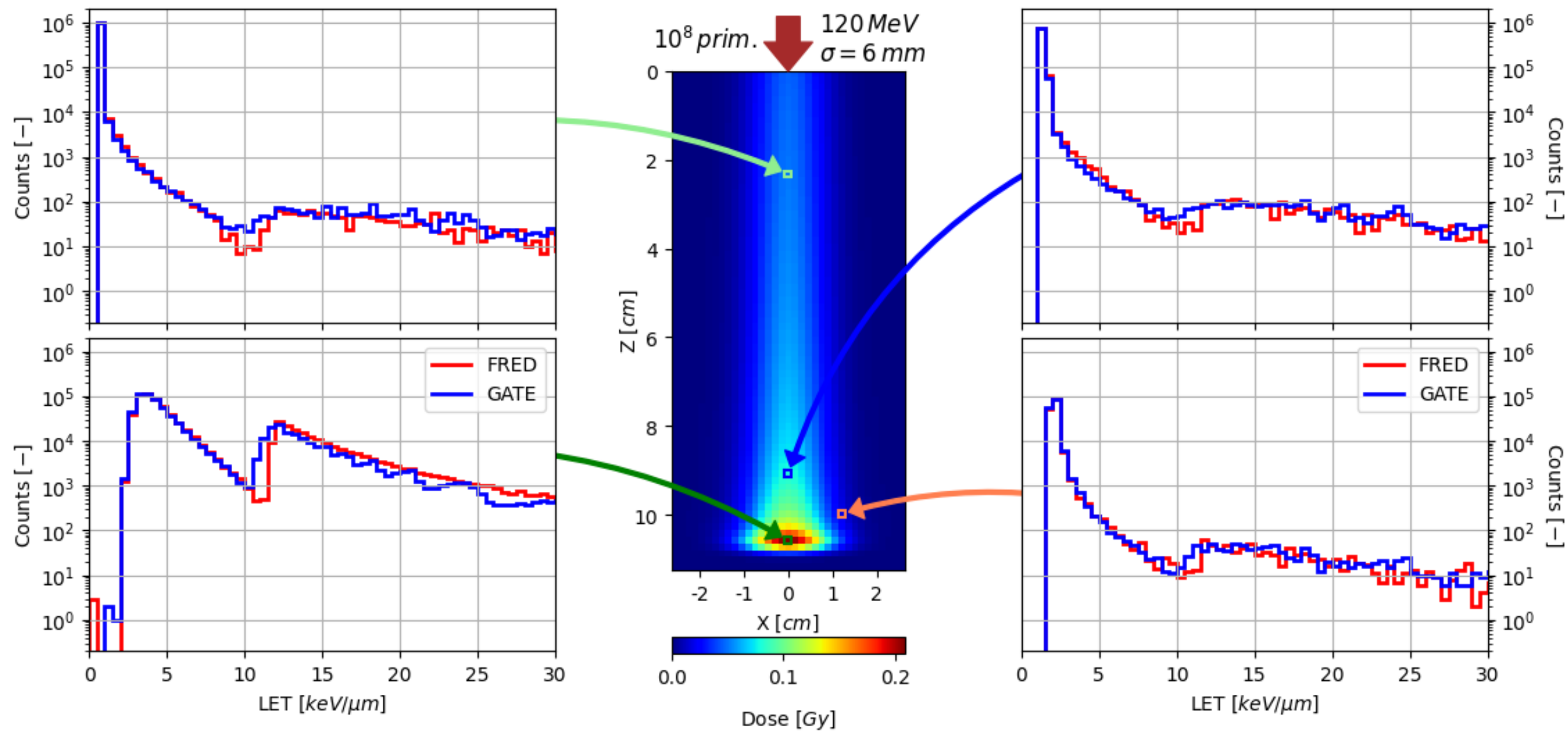
## • Impact/Application

- Commissioning and validation of MC-TPS
- Biologically and LET-weighted treatment planning

\*Dose computed with GPU-accelerated MC code (FRED) according to variable RBE model proposed by (A McNamara et al. 2015, PMB)

# Computational LET QA

Not only the averaged LET (e.g. LETd) but the whole LET spectra in a voxelized geometry



Collaboration




SAPIENZA  
UNIVERSITÀ DI ROMA



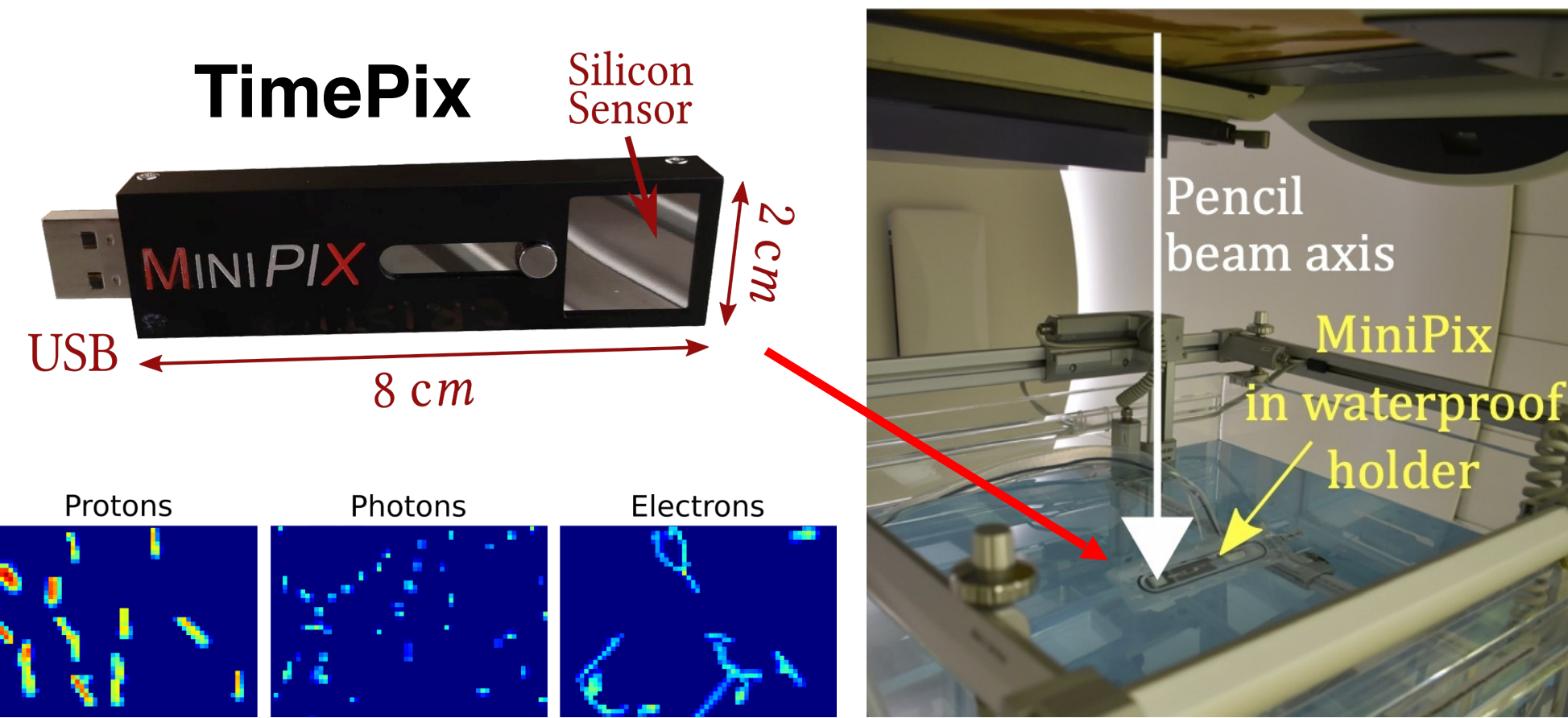
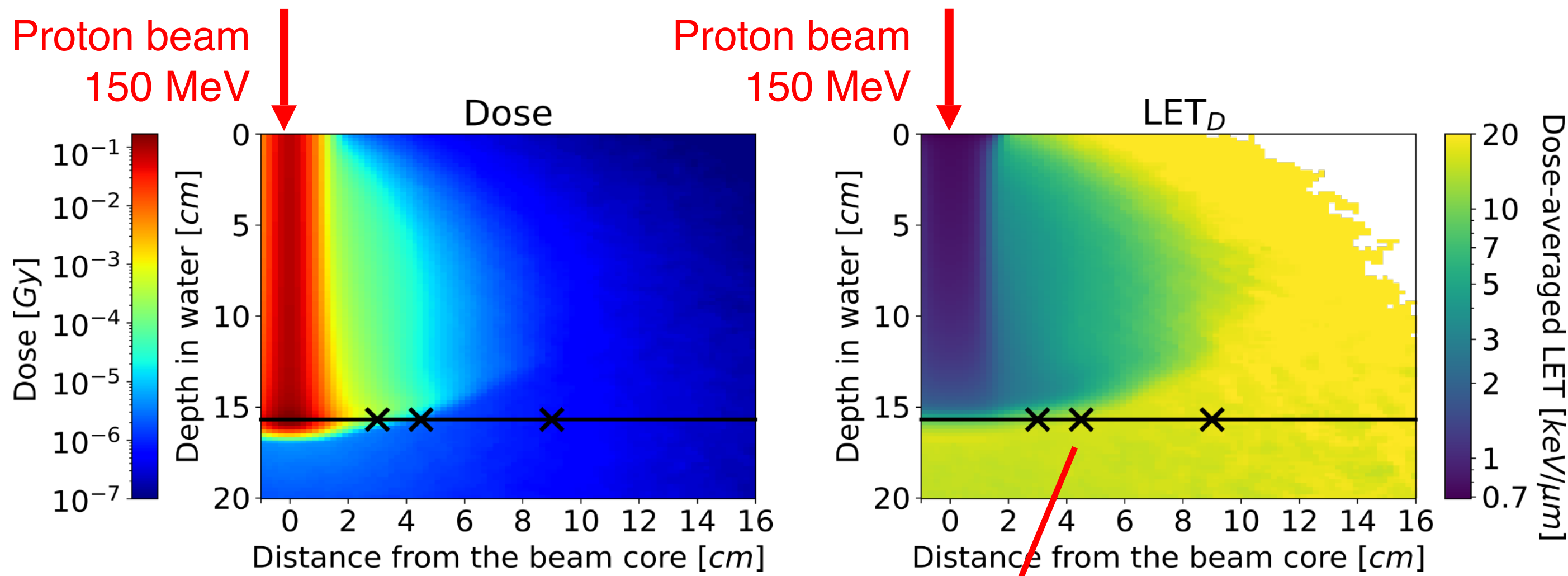
Grant for characterisation of mixed radiation fields in proton therapy



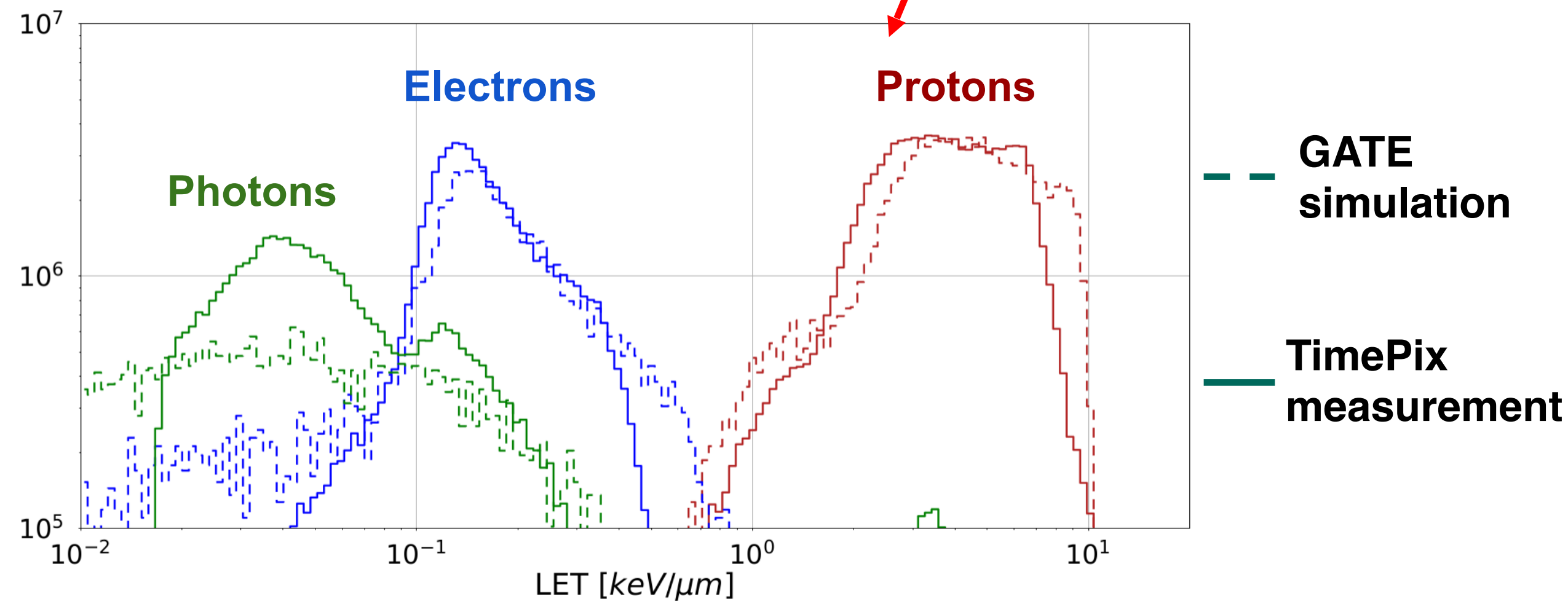
antoni.rucinski@ifj.edu.pl



# Experimental LET QA



- Monte Carlo for detector development
- Depositions in TimePix simulated in GATE and FRED
- TimePix response with Geant4-based AllPix<sup>2</sup>



Collaboration

ADVACAM  
Imaging the Unseen



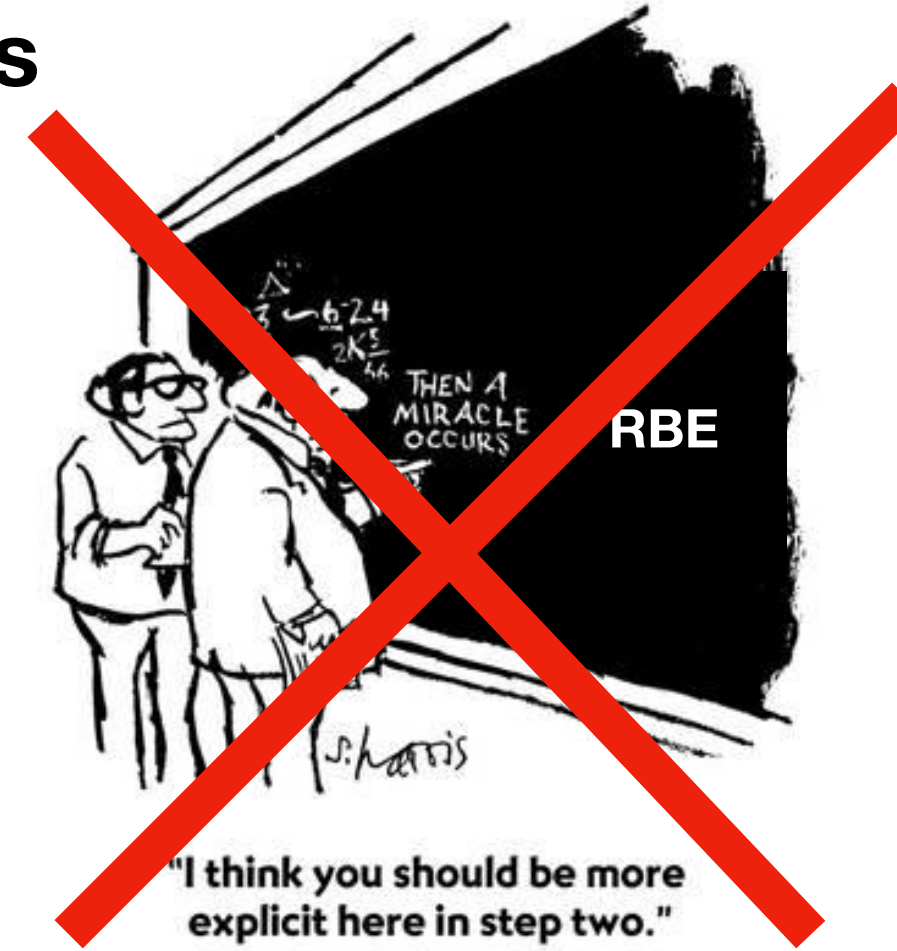
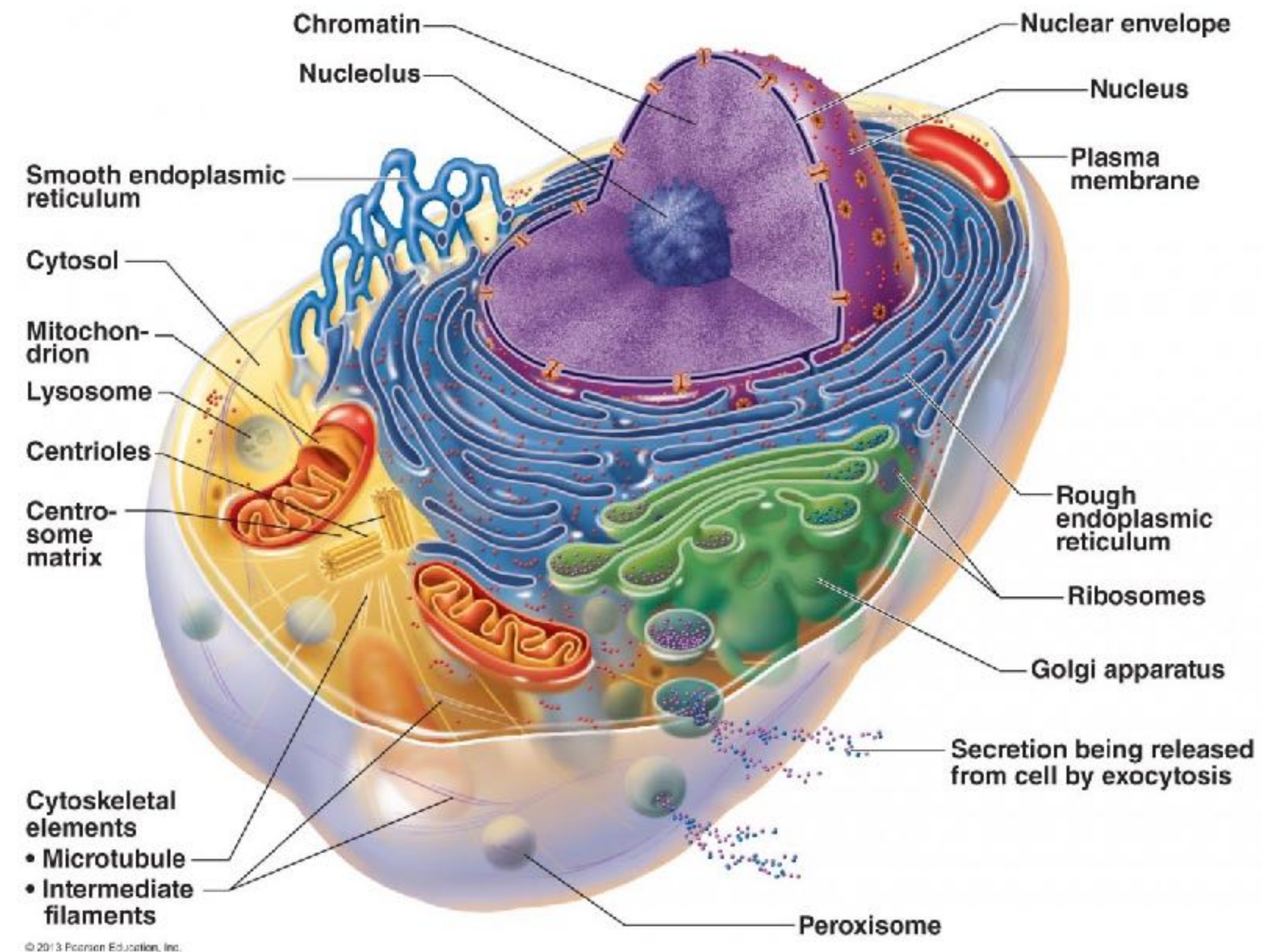
The National Centre  
for Research and Development

Grant for characterisation of  
mixed radiation fields in proton  
therapy

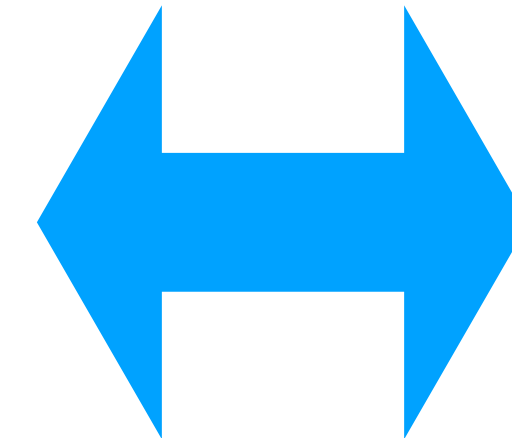
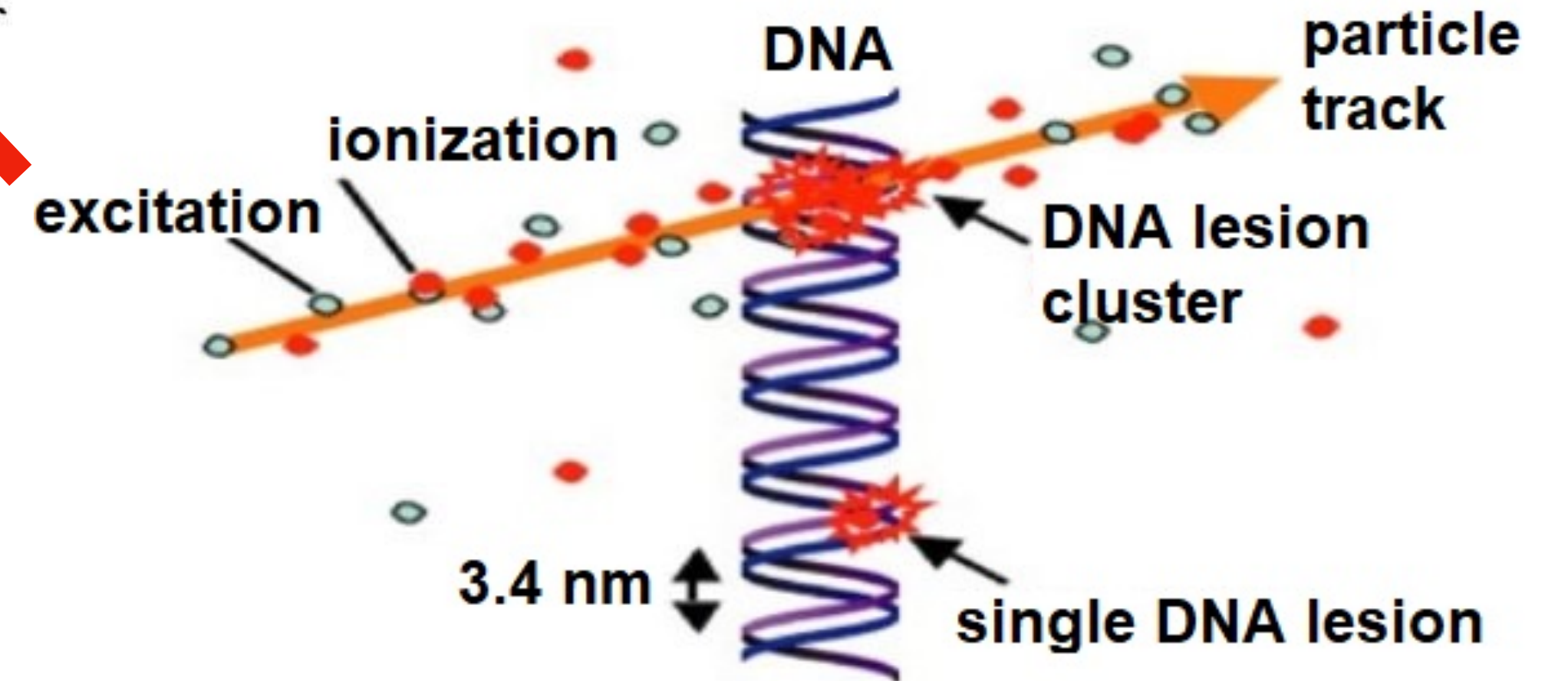
# Dose becomes inadequate at microscopic levels

The hypothesis: Similar clustering leads to similar biological effect

Cell/DNA damage: Micro/nano-scopic concepts



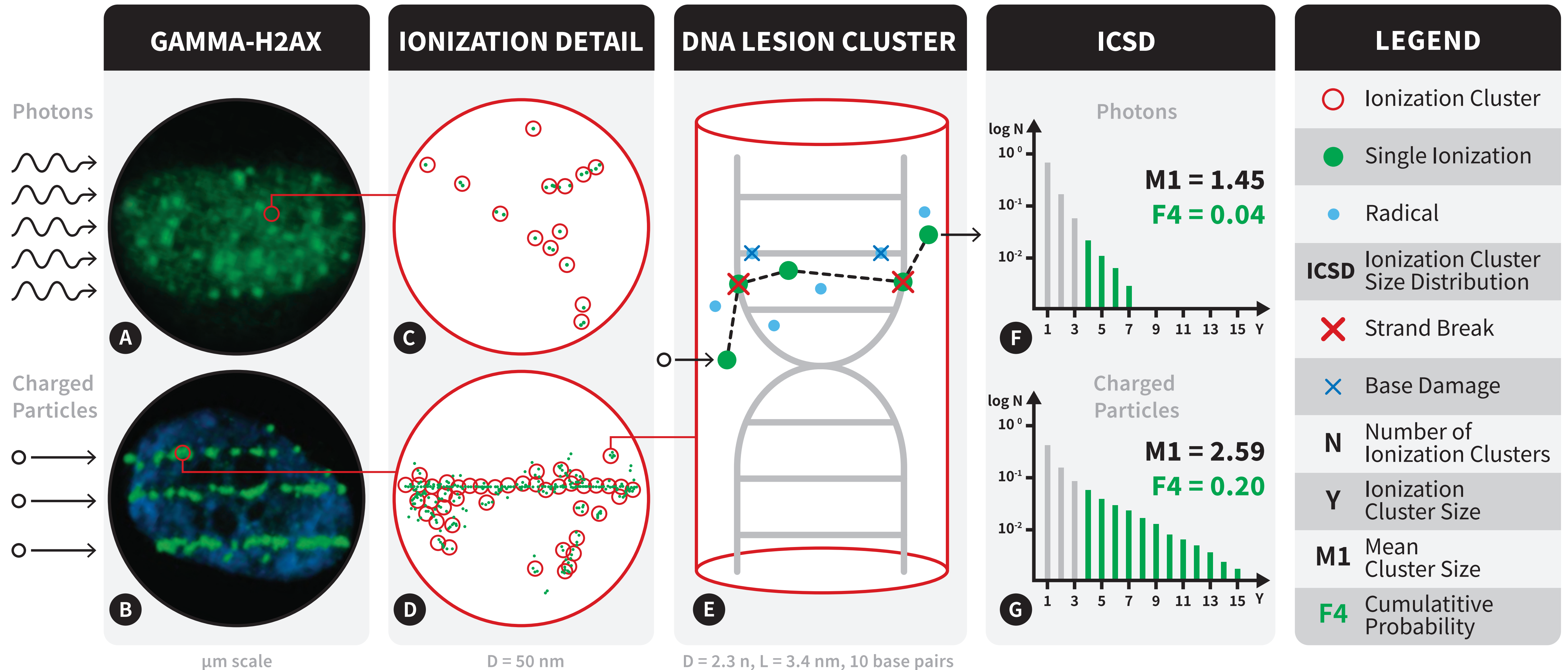
Local ionization cluster size on the nanometer scale is the starting condition for all subsequent processes that lead to the observed DNA damage



- Radiation damage is a stochastic quantity on the micro- and nanoscopic level, therefore the modeling solution must use micro- and/or nanodosimetric distributions rather than average quantities (LET, absorbed dose).

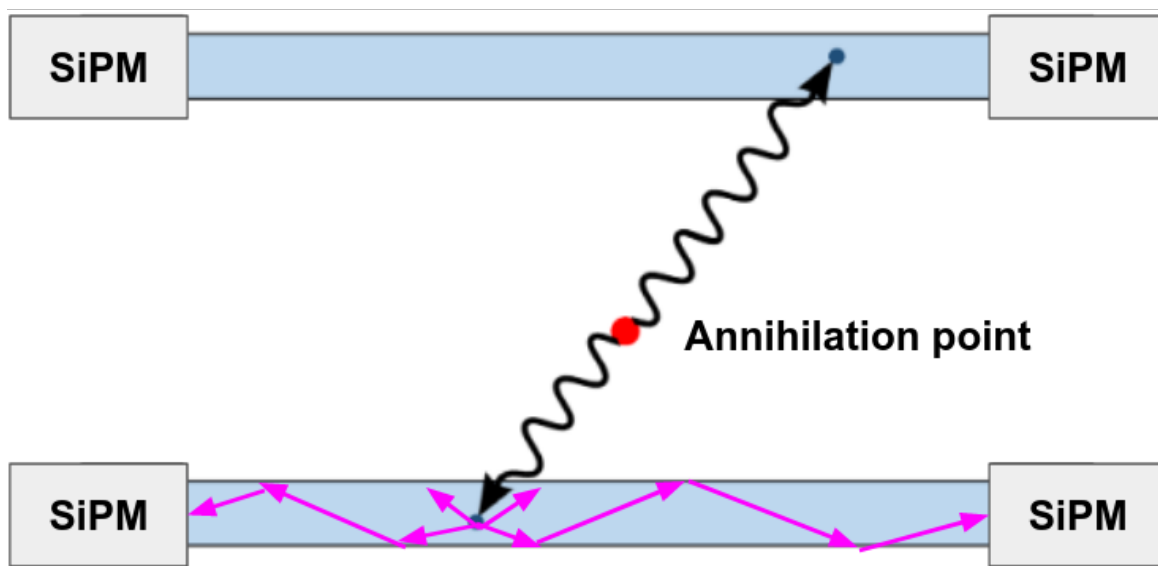
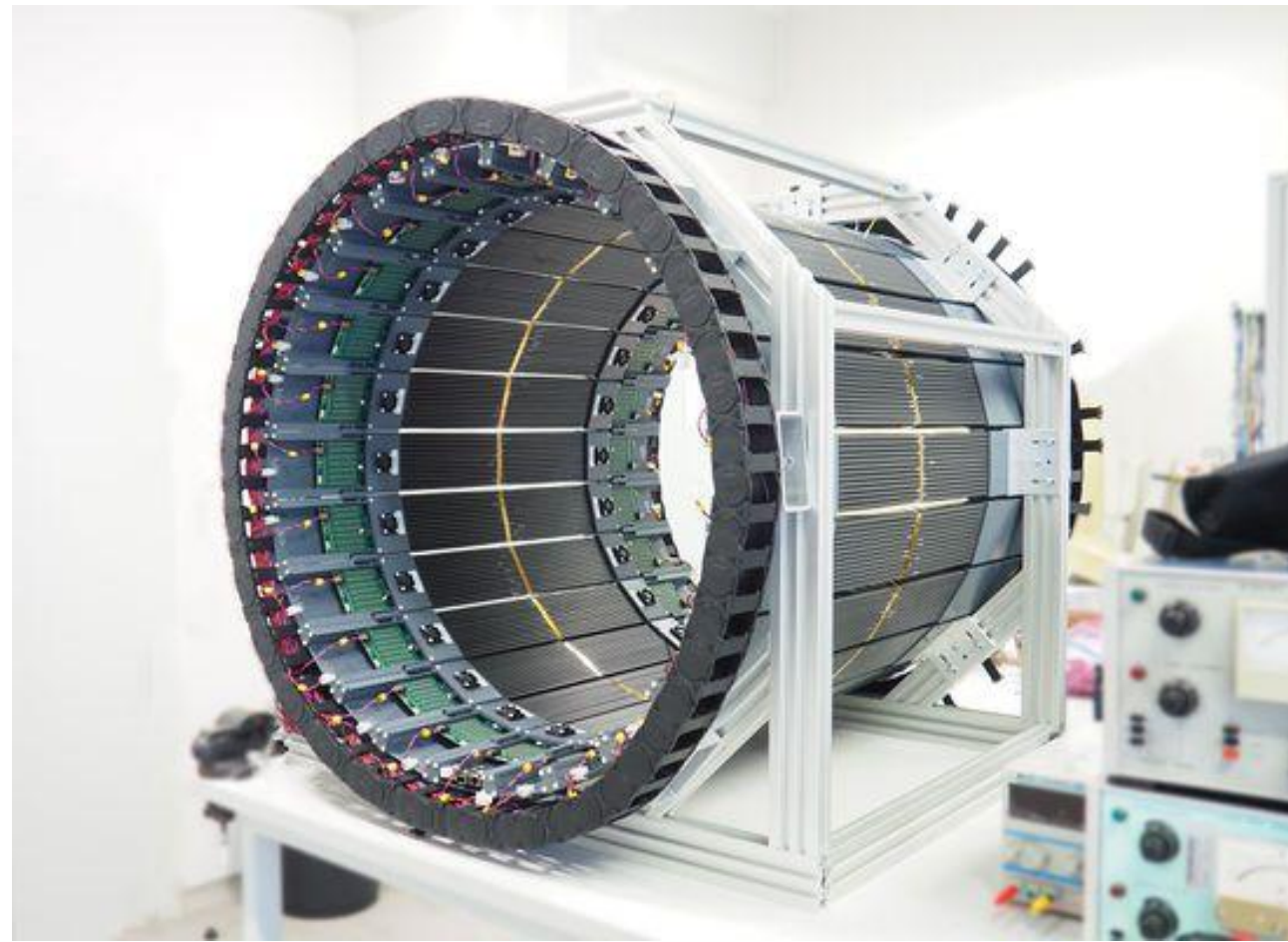
# Ionization detail: principles and concepts

ProtoTerra PhD students work with LLU, CA, USA & Krakow on ID detector & TP approach

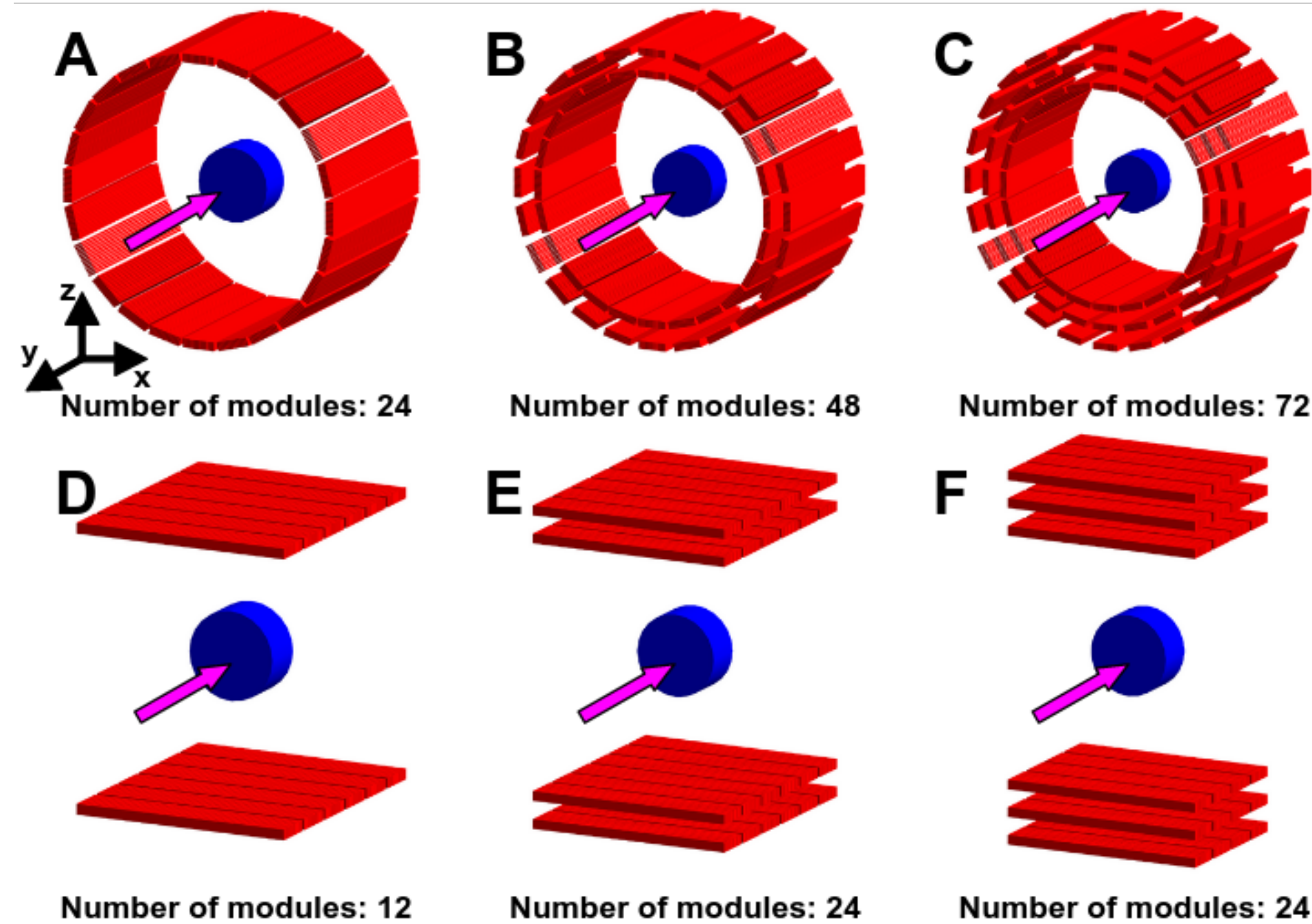


# J-PET for range monitoring in PT

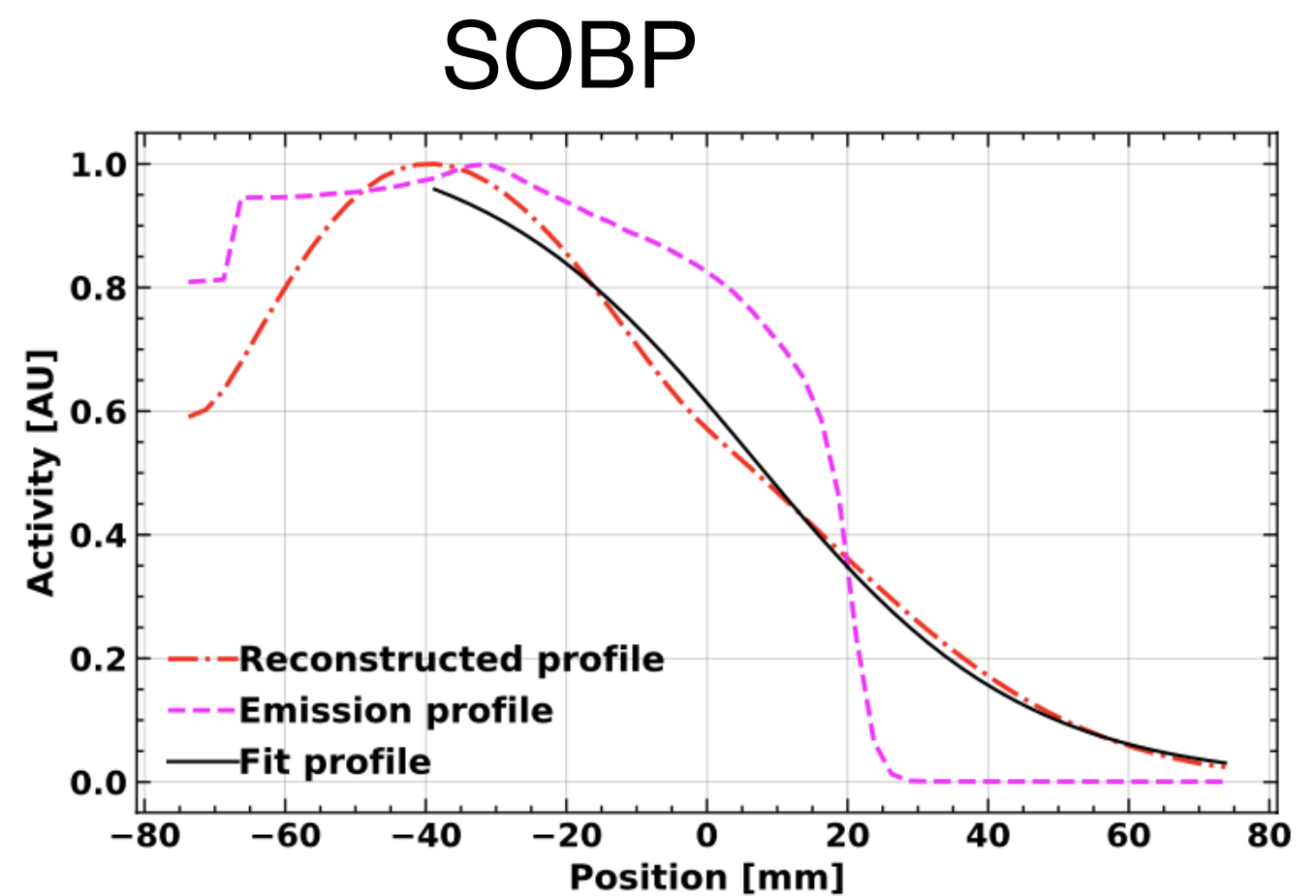
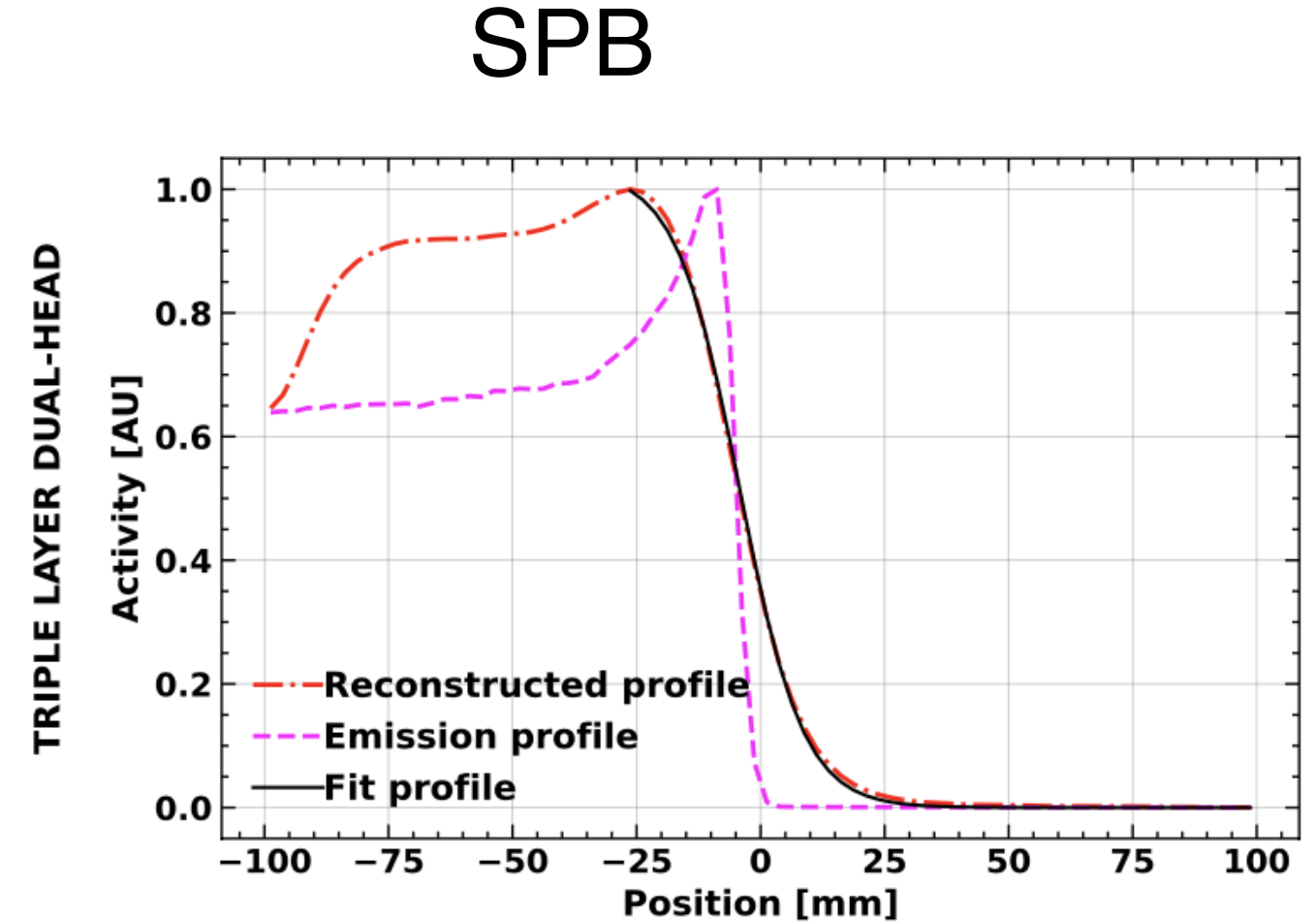
J-PET operation principle



Sensitivity

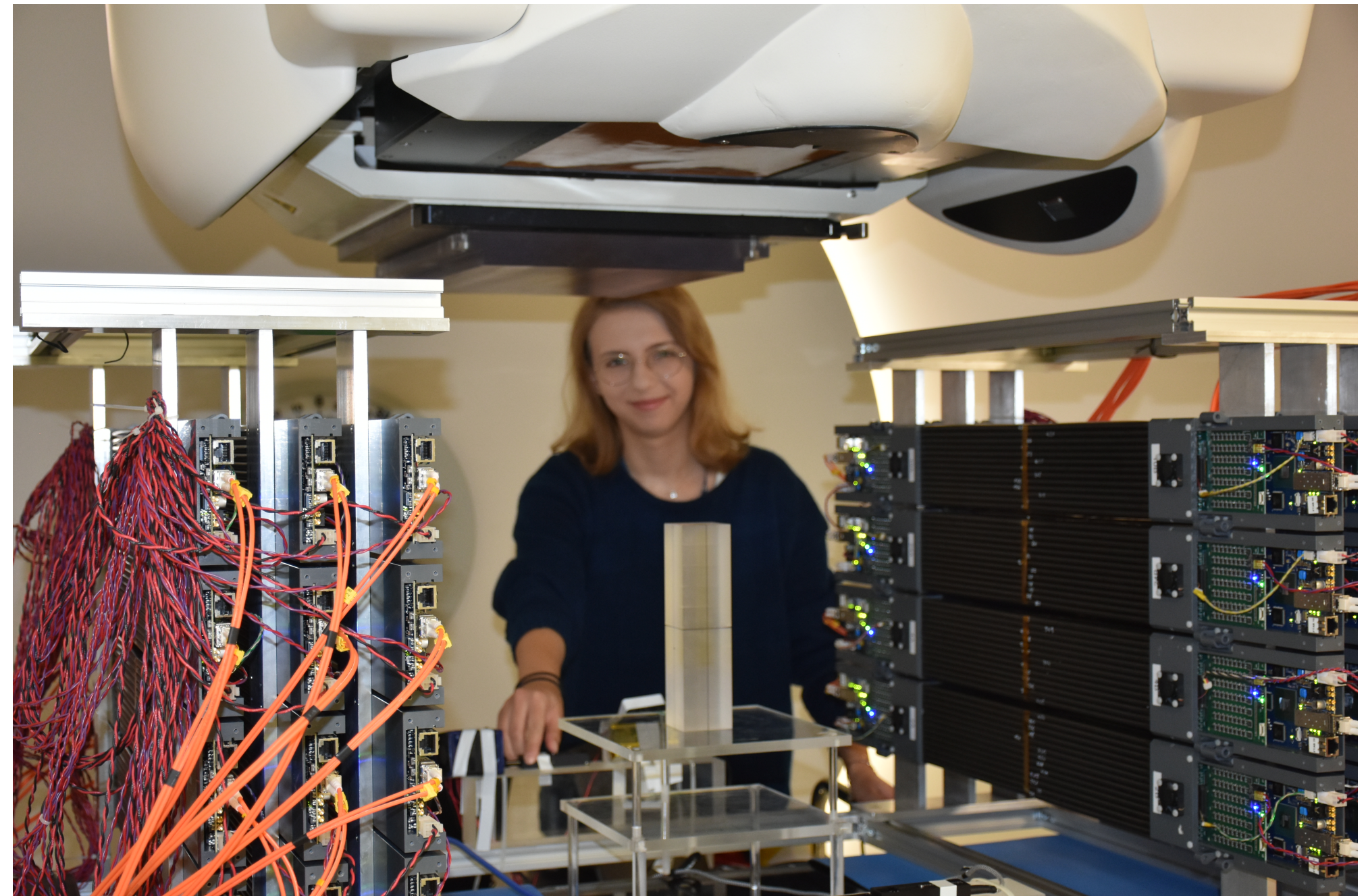
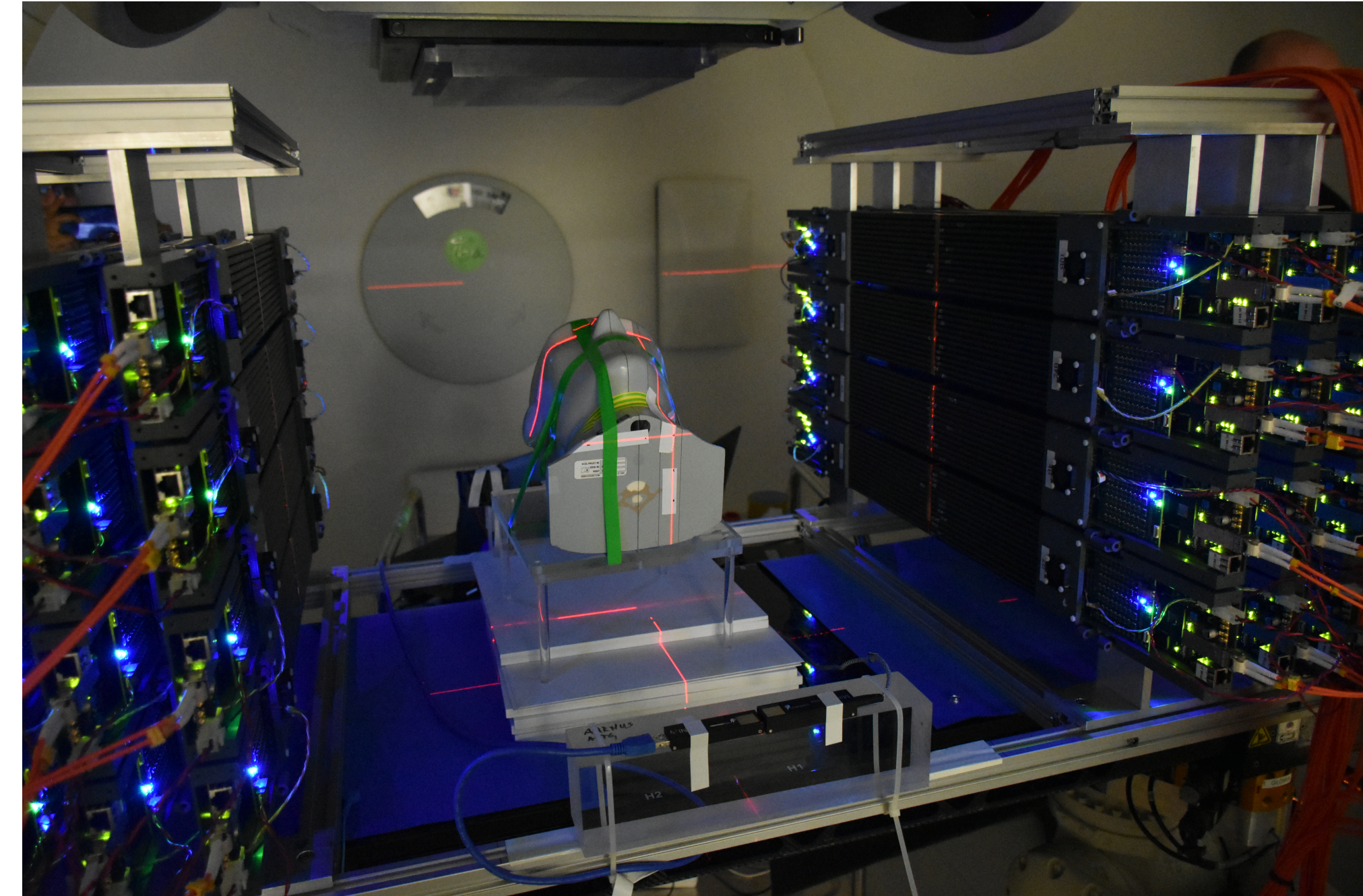


Range estimation



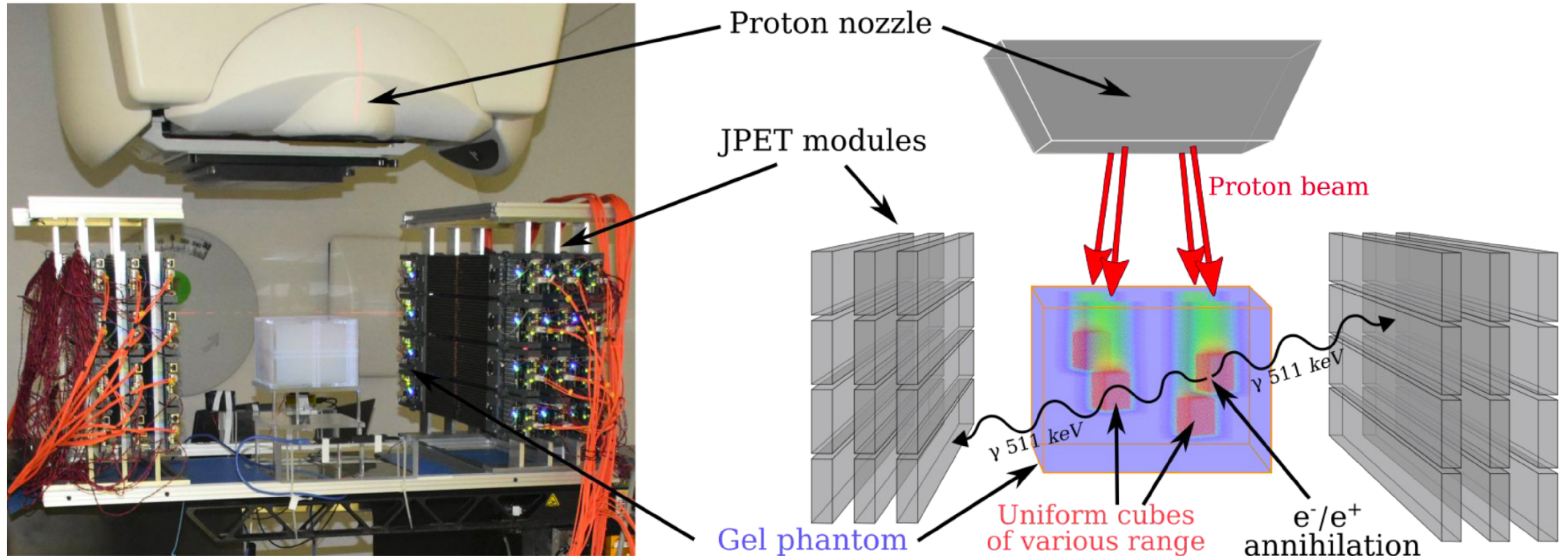
Setup	SPB			SOBP		
	$\eta[10^{-6}]$	$\sigma(\eta)[10^{-6}]$	$\bar{H}$	$\eta[10^{-6}]$	$\sigma(\eta)[10^{-6}]$	$\bar{H}$
Single layer cylindrical	9.45	0.29	1.0	3.64	0.22	1.0
<b>Double layer cylindrical</b>	<b>27.41</b>	<b>0.80</b>	<b>2.9</b>	<b>10.76</b>	<b>0.65</b>	<b>2.9</b>
Triple layer cylindrical	45.72	1.26	4.8	18.00	1.11	5.0
Single layer dual-head	3.79	0.13	0.4	2.45	0.19	0.7
Double layer dual-head	10.55	0.35	1.1	7.21	0.56	2.0
Triple layer dual-head	10.22	0.26	1.1	8.92	0.78	2.4

# J-PET & proton beams 2021



# J-PET for range monitoring in PT

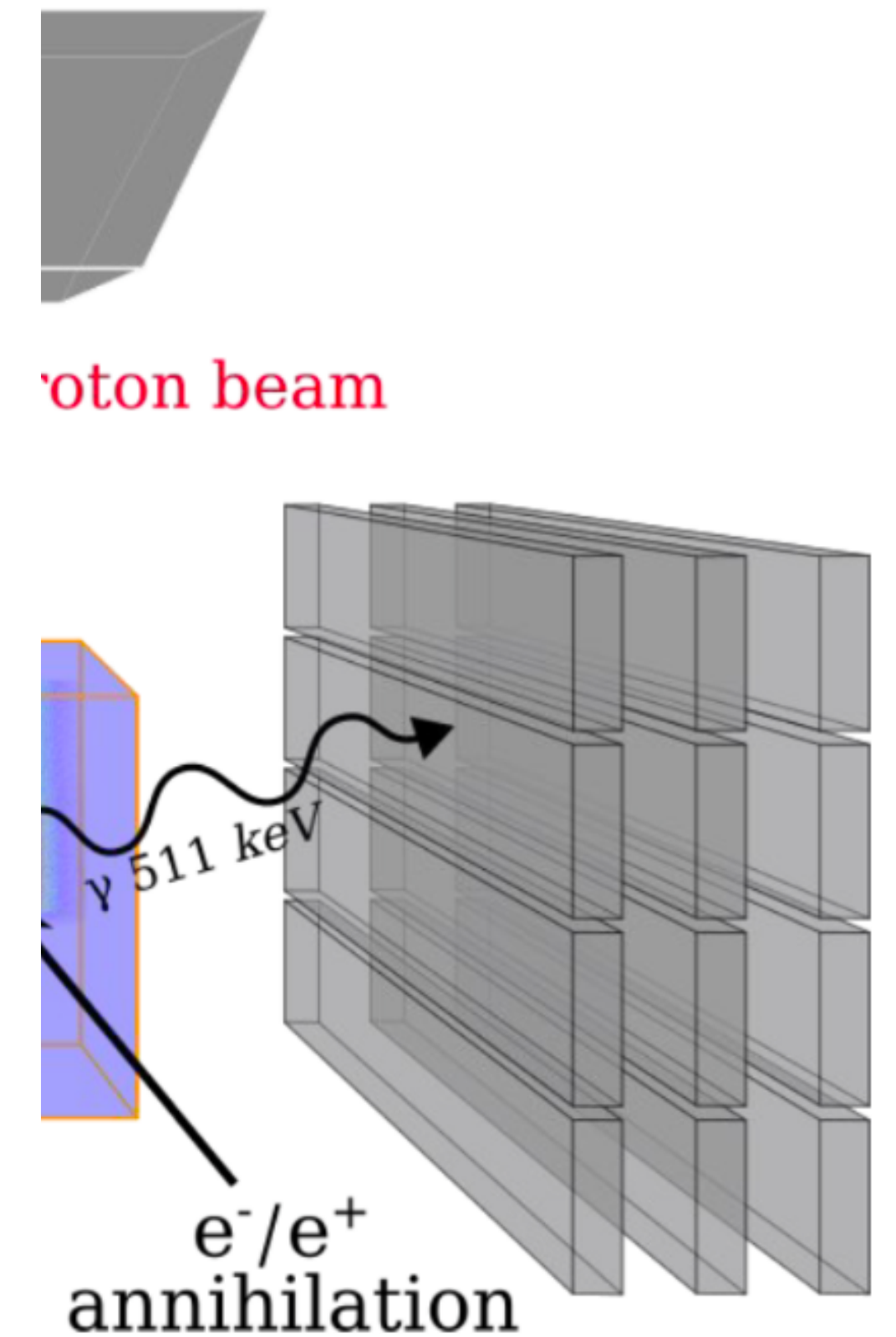
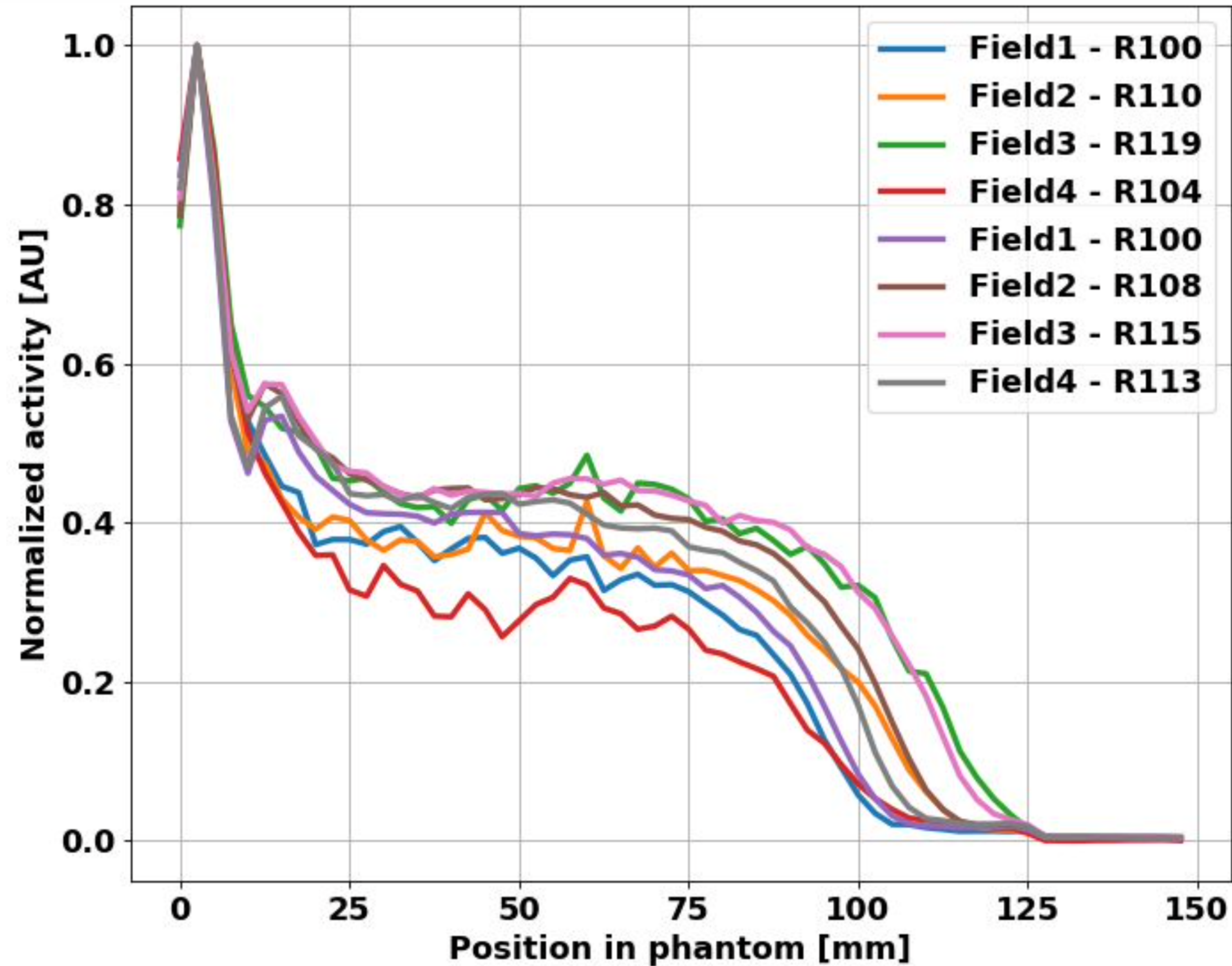
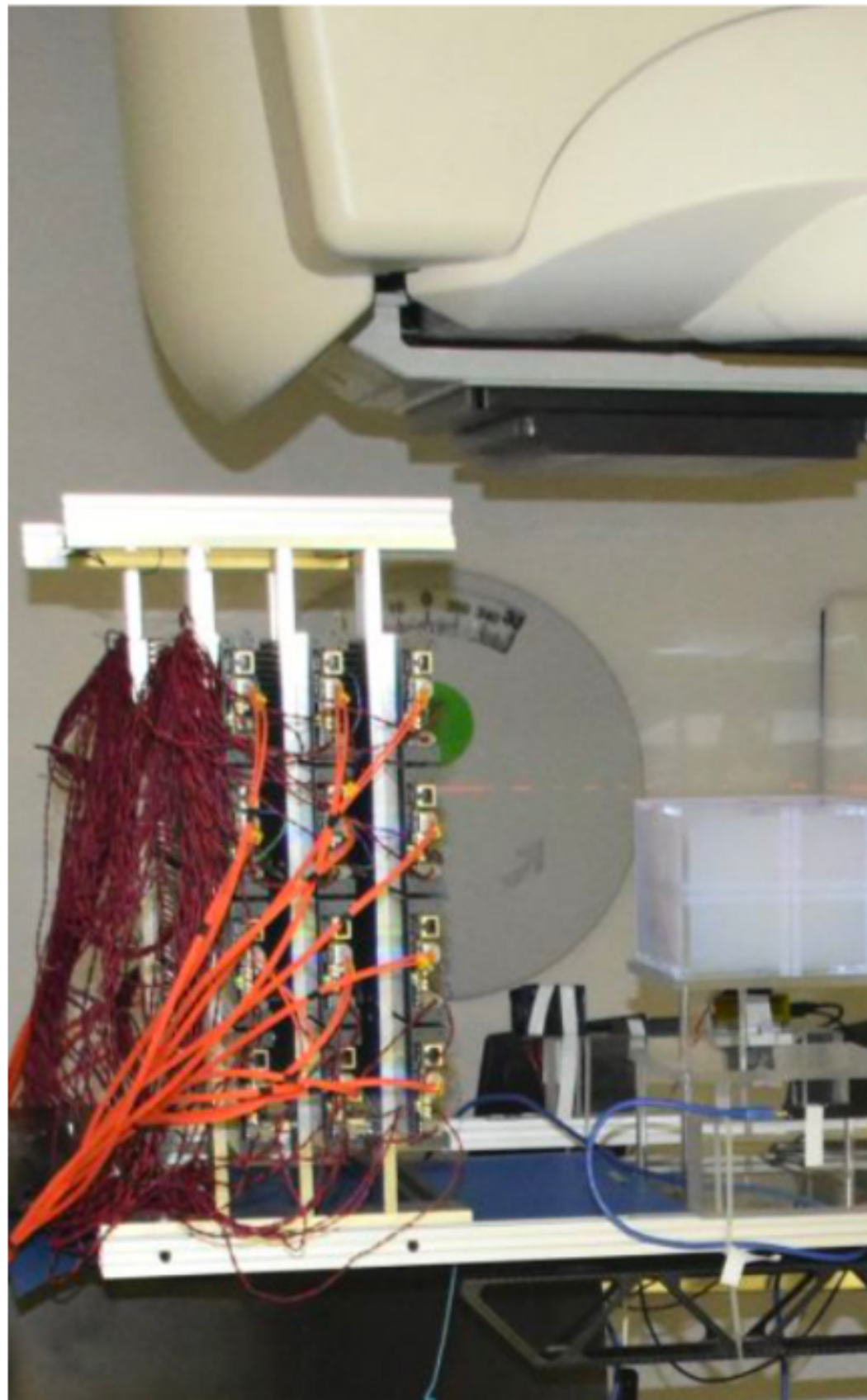
## Experimental validation with proton beams at CCB





# J-PET for range monitoring in PT

## Experimental validation with proton beams at CCB

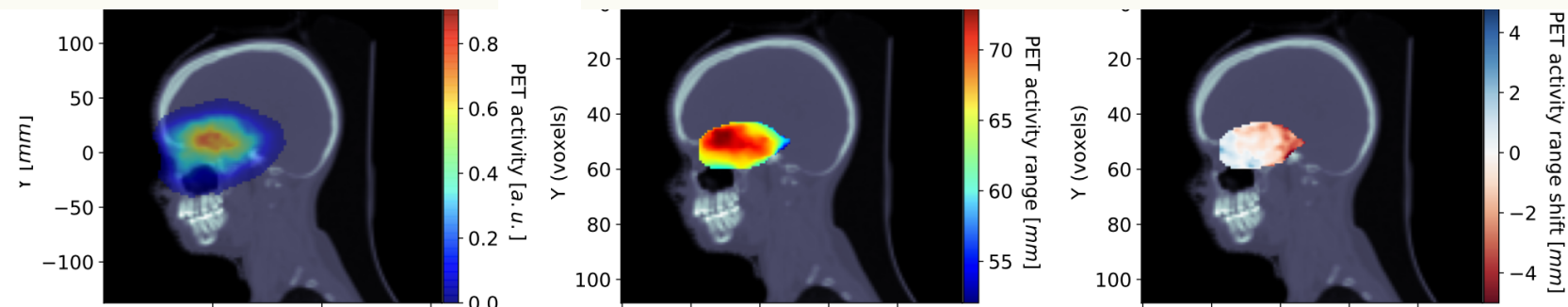


# Range shift detection in patients

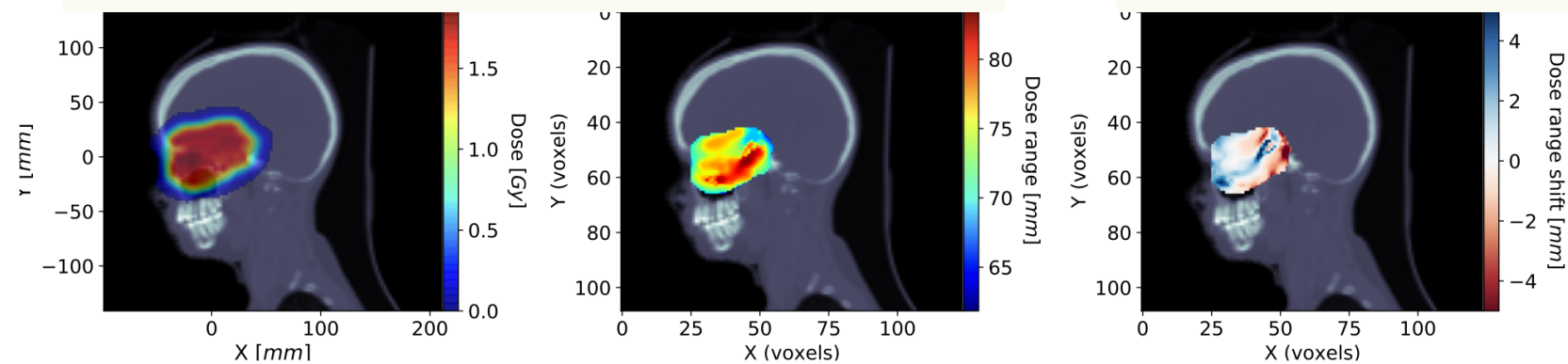
## Patient simulation studies with J-PET

94 patients x 27 scenarios  
(24 shifts + 2 CT cal. + reference)

Activity BEV → Range → Range diff.



Dose BEV → Range → Range diff.

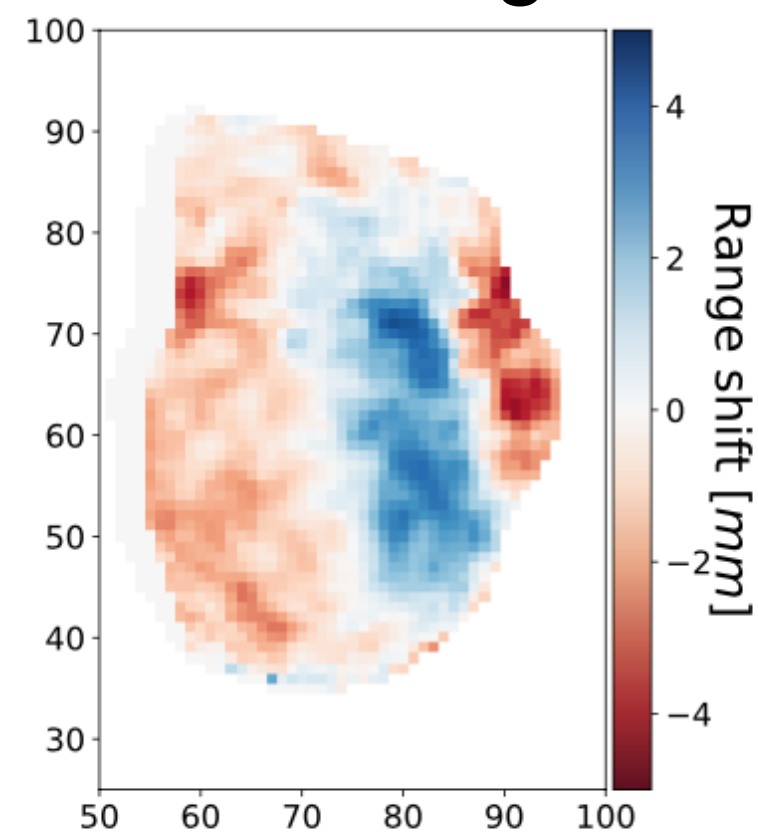


(d)

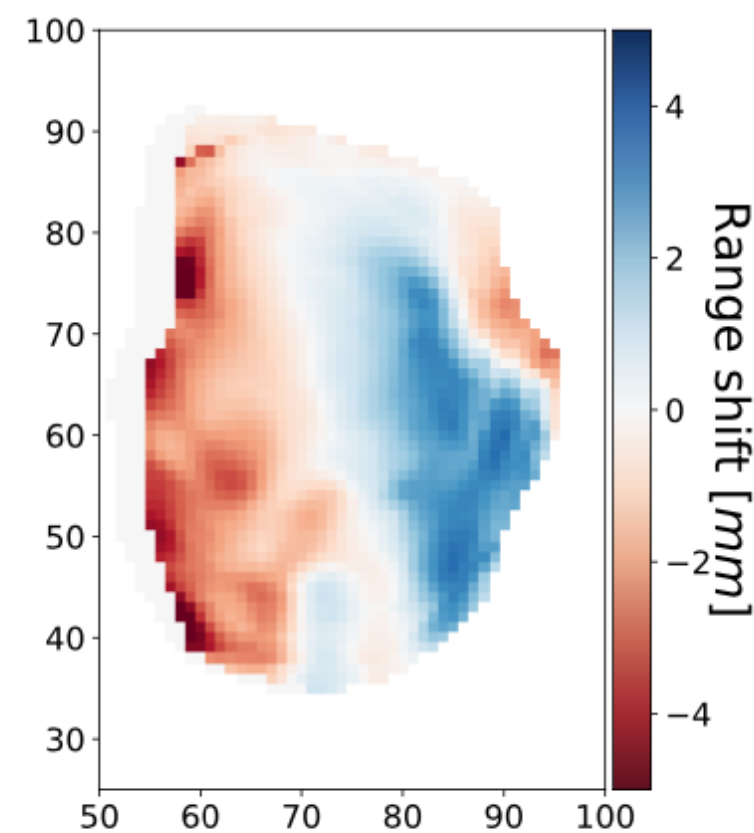
(e)

(f)

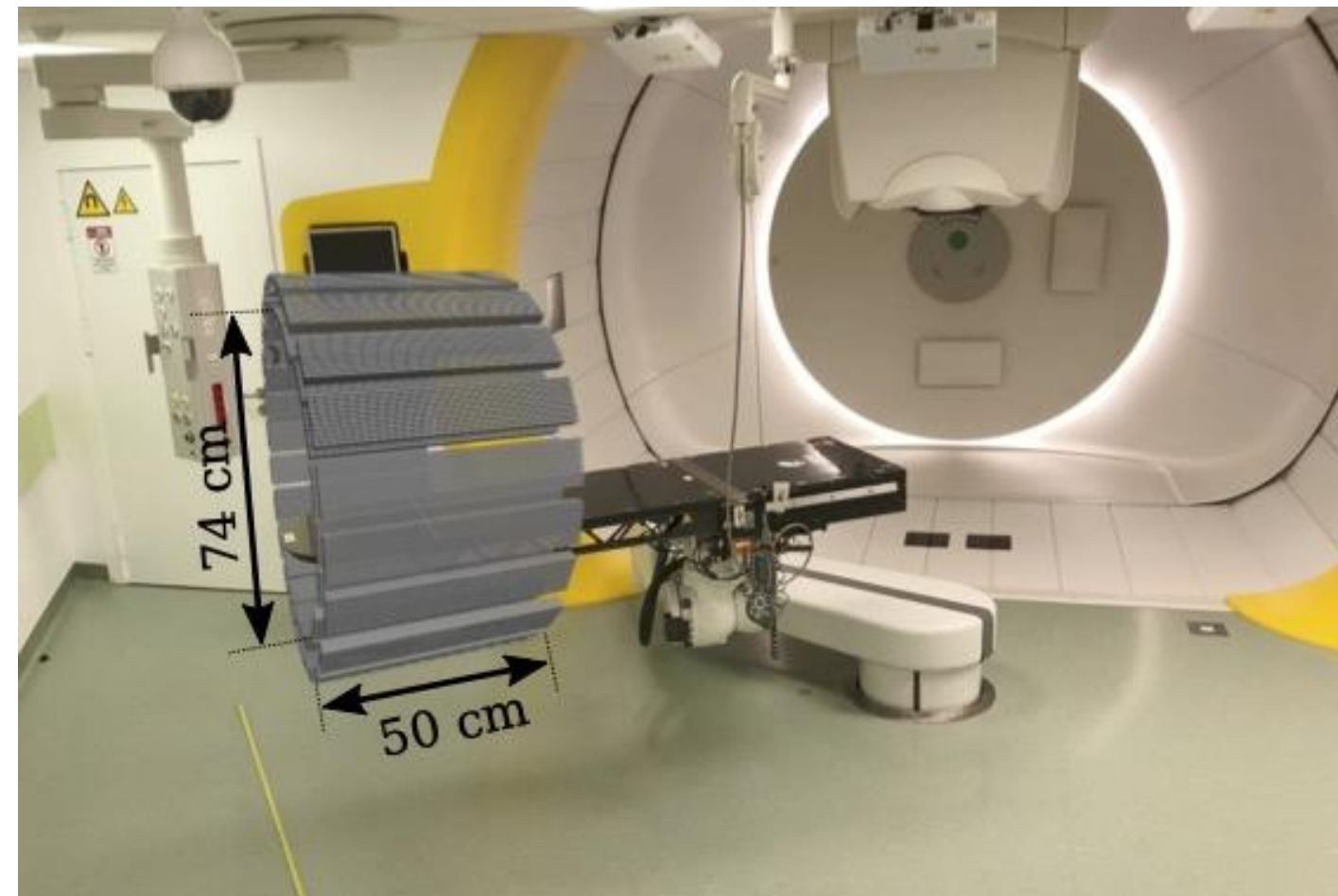
PET image



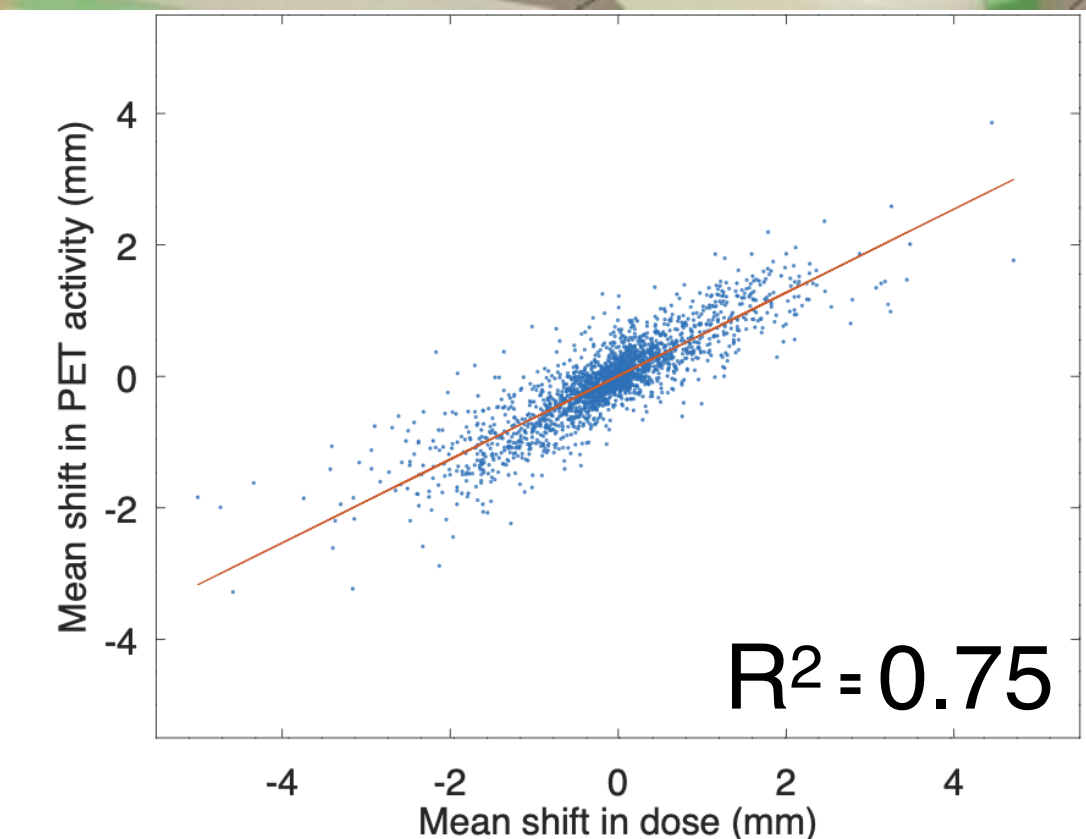
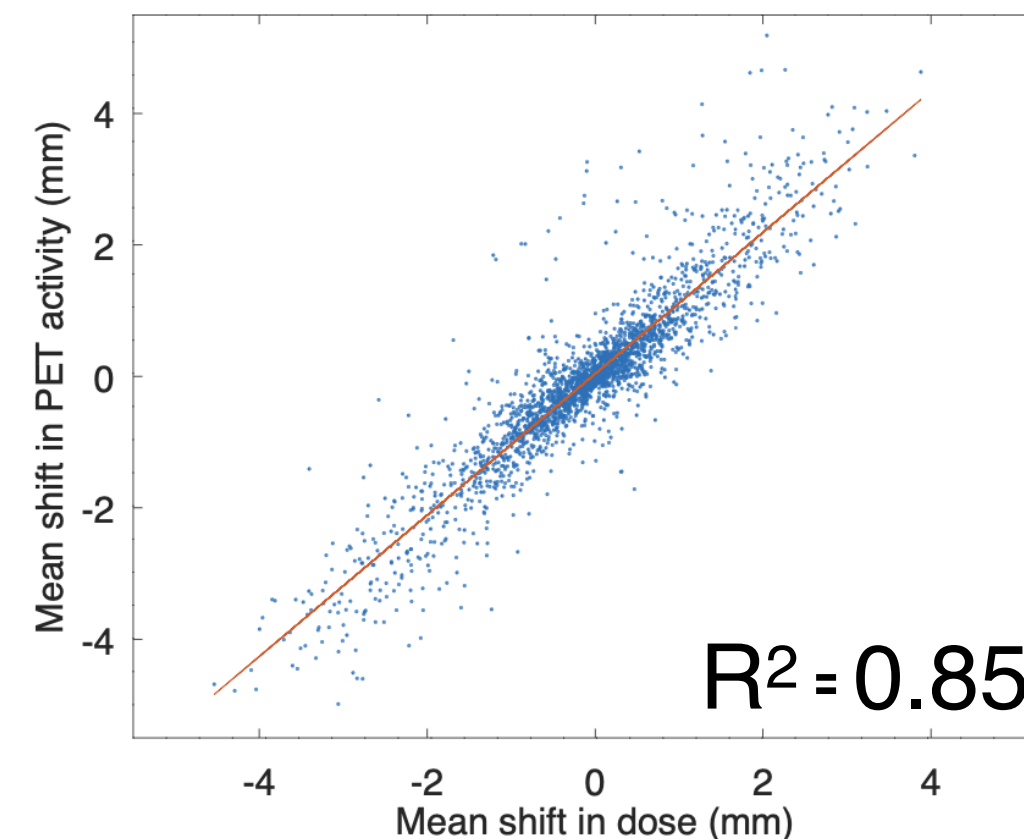
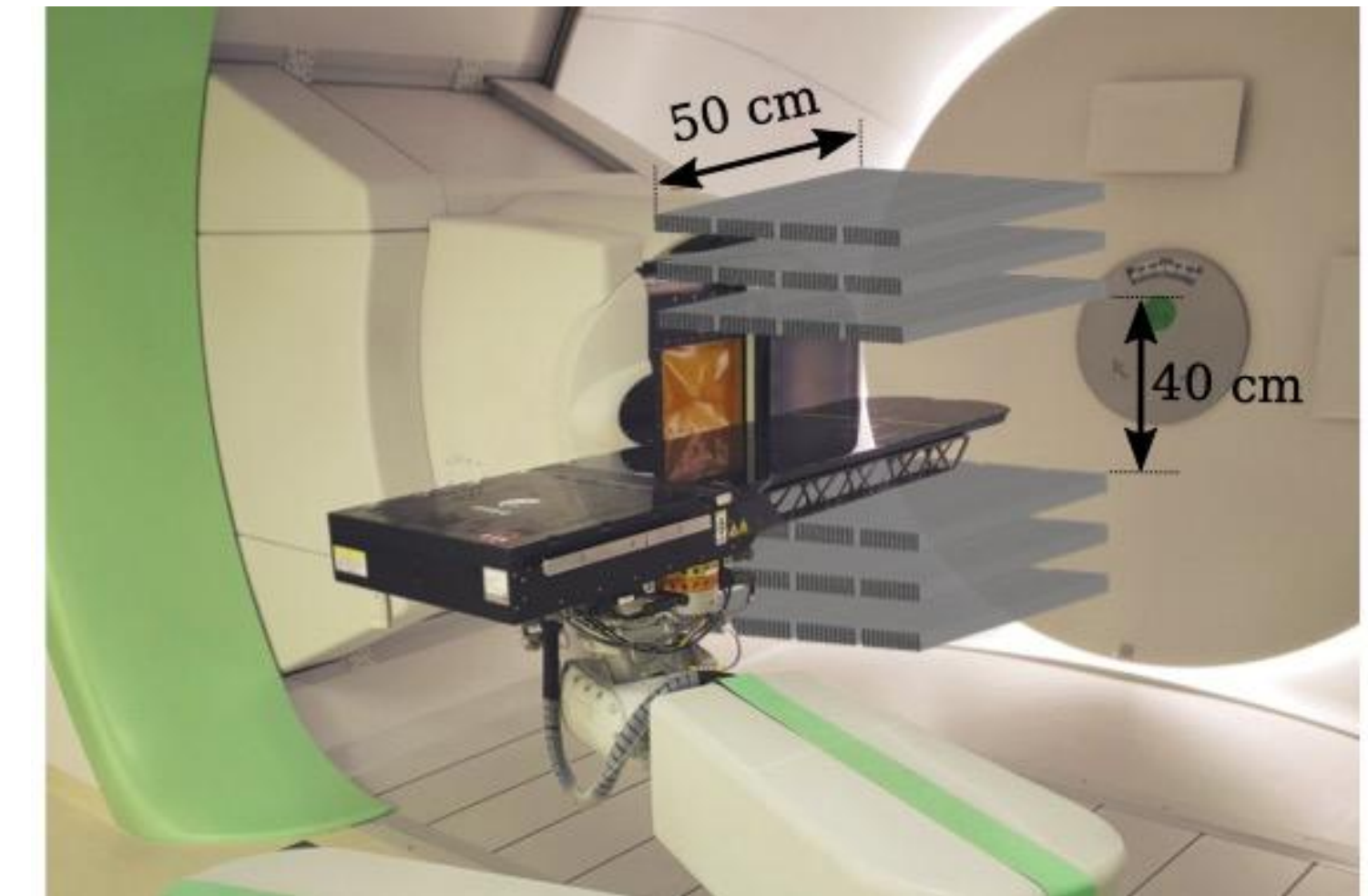
Dose image



In-room PET imaging protocol



In-beam PET imaging protocol



K. McNamara et al. 2022 <https://doi.org/10.1088/1361-6560/aca515>

D. Borys et al. 2022 <https://doi.org/10.1088/1361-6560/ac944c>

K. Brzezinski, et al., PMB (2023) provisionally accepted

# Summary

- **Proton therapy facility offers to medical physics research:**
  - Access to proton beams in Gantry rooms for experiments
  - Anonymised patient data for TP studies and protocols development
  - Radiobiology labs equipped to perform in vitro experiments
- **In CCB Krakow proton center**
  - MC based patient QA - Dose evaluation
  - MC to Support treatment planning - LET evaluation
  - Detector development - J-PET and TimePix
  - Ionisation detail based treatment planning

# Thank you

M. Bałamut, D. Borys, K. Brzeziński, J. Gajewski, M. Garbacz, W. Komenda,  
R. Kopeć, M. Kozani, D. Krzempek, K. Krzempek, G. Mierzwińska, J. Mischczyk,  
N. Mojżeszek, M. Rydygier, P. Stasica, A. Zając-Grabiec



THE HENRYK NIEWODNICZAŃSKI  
INSTITUTE OF NUCLEAR PHYSICS  
POLISH ACADEMY OF SCIENCES

M. Pawlik-Niedźwiecka,  
Sz. Niedźwiecki, G. Korcyl,  
J. Baran, P. Moskal



JAGIELLONIAN UNIVERSITY  
IN KRAKOW

V. Patera, A. Schiavi



SAPIENZA  
UNIVERSITÀ DI ROMA

C. Chang, L. Lin



EMORY | WINSHIP  
CANCER  
INSTITUTE

E. Scifoni, F. Tommasino



Istituto Nazionale di Fisica Nucleare

K. Schubert, N. Hanh



E. Pluta, T. Skóra,  
D. Kabat



MARIA SKŁODOWSKA-CURIE  
INSTITUTE - ONCOLOGY CENTER

I. Rinaldi,  
G. Vilches-Freixas



C. Granja, C.  
Oancea,  
L. Marek



N. Krah



D. Borys



C. Winterhalter,  
K. McNamara,  
A. Lomax  
D. C. Weber  
S. Zürrer



M. Durante



# The hypothesis: Similar clustering leads to similar biological effect

- Local **ionization cluster size** on the nanometer scale is the starting condition for all subsequent processes that lead to the observed DNA damage
- The **complexity of the damage determines the repair pathways** chosen by cells and, in turn, will dictate the **final biological outcomes** that are important for RT
- **Complex DNA damage** caused by ionization clusters in short DNA segments is responsible for **cell death/dysfunction/DNA mutation**

