## TOPAS-nBio Simulation of Radiation Chemistry Following FLASH Irradiation Including Reactions of Biological Importance

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## Presentation outline



- 1. INTRODUCTION
- 2. PURPOSE
- 3. METHODS
- 4. RESULTS

## Electron FLASH methodology Oxygen depletion hypothesis

- Delivering ≥ 10 Gy dose in a limited number of 1-2 Gy pulses
- Overall time ≤ 100 ms
- These high dose rates of irradiations have been shown to reduce radiation damage of healthy tissues, but not the tumor

- **Oxygen depletion** is one of the most commonly mentioned hypothesis to elucidate the FLASH effect
- In healthy tissues, the O<sub>2</sub> is depleted in a great extent, so that cells ares made transiently hypoxic and thus radioresistant
- In **tumor cells**, the O<sub>2</sub> difference is much smaller, therefore the TCP is maintained at similar level than in conventional radiotherapy CONV-RT



R. Labarbe et al. (2020) "A physicochemical model of reaction kinetics supports peroxyl radical recombination as the main determinant of the FLASH effect", Radiotherapy and Oncology 153: 303 J. D. Wilson et al. (2020) "Ultra-High dose Rate (FLASH) Radiotherapy: Silver Bullet or Fool's Gold?", Front. Oncol. 9: 1563

## Purpose

• Does the addition of biologically relevant moieties change the final O<sub>2</sub> concentration under FLASH vs. CONV conditions?

## PURE WATER MODEL





Adapted from: Wardman P. Radiotherapy Using High-Intensity Pulsed Radiation Beams (FLASH): A Radiation-Chemical Perspective. *Radiat Res.* 2020 May 20.

## Purpose

• Does the addition of biologically relevant moieties change the final O2 concentration under FLASH vs. CONV conditions?



#### Methods: Introducing biological models into TOPAS-nBio simulations



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• Water radiolysis reactions <sup>(1)</sup> and:

$$e_{aq}^{-} + O_2 \rightarrow O_2^{-}$$
  
 $H^{\bullet} + O_2 \rightarrow HO_2^{-}$ 

Reactions for simulation of radiolysis in pure liquid water.

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Reaction	k <sub>obs</sub> (/M/s)						
$e_{aq}^{-} + e_{aq}^{-} \rightarrow H_2 + OH^{-}$	5.5 x 10 <sup>9</sup>						
$e^{aq} + H_3O^+ \rightarrow H^{\bullet}$	2.3 x 10 <sup>10</sup>						
$e_{aq}^{-} + H^{\bullet} \rightarrow H_2 + OH^{-}$	2.5 x 10 <sup>10</sup>						
$e_{aq}^{-} + OH \rightarrow OH^{-}$	3.0 x 10 <sup>10</sup>						
$e_{aq}^{-} + H_2O_2 \rightarrow OH^{-} + OH^{-}$	1.1 x 10 <sup>10</sup>						
$H_3O^+ + OH^- \rightarrow H_2O$	14.3 x 10 <sup>10</sup>						
$H^{\bullet} + H^{\bullet} \rightarrow H_2$	7.8 x 10 <sup>9</sup>						
$H^{\bullet} + {}^{\bullet}OH \rightarrow H_2O$	1.55 x 10 <sup>10</sup>						
$H^{\bullet} + H_2O_2 \rightarrow {}^{\bullet}OH + H_2O$	9.0 x 10 <sup>7</sup>						
$^{\circ}OH + ^{\circ}OH \rightarrow H_2O_2$	5.5 x 10 <sup>9</sup>						

<sup>(1)</sup> Pimblott S (1992) *J. Phys. Chem.* **96** 4485–91 <sup>(2)</sup> Howard B and Michaels (1978) *Rad. Res.* **74** 23-34

### Methods: Introducing biological models to TOPAS-nBio simulations



#### Methods: Introducing biological models to TOPAS-nBio simulations



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DNA + e_{aq} \rightarrow DNA^{\circ}
DNA + H<sup>•</sup> \rightarrow DNA^{\circ}
DNA + OH<sup>•</sup> \rightarrow DNA-OH adduct
DNA-OH adduct + O2 \rightarrow DNA-OO
```

 $\begin{array}{l} \mathsf{RNA} + \mathsf{e}_{\mathsf{aq}} \to \mathsf{RNA}^{*} \\ \mathsf{RNA} + \mathsf{H}^{*} \to \mathsf{RNA}^{*} \\ \mathsf{RNA} + \mathsf{OH}^{*} \to \mathsf{RNA}^{*} \end{array}$ 

```
Proteins+ e_{aq} \rightarrow Proteins *
Proteins + H* \rightarrow Proteins *
Proteins + OH* \rightarrow Proteins *
```

Amino Acids +  $e_{aq} \rightarrow$  Amino Acids \* Amino Acids + H<sup>•</sup>  $\rightarrow$  Amino Acids \* Amino Acids + OH<sup>•</sup>  $\rightarrow$  Amino Acids

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Free Nu + e_{aq} \rightarrow Free Nu <sup>•</sup>
Free Nu + H<sup>•</sup> \rightarrow Free Nu <sup>•</sup>
Free Nu + OH<sup>•</sup> \rightarrow Free Nu <sup>•</sup>
```

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(Nu = Nucleotides)
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<sup>(1)</sup> Pimblott S (1992) J. Phys. Chem. 96 4485–91
 <sup>(2)</sup> Howard B Michaels and John Hunt (1978) Rad. Res. 74 23-34

#### Methods: TOPAS-nBio Monte Carlo track-structure simulations

- An extension of TOPAS tool for sub-cellular simulations.<sup>1</sup>
- Simulates the physical and chemistry stages of water radiolysis.
- The physical and pre-chemical stage of irradiation inherit the parameters provided by Geant4-DNA.<sup>2</sup>
- For chemistry, TOPAS-nBio version for this project uses Independent Reactions Time (IRT) with inter-track simulation capability.<sup>3</sup>



credits:https://gray.mgh.harvard.edu/research/software/258-topas-nbio

#### PULSE PARAMETERS

- The simulations were carried out for the three different models
- Total doses **up to 60 Gy** were used in 10 Gy steps<sup>4</sup>
- The system was irradiated by **1 MeV electron beam.**

Mode	Dose Rate (Gy * s <sup>-1</sup> )	Dose (Gy)	Pulse frequency (Hz)	Pulse width (µs)	Number of pulses	Treatment time (s)
CONV	0.29	10-60	10	1.0	350-2075	36-210
FLASH	500	10-60	100	1.75-1.9	2-12	0.01-0.11

<sup>1</sup> Schuemann J et al., (2019) Rad. Res. 191 125-138.

- <sup>2</sup> Ramos-Méndez J et al., (2018) Phys. Med. Biol. 63 105014 12pp.
- <sup>3</sup> Ramos-Méndez J et al., (2020) Rad. Res. **194** 351-362.
- <sup>4</sup> P. Montay-Gruel et al. (2019) Proc Natl Acad Sci. USA. **116**(22):10943-10951





# RESULTS

### Results: time evolution of O2



#### THE RESULTS FOR MODEL2 ARE STILL BEING ELABORATED

Ramos-Mendez et al., "LET-dependent intetrack yields in proton irradiation at ultra-high dose rates relevant for FLASH radiotherapy" Rad. Res. 194:351-362 (2020).

## Results: time evolution of H2O2





## Results: O2 depletion

Model 0 CONV-RT
Model 0 FLASH-RT



## Conclusions/summary



SIMULATIONS PERFORMED IN PURE WATER DO NOT REFLECT THE RADIATION CHEMISTRY GOING ON IN BIOLOGICAL SYSTEMS

- In this work we implemented three models to evaluate the chemical yields produced by low and high-dose rates.
- It was found that additional biological material affects significantly the yields compared to a pure liquid water model.
- The addition of a more detail model including more biological material affects the total yield of products like H<sub>2</sub>O<sub>2</sub> and O<sub>2</sub>.



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