#### The MC simulation of WIDMApp: an innovative approach for individual dose monitoring in Molecular Radiotherapy

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#### Molecular Radio-Therapy

- MRT dose distribution depends on:
  - Pharmaceutical biokinetics
  - Competition with supportive drugs

Uptake and clearance vary among individuals

 The state of the art monitoring is done with a few SPECTs
 Low accuracy (uncertainty > 30%)

Dose limited by organs at risk dose knowledge

### The WIDMApp approach

Wearable Individual Dose Monitoring Apparatus

WIDMApp aims to provide a <u>continuos</u> real-time measurement of radiopharmaceutical accumulation and transit in MRT treatments.



Detector system

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MC simulation

Detector system

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Detector system

MC simulation

Unfolding algorithm<sub>5</sub>

#### Monte Carlo simulation needs for WIDMApp

- Indispensable to estimate the probability that a photon produced in each organ produce a signal in all the detectors
- It will be tailored on each patient
  - Importing the CT
  - ROIs to define the relevant organs
  - ROIs to define detectors (placeholders during the CT)

#### Feasibility study with NEMA phantom

- NEMA a human abdomen phantom (filled with water)
- 3 spheres filled with 3 different isotopes
- to mimic 3 organs with different washout <sup>18</sup>F ( $\beta^+$ ,  $t_{1/2}$ =1.83 h, A= 54 MBq) <sup>99m</sup>Tc (140 keV  $\gamma$ ,  $t_{1/2}$ =6.01 h, A= 53 MBq) <sup>64</sup>Cu ( $\beta^+$ ,  $t_{1/2}$ =12.7 h, A= 28 MBq)
- 3 detectors placed nearby the 3 spheres





#### MC simulation

- Developed thinking to the WIDMApp pipeline
- Imports the CT of the NEMA phantom for the geometry
- Sample primaries generation points from "organ" ROIs designed in the DICOM files
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- Hit or miss approach
- Winding number algorithm to check if the point is inside the ROI
- Place the detectors in a G4ParallelWorld

#### Sensitive detectors

- The detectors are placed in a G4ParallelWorld
- Can be superimposed to the CT geometry
- It's possible to use all the possible voxel tracking
- It will be possible to define detectors from the ROIs
  - We are working on using CGAL to find a tasselated solid from each DICOM ROI
  - and importing that as a G4TasselatedSolid

#### Checking the geometry

- A dedicated run to check the geometry has been developed
- Geantinos are shot from one of the CT planes
  - One per voxel
  - In the direction orthogonal to the plane
- At each step, density and material index are scored

#### density3D yx projection



#### MC output

The MC output is the probability of each organ to produce a signal in all the detectors



Such matrix is needed by the unfolding algorithm

$$\chi^{2}\left(A'_{0j}, t'_{1/2j}^{eff}\right) = \sum_{i=1}^{N_{det}} \sum_{k} \left( \underbrace{S_{i}(t_{k}) - S_{i}(t_{k}; A'_{0j}; t'_{1/2j})}_{\sigma_{i}(t_{k})} \right)^{2}$$

$$S_{i}(t_{k}; A'_{0j}, t'_{1/2j}) = \sum_{j=1}^{N_{org}} p_{ij} A_{0j} e^{-t_{k}/(t_{1/2j})/\ln(2)}$$

#### Results

- Estimation of the total absorbed dose with an error of 3-7% on  $t_{1/2}$
- Larger error on <sup>99m</sup>Tc A0
  - WIDMApp is not designed to estimate A<sub>0</sub>s
  - <sup>99m</sup>Tc emits a **y** with lower energy

	t <sub>1/2</sub> [h]	err	A <sub>0</sub> [MBq]	err
<sup>18</sup> F	1.97 ± 0.06	7%	54.8 ± 3.3	1%
<sup>99m</sup> Tc	6.20 ± 0.35	3%	43.7 ± 3.8	-18%
<sup>64</sup> Cu	13.34 ± 0.08	5%	28.3 ± 1.9	<1%



#### Container

- The WIDMApp MC simulation has several external dependencies
  - Geant4
  - DCMTK (for each patient a simulation is needed -> reading DICOM is mandatory)
  - CGAL (in future)
- A Docker container has been developed with all the dependencies installed

#### Conclusions

- WIDMApp is a project aimed at monitoring MRT
- MC simulation is a key point to infer the organs dose from the detector measures
- It will have to be tailored for each patient
- We developed a Geant4 application designed to use ROIs to identify organs and detectors
- A paper will be submitted soon

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thank you for your attention!

#### BACKUP SLIDES

#### First feasibility study

- MIRD Phantom
  - 6 activated organs



- Source:
  - 1131 uniform emission;
  - 2.5 GBq treatment (30% in thyroid, 60% on other organs: Liver, Kidneys, Spleen, Bladder)
- Detectors:
  - Radius 1.3 cm, thickness 0.3 cm;
  - Efficency = 100%;
  - One per organ (two on thiroid);

#### First feasibility study

- The algorithm manages to converge to the true values
- We tested also its stability with respect to priors and background



• S. Morganti et al., A wearable radiation measurement system for collection of patient-specific time-activity data in radiopharmaceutical therapy: system design and Monte Carlo simulation results. Med Phys. 2021;1-10. https://doi.org/10.1002/mp.15311

The algorithm converges up to fluctuations on the measured points of 15%  $\frac{1}{7}$ 

#### Primaries generation

- The generation of primaries is done sampling (with a hitor-miss approach) points within the "organs" ROIs
- Firstly selecting a slice of the ROI and so its Z
- Then sampling a points within
   (Xmin : Xmax, Ymin : Ymax, Z-dZ : Z+dZ)
  - where X/Ymin and X/Ymax are the ROI extremes
  - Z is the ROI slice one and dZ its thickness

### Checking if the point is inside the ROI

- To accept the sampled point we have to check if it lies inside the selected ROI slice
- The fastest and precisest method is the *winding number* algorithm:
  - Find all the edges of the polygon that cut through the line passing through the query point and is parallel to one axis.
  - For these edges, check if the query point is on the left or right side of the edge when looking at all the edges in anticlockwise direction
  - Increase the value of winding number by one if query point is on the left side and decrease by one if query point is on the right side
  - If the final winding number is non zero, the point lies inside the polygon

### The WIDMApp detector

# Requirement for the **wearable y** detecting sensor:

- thin, lightweight, comfortable, mechanical robust, simple, compatible device;
- linear response over a wide
   counting rate range (from few CPS to ~ 10kCPS);
- Iow power consuming and battery operated



### The first prototype

First Sensor Prototype: Para-tephernyl 15x15 mm<sup>2</sup>, h= 3mm;

2x2 6mm C-Series SiPM array.



- Shape of the sensor and read-out are not yet optimized
  - A simulation will be perfomed for this
- It has a known non linearity for high rate values
  - The saturation in data has been corrected
- . The efficiency have been calculated with a dedicated test

#### Detector efficency

 $CPS(t) = A(t) \times \epsilon_{Geom} \times \epsilon_{Int} \times \epsilon_{Det} = A(t) \times \epsilon_{MC} \times \epsilon_{Det},$ 

- Measurements with only one active sphere per time inside the phantom have been acquired
- Single isotope measurements have been used to evaluate the efficiency of each detector, according to:

$$\epsilon_{Det}^{ij} = \frac{CPS(0)^j}{A(0)^i \times \epsilon_{MC}^{ij}},$$

• T=0, i.e. the test start time.

#### Clinical case of interest

- Neuroendocrins tumor with Lu177
- PROS:



- Great interest on this isotope only lately used in the clinical practice;
- Many questions on the dose absorbed by kidneys;
- . CONS
  - The literature about is poor
  - These kind of tumors are mostly in the abdomen region

# **Targeted Radionuclide Therapy**

- Targeted Radionuclide Therapy How does it work?
  - A **radiopharmaceutical** is injected in the patient
  - Due to its **specificity** for the tumor, the drugs binds to the tumor
  - **Radiation** emitted by the radionuclide damages the tumor



### **Dose Monitoring in TRT**

- A systematic measurement of the internal activity distribution in all the patient undergoing TRT would enable improvements in treatment control and an individualization of the activity to be administered;
- This personalization would increase the local tumor control while sparing healthy tissue and lowering toxicity.

When the hospital resources are available, the state-of-the-art method utilizes SPECT scan imaging:



### **Dose Monitoring in TRT**

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- This personalization would increase the **local tumor control** while **sparing healthy tissue** and **lowering toxicity**.

Need for a system for dosimetry measurements capable of a continuous mapping of the organs TAC



# **Feasability Study - Simulation**

MC Simulation



contribution from each organs emission for all the placed detectors



#### • MIRD Phantom:

 Simplified, reference in radio protection Organs studies

#### • Detectors:

- Radius 1.3 cm, thickness 0.3 cm;
- Efficency = 100%;
- Placed in correspondance with the organs of interest
- Source:
  - Organ source: I<sup>131</sup> uniform emission



# **Feasability Study - Simulation**

MC Simulation



contribution from each organs emission for all the placed detectors



we obtain the values of the weight  $\varepsilon_{ij}$  for each detector-organ combination

## **Feasability Study - Dataset creation**







Input for the definition of a realistic data set

# **Feasability Study - Dataset creation**



Assumptions on <sup>131</sup> I Biokinetic  $(A_0 and \tau_{organ})$ 



Input for the definition of a realistic data set

- A **2.5 GBq** administration is assumed;
- We roughly assigned 30% of the activity to the **thyroid**, given its high specificity for Iodine, and 70% equally divided;
- We further divided this second part among the 5 other active organs assuming for them the same specific activity of radiopharmaceutical.

## **Feasability Study - Dataset creation**



Assumptions on <sup>131</sup>Ι Biokinetic (A<sub>0</sub> and **τ**organ)



Input for the definition of a realistic data set

• The number of total counts (i.e. from all the organs) is obtained as a function of the time at the given detector:

$$S_{MC}^{i} = \sum_{i=1}^{6} A_{0_{j}}^{theo} e^{-t/\tau_{j}^{theo}} \epsilon_{ji}$$

- As described in the equation, considering the contribution of each organs (j) we obtain a Time Counts Curves (TCC) for each detector (i);
- In order to simulate the detector response, we sampled the detector responses assuming a Poissonian distribution;

### **Feasability Study - Deconvolution Algorithm**



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The role of the WIDMApp algorithm is to infer from these 7 TCCs the 6 TACs, one per organ, that originated them.

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$$S_{MC}^{i} = \sum_{i=1}^{6} A_{0_{j}}^{theo} e^{-t/\tau_{j}^{theo}} f_{ji}$$

- The algorithm has thus to identify the combination of 6 pairs of parameters A<sub>0j</sub> and τ<sub>j</sub> that best reproduces all the 7 TCCs data points at the same time;
- This is achieved by defining a merit function  $\chi^2(A_{0j}, \tau_j)$  that considers all the points from all the 7 TCCs:

$$\chi^{2}(A_{0j},\tau_{j}) = \sum_{i}^{N_{det}} \sum_{k} \left( \frac{S_{i}^{sim}(t_{k}) - S_{i}(t_{k};A_{0j},\tau_{j})}{\sigma_{i}(t_{k})} \right)^{2}$$

- Given the strong correlation between the unknown parameters, different minimization algorithms were compared in terms of robustness, varying:
  - Initial parameters of the fit;
  - Uncertainties applied to the data;