

# *In vitro* dosimetry for assessment of Targeted- $\alpha$ -Therapy (T $\alpha$ T)



## XXIInd Colloque GANIL – Applications

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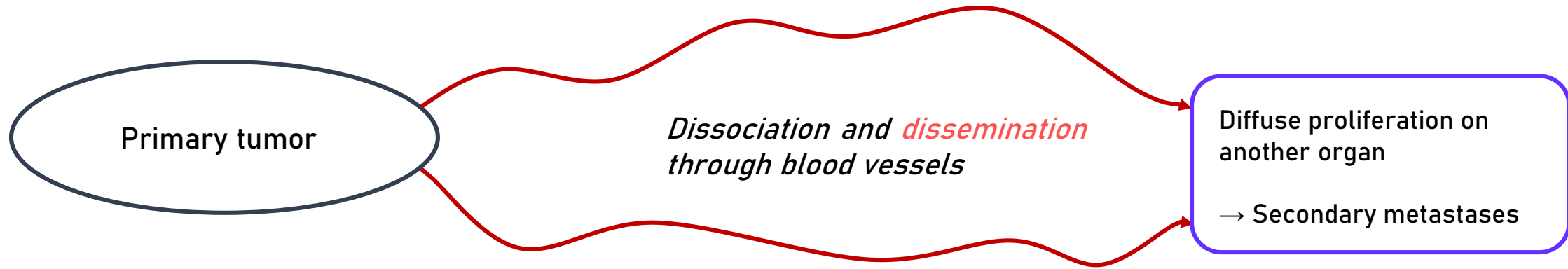
A. Doudard<sup>1</sup>, A. Corroyer-Dulmont<sup>2,3</sup>, C. Jaudet<sup>2</sup>, M. Bernaudin<sup>3</sup>, S. Valable<sup>3</sup>, A.M. Frelin-Labelme<sup>1</sup>

<sup>1</sup>Grand Accélérateur National d'Ions Lourds (GANIL), CEA/DRF CNRS/IN2P3, 14076 Caen, France

<sup>2</sup>Medical Physics Department, CLCC François Baclesse, 14000 Caen, France

<sup>3</sup>Normandie Univ, UNICAEN, CEA, CNRS, ISTCT/CERVOxy Group, Caen, France

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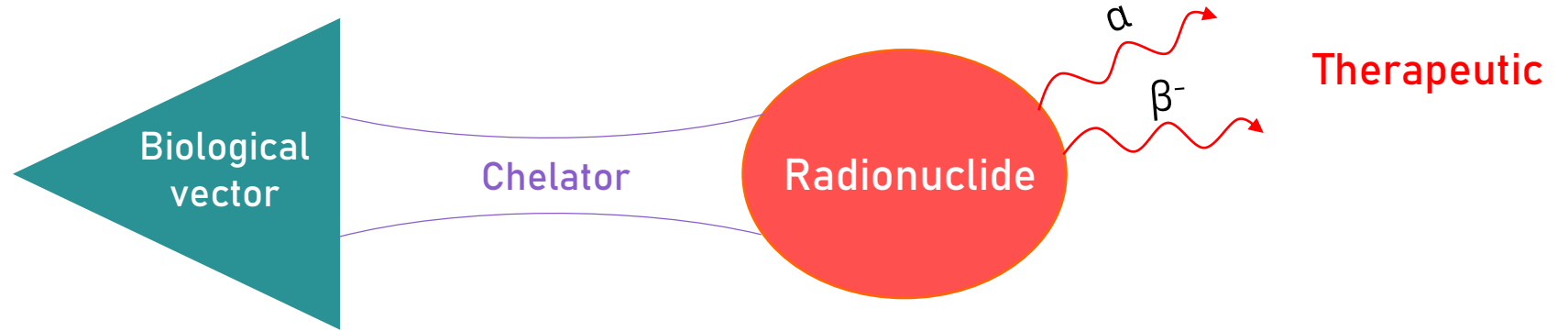


- Metastases are detectable once formed, treatment via radiosurgery, radiotherapy and/or chemotherapy.
  - ↳ Multiple locations + potential radio-induced brain damage → poor prognostic (brain metastases: 6 months)
- Detection and treatment of early stages of brain metastasis formation: Targeted Radionuclide Therapy (TRT)

# VCAM-1-based TaT

Overview · Detection system  
Deconvolution methods  
Application to  $^{223}\text{Ra}$  · Outlooks

· Radiopharmaceutical structure:



· Over-expression of VCAM-1 observed near breast-cancer-induced brain metastases

→ Anti-VCAM-1 antibody

· Spatially restricted target + sensible environment (few dozens  $\mu\text{m}$  in brain)

· Compromise between clinical usability and short physical period (radioprotection)

$^{212}\text{Pb}-\alpha\text{VCAM-1}$

*A. Corroyer-Dulmont et al., Neuro-Oncology (2020)*

*N. Falzone et al., Theranostics (2018)*

$^{212}\text{Pb}$  (10.1 h)

↳  $^{212}\text{Bi}$  (60.5 min): 6.1 & 8.8 MeV  $\alpha$   
= 50 & 91  $\mu\text{m}$  range in water

➤ Goal of *in vitro* tests: measure biological effectiveness as a function of the delivered dose to the cells.

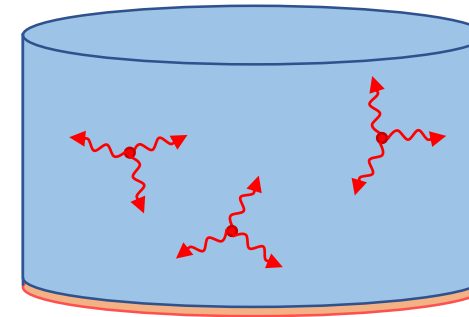
· MIRD formalism:

$$\text{Dose} \propto A_0 \times \phi(\text{target} \leftarrow \text{source})$$

Injected activity

Fraction of energy emitted by the source absorbed by the target

· *In vitro* configuration:



~2 mm : Culture medium

◇ ~20 μm : Cell medium

Simulated dosimetry under homogeneous distribution hypothesis

β ✓

(range > mm)

α ✗

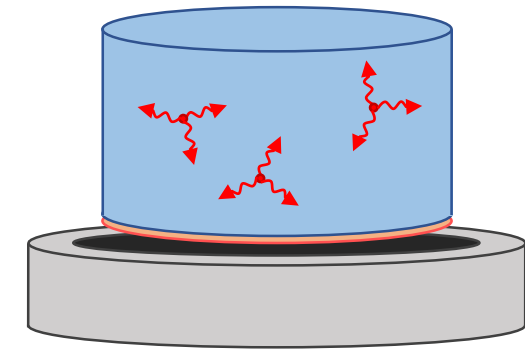
$^{212}\text{Bi}$  : 6.1 & 8.8 MeV  
= 50 & 91 μm in water

→ Need to determine the spatial distribution of the radionuclides in the culture medium and its evolution during the time of the irradiation.

A.M. Frelin-Labalme et al., Med. Phys. (2020)

## ➤ Development of a new TaT dosimetry system:

→ Acquisition of energy spectra of the alpha particles emitted through the culture medium and cell layer ;



*α-detector under culture well*

→ Application of a spectral deconvolution method to estimate the spatial and temporal distribution of the radionuclides ;

→ Reconstruction of the dose deposited on the cells through Monte-Carlo simulations.

# Detection of $\alpha$ -particles

• Dimensions compatible with in vitro cultures

• High efficiency (low activities)

Silicon semiconductor detectors



(PIPS, Mirion, U.S.A.)

• Specific conditions of T (noise, stability)

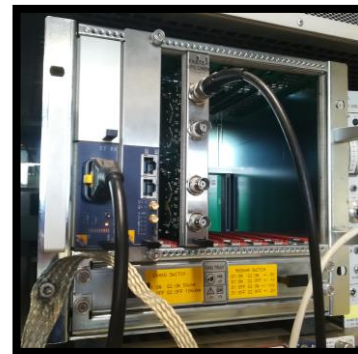
• Good energy resolution for  $\alpha < 10$  MeV and response linearity

$$\sigma_E(E = 5\,486 \text{ MeV}) = 12,3 \text{ keV}$$

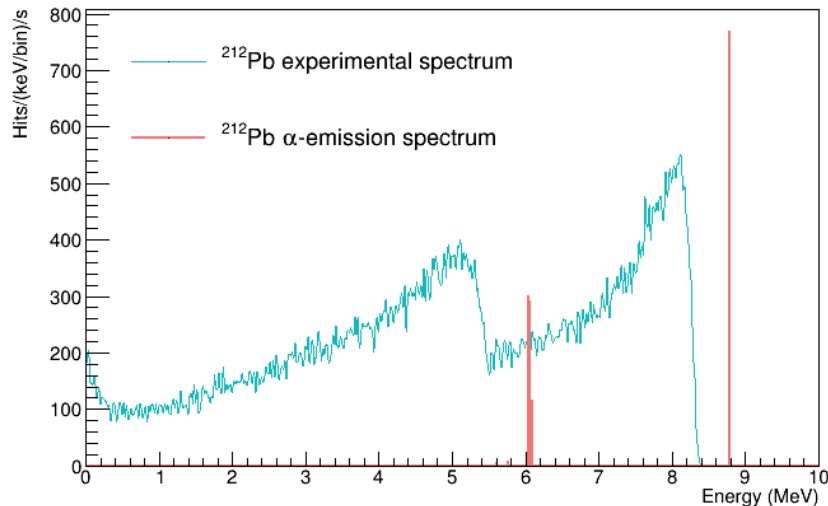
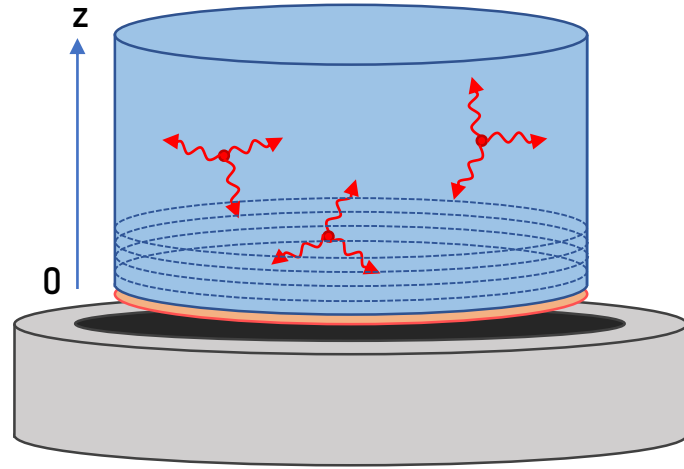
⇒ spatial resolution < 0,5  $\mu\text{m}$

(cell medium: 20  $\mu\text{m}$ , range: 50 to 100  $\mu\text{m}$ )

• Coupled with FASTER (LPC Caen) acquisition system



# Deconvolution following a parametric model



- Time evolution: **t-discretization**
- Decomposition of experimental spectra as a sum of elementary spectra: **z-discretization**

$$\hookrightarrow SP_{exp}(t_k, E) = \sum_i A(t_k, z_i) \cdot SP_{elem}(z_i, E)$$

- Constraints: **Parametric description of activities**

$$\hookrightarrow A(t_k, z_i) = A(z_i, p_1(t_k), p_2(t_k), \dots)$$

- A satisfying model: **exponential distribution**

$$\hookrightarrow A(t_k, z_i) = a(t_k) \cdot e^{-b(t_k) \cdot z_i} + h(t_k)$$

*A.M. Frelin-Labalme et al., Med. Phys. (2020)*

**3 parameters**

•  $SP_{exp}(t, E) = \sum_i A(t, z_i) \cdot SP_{elem}(z_i, E)$   $\xrightarrow[\text{time interval}]{\text{For each}}$   $Y = Xa, \quad a_{sol} = \min_a \|Xa - Y\|^2$

• Additional **physical constraints** on  $a$   
 (positivity, limited sum of activities, ...)

$\longrightarrow$   $\left\{ \begin{array}{l} Ca_{sol} = c \\ d_{min} \leq Da_{sol} \leq d_{max} \\ a_{min} \leq a_{sol} \leq a_{max} \end{array} \right.$  (Quadratic programming)

• Examples

No constraints on  $a$

Least Squares (LS)

Algebraic resolution

Pseudo inverse :  $a_{LS} = (X^T X)^{-1} X^T Y$

$a \geq 0$

Non-Negative Least Squares (NNLS)

Iterative resolution

*Law & Hanson algorithm*



- The objective function can be modified to adapt a particular modelization :

$$A(t_k, z_i) = a(t_k) \cdot e^{-b(t_k) \cdot z_i} + h(t_k)$$

→ Monotonicity of successive derivatives, property of the exponential model

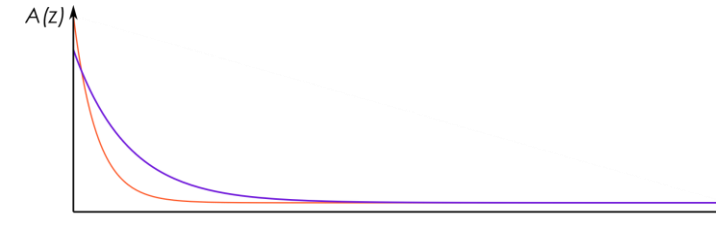
- Other possibilities (can be combined):

→ Removal of constraints near the bottom culture wells



→ Likelihood maximisation criterion instead of least squares

→ Separation of the spatial distributions of different radionuclides



→ Irregular spatial sampling of the elementary spectra (adaptive resolution)

# Application to a clinical radiopharmaceutical

Overview · Detection system  
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➤ **Goal** : assessment of the experimental set-up and of the deconvolution methods



*FASTER module and voltage supplies*



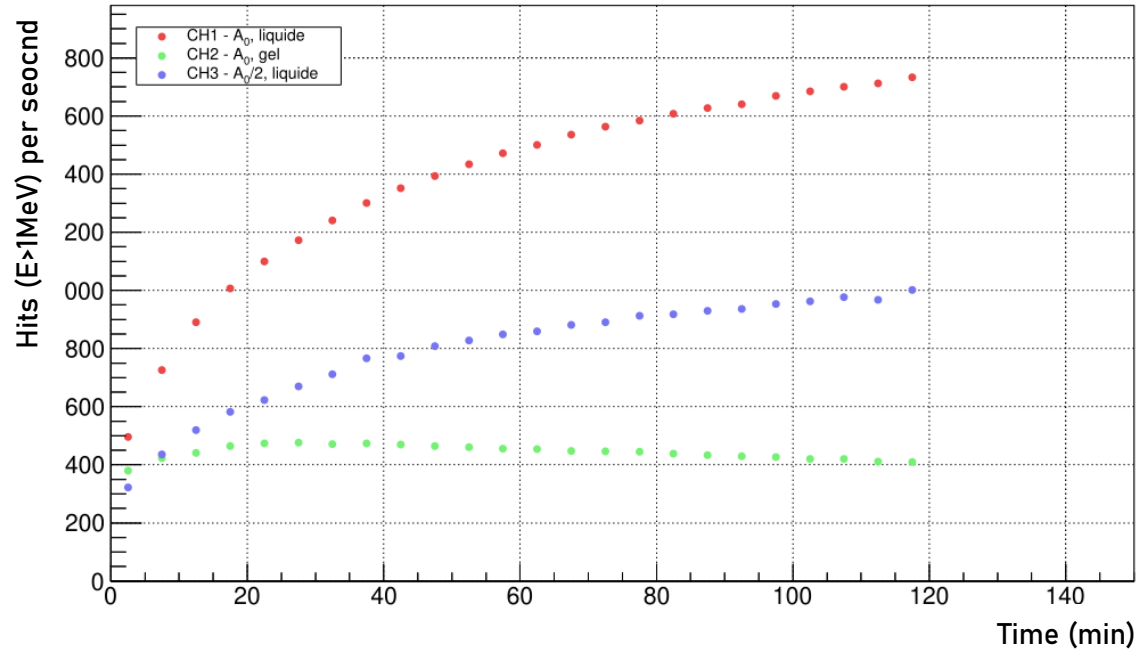
*Light-tight container*



*Diode support and culture wells*

- Measurements with an  $\alpha$ -emitter radiopharmaceutical: Xofigo ( $\text{Cl}_2^{223}\text{Ra}$ ), at the CLCC François Baclesse (Caen, France).
- $4\alpha$ -emission spectrum (5.6 MeV, 6.7 MeV, 7.4 MeV, 6.4 MeV) ( $^{212}\text{Pb}$  :  $1\alpha$ , 2 decay paths, 6.1 MeV & 8.8 MeV).

## Hitrates comparison



• 3 configurations studied:

- Liquid solution of activity  $A_0$
- Liquid solution of activity  $A_0/2$
- Gelified solution of activity  $A_0$  with SuperAbsorbent Polymer (SAP, Curas)

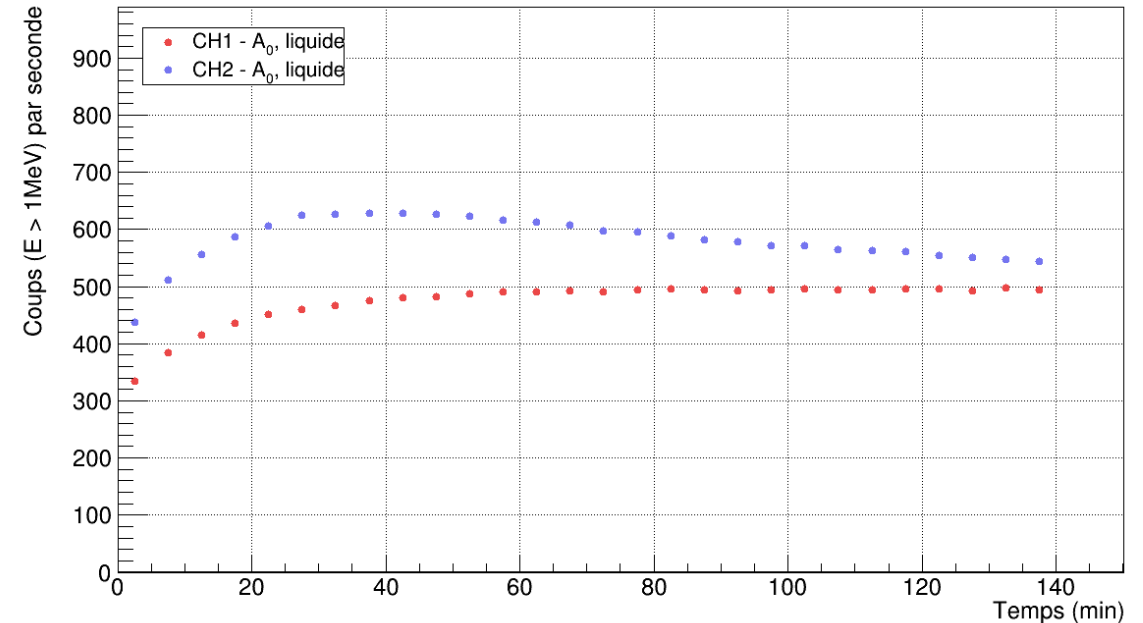
•  $A_0 = 9,3 \text{ kBq}$

• Higher stability of hitrate in time with the gelified solution

→ Showcases the displacement of radionuclides during *in vitro* experiments

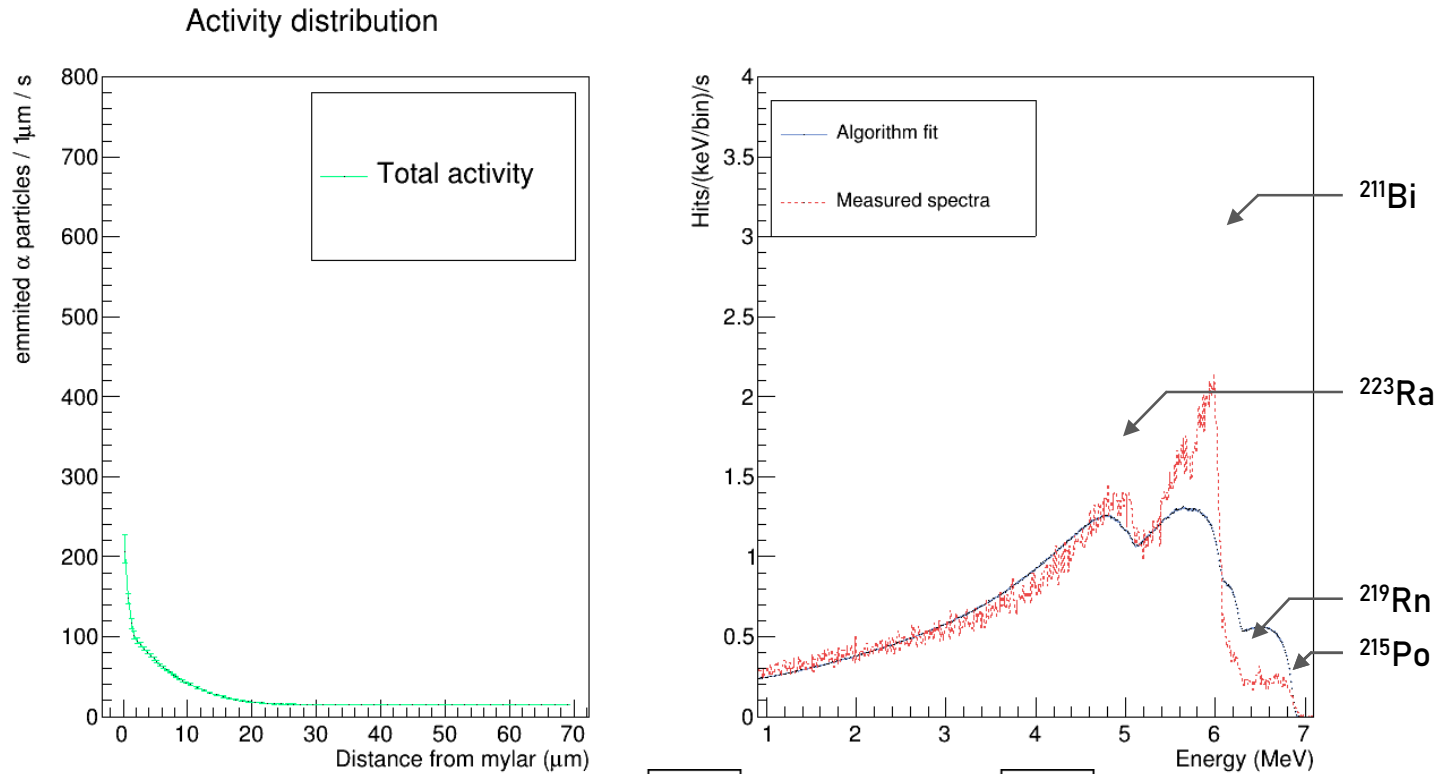
- Both spectra sets correspond to two separate wells filled with 9,3 kBq of the same liquid solution of  $^{223}\text{Ra}$ .
- Different hitrate profiles hint at different distribution kinetics for each well.

Hitrates comparison



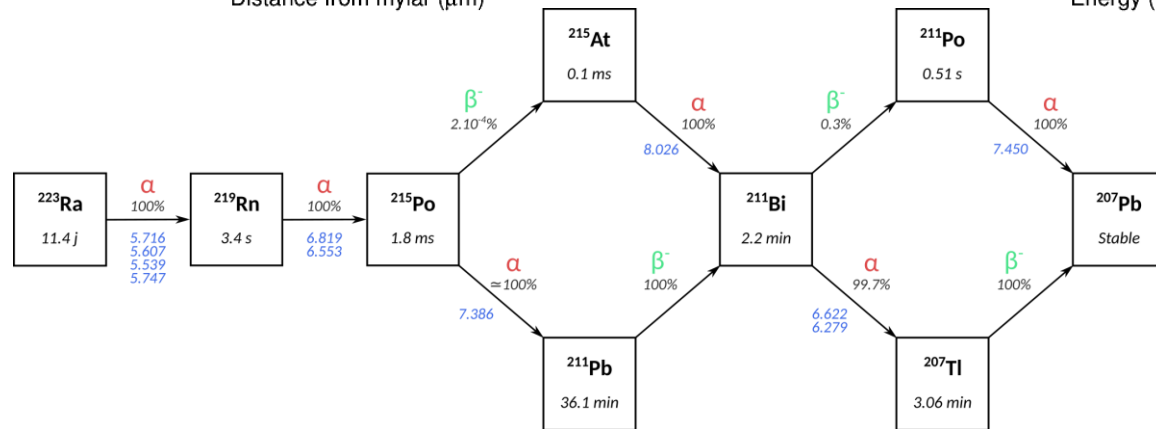
→ Dose uncertainty estimations must also consider potential lack of reproducibility of the tests

# Measurements with $^{223}\text{Ra}$ (3/4) – Deconvolutions

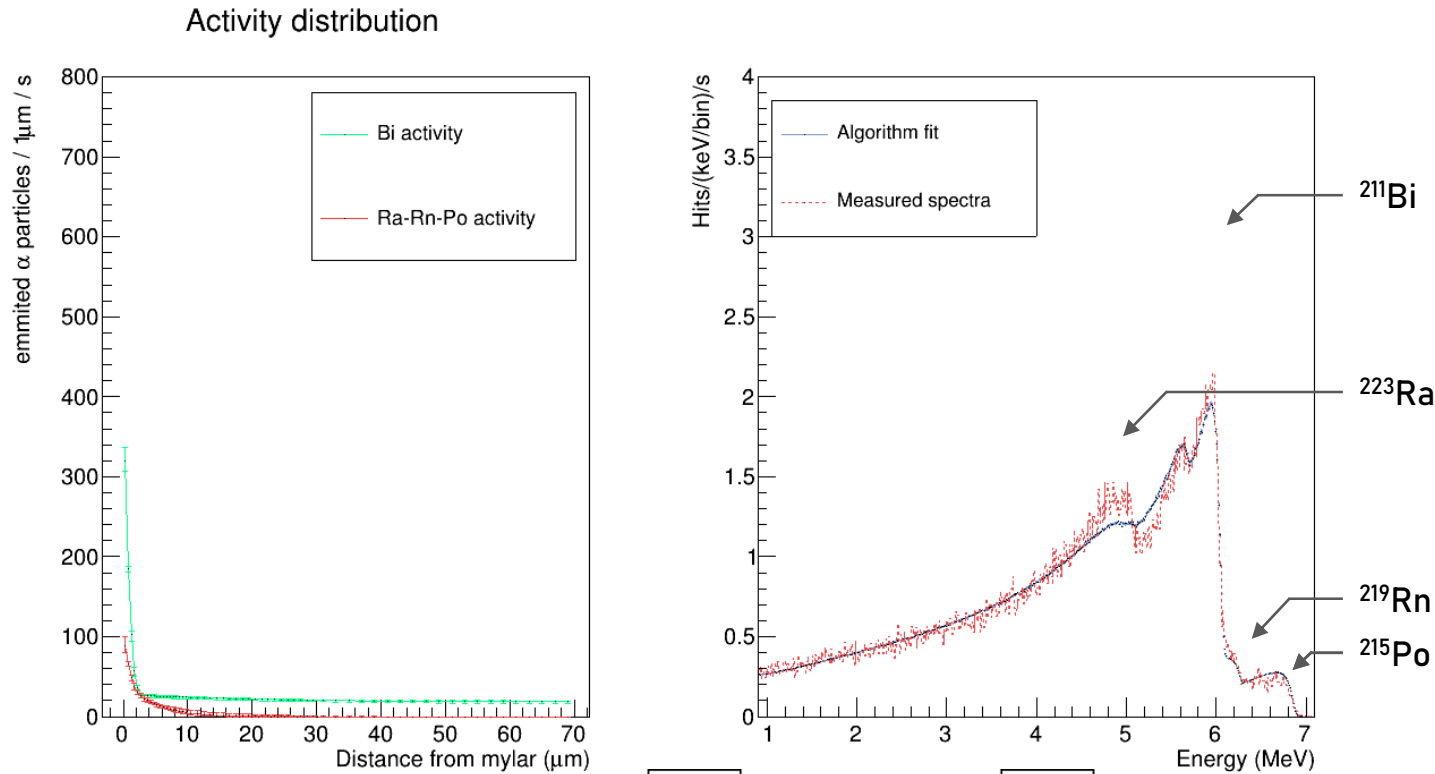


## ➤ Matrix deconvolution methods application

- Set of spectra obtained for a well filled with an activity  $A_0 = 9,3 \text{ kBq}$  of  $^{223}\text{Ra}$
- $^{211}\text{Bi}$  distribution strongly differs to the rest of the decay chain ( $T_{1/2, ^{211}\text{Pb}} = 36 \text{ min}$ )
- A three-way separation of the spatial distributions leads to the best experimental spectra reconstructions (based on information criteria)
- Majority of the information lies in a short energy range : overfitting issues

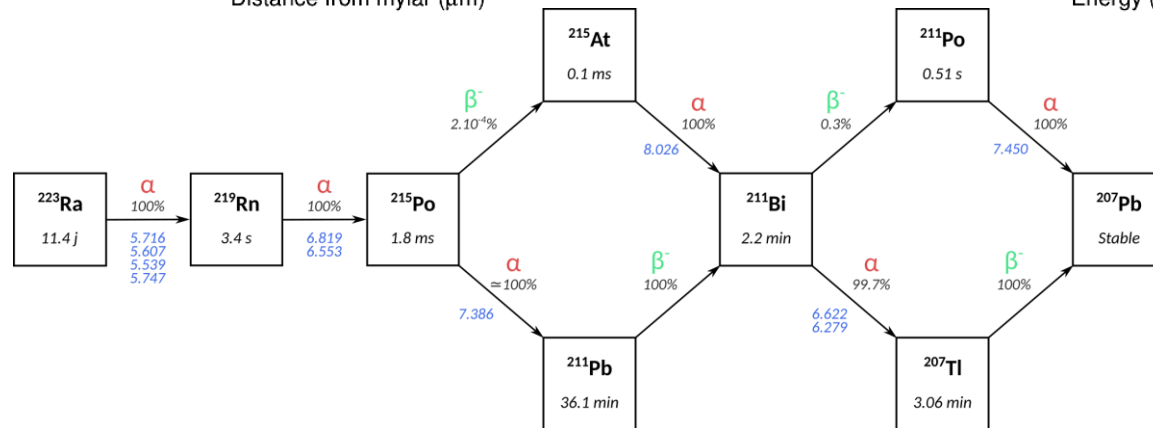


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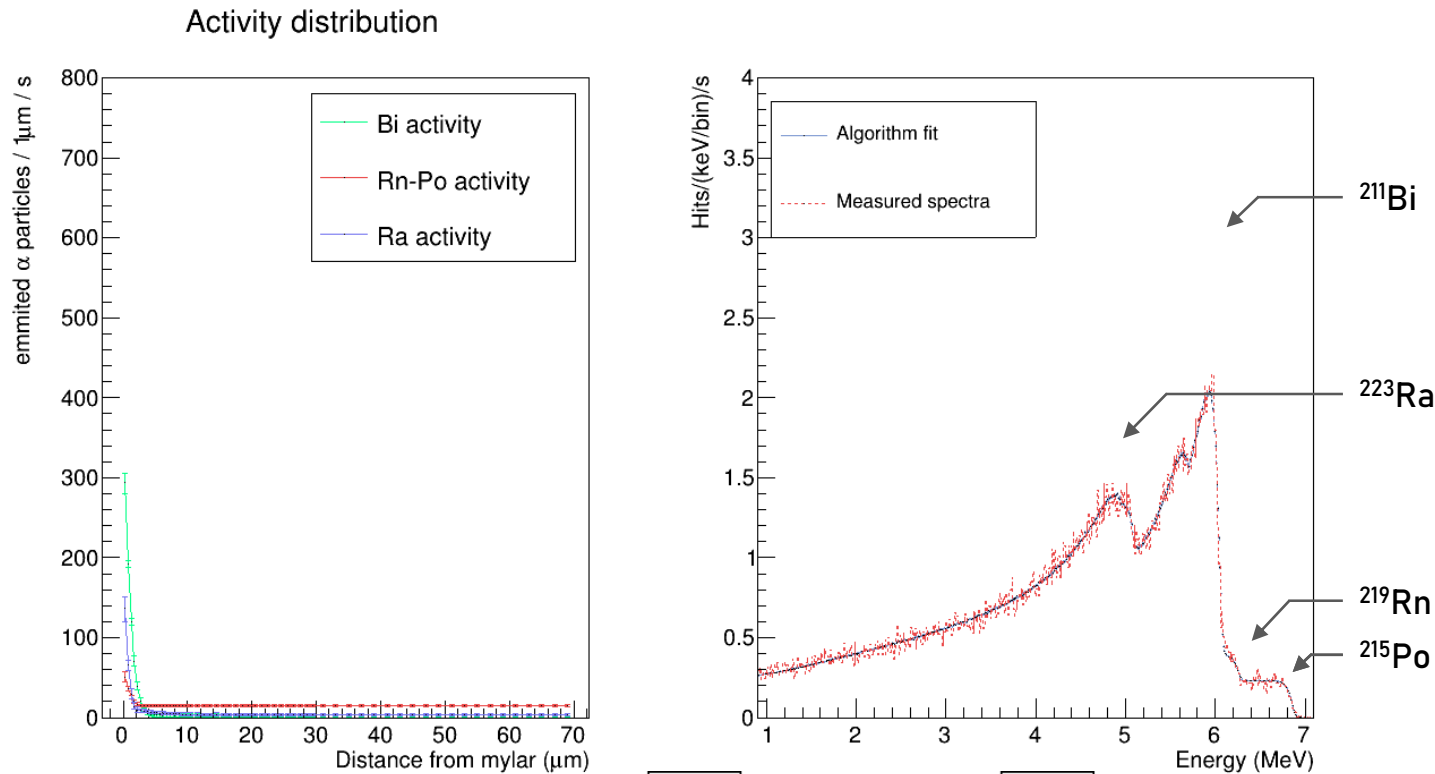


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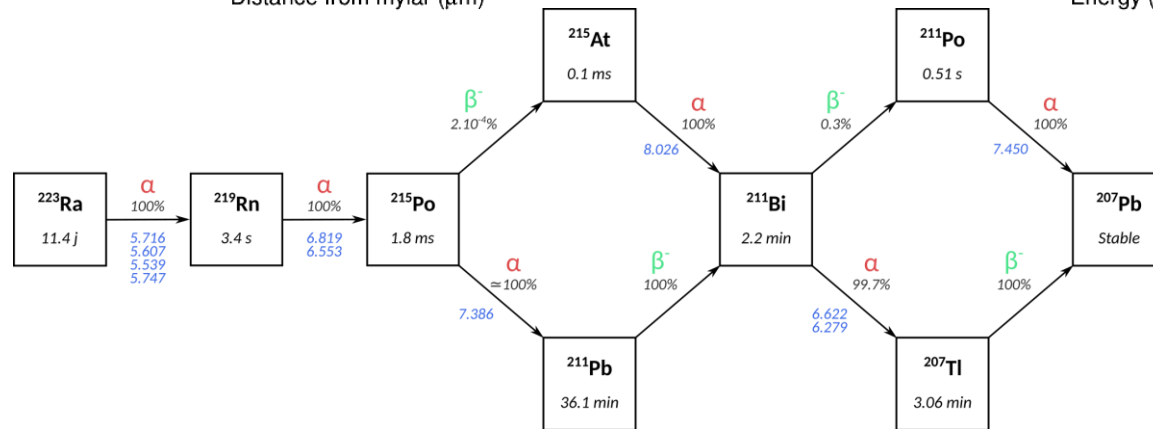


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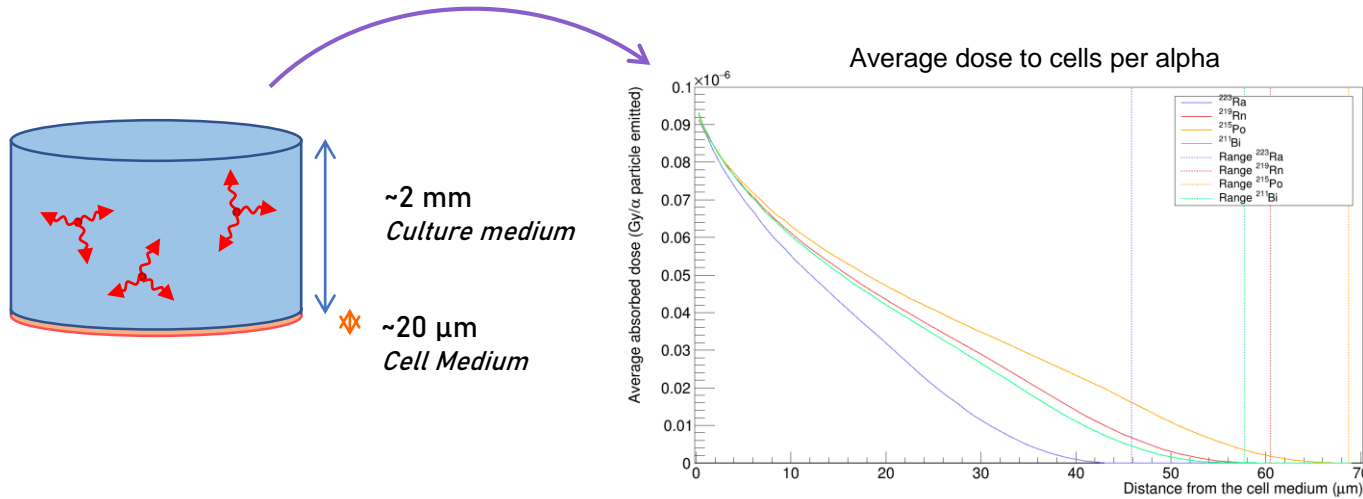


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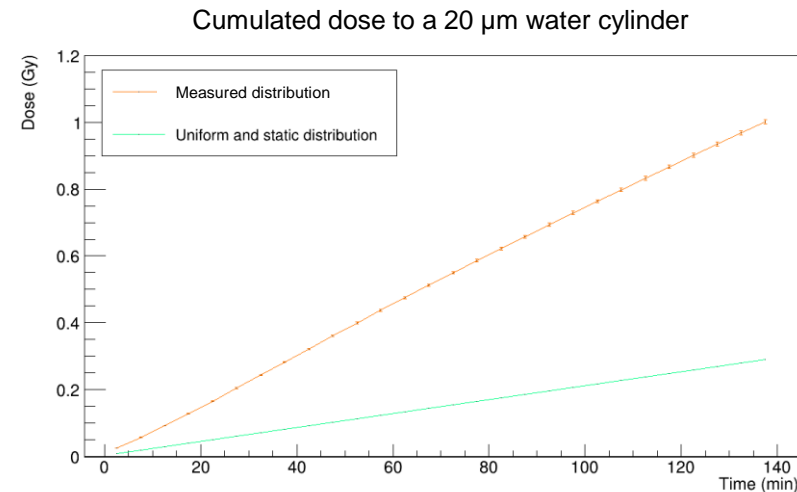
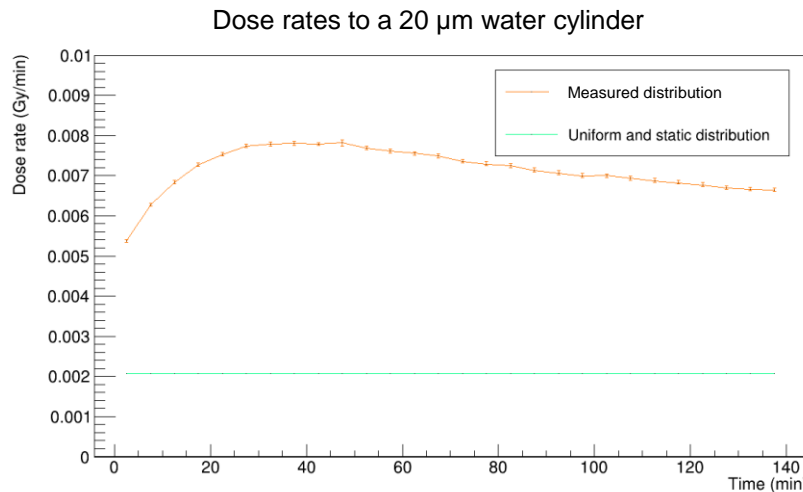
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# Measurements with $^{223}\text{Ra}$ (4/4) – Dose computation



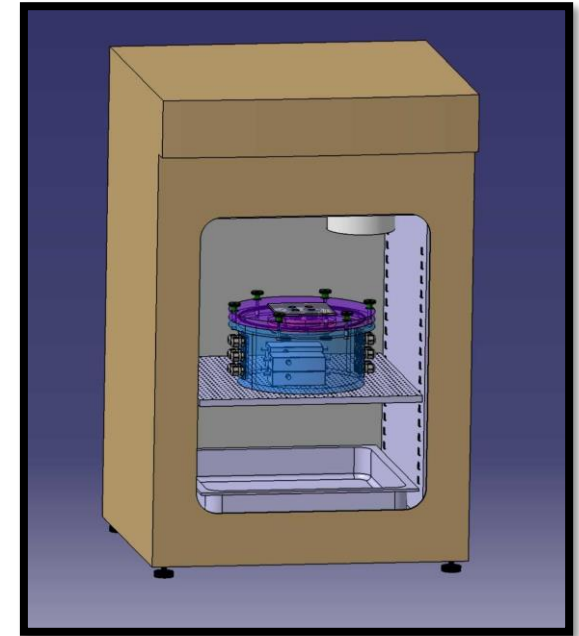
- Simulated cell geometry: 20 μm-high water cylinder.
- For every sampled height, mean delivered dose to the cells per α emitted is computed.
- Continuous aspects of dose rate and cumulated dose graphs: limited overfitting impact.
- Ongoing study.



3 to 4-fold underestimation



- ❑ Short  $\alpha$  emissions range: source of important *in vitro* dosimetry errors under homogeneity assumptions.
- ❑ Matrix deconvolution easily adjusts physical or hypothetical needs. Computational speed and portability of the set-up: on-site dose estimations.
- ❑ Noticeable overfitting for complex decay schemes. Consequences on dose computation uncertainties are currently evaluated.
- ❑ Cell modelling impact on dose computation is currently studied.
- ❑ First measurements of  $^{212}\text{Pb}$  injected in cell cultures under *in vitro* conditions: end of 2021.



Thank you for your attention.