



University of California
San Francisco

TOPAS-nBio for water radiolysis simulation

José Ramos-Méndez, PhD and Bruce Faddegon, PhD

Department of Radiation Oncology.

8/14/18



Outline



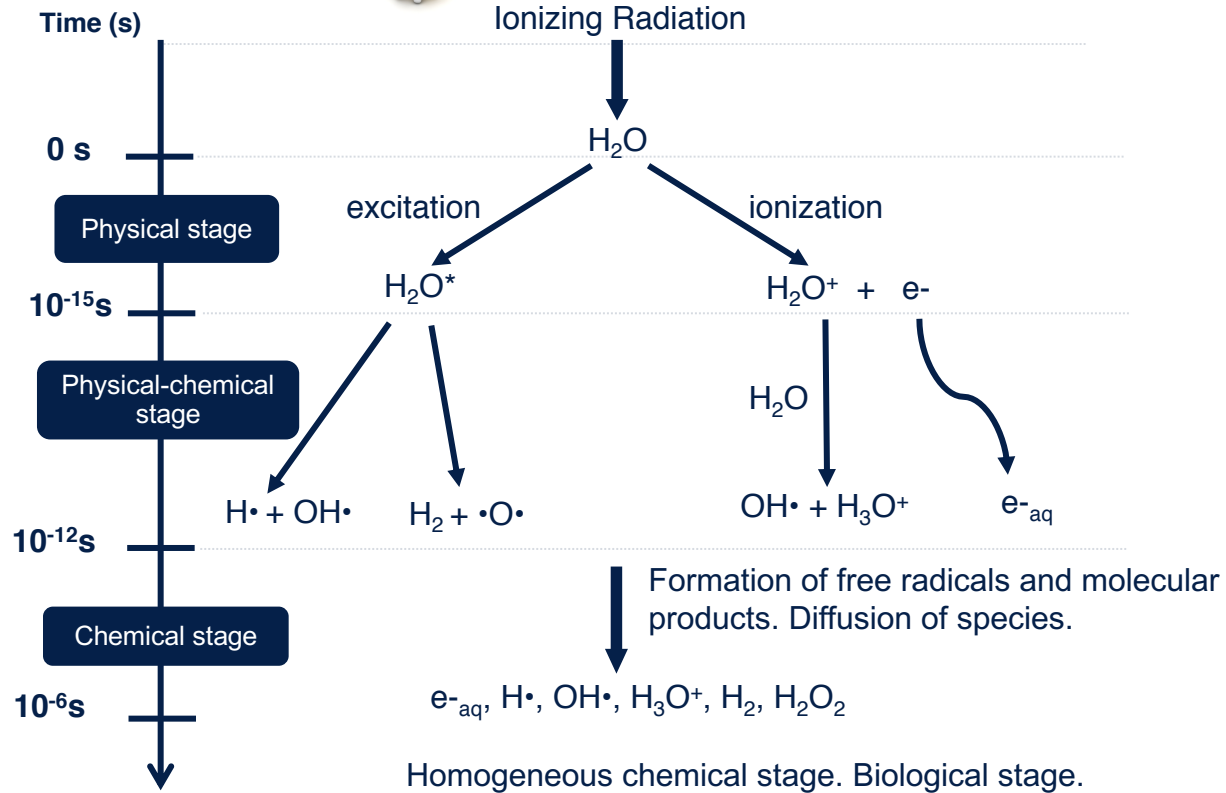
What?



Why?



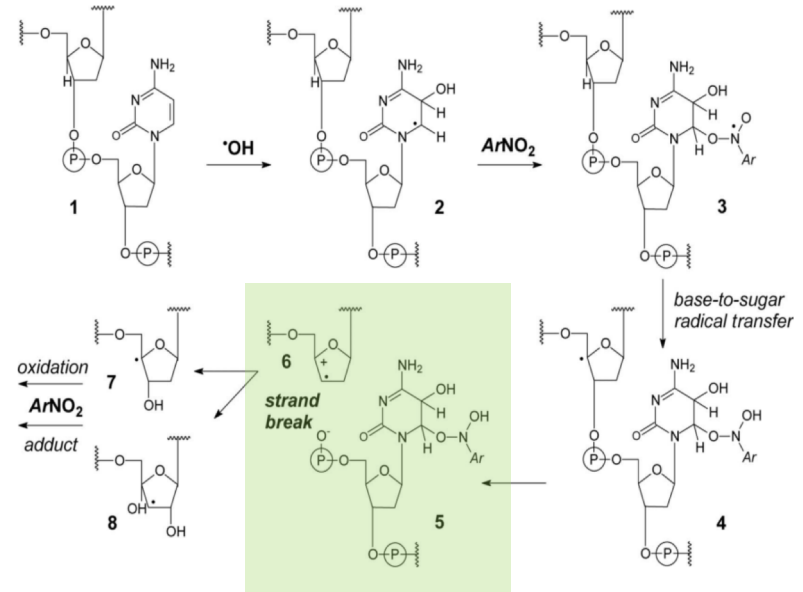
How?



Radiolysis of water

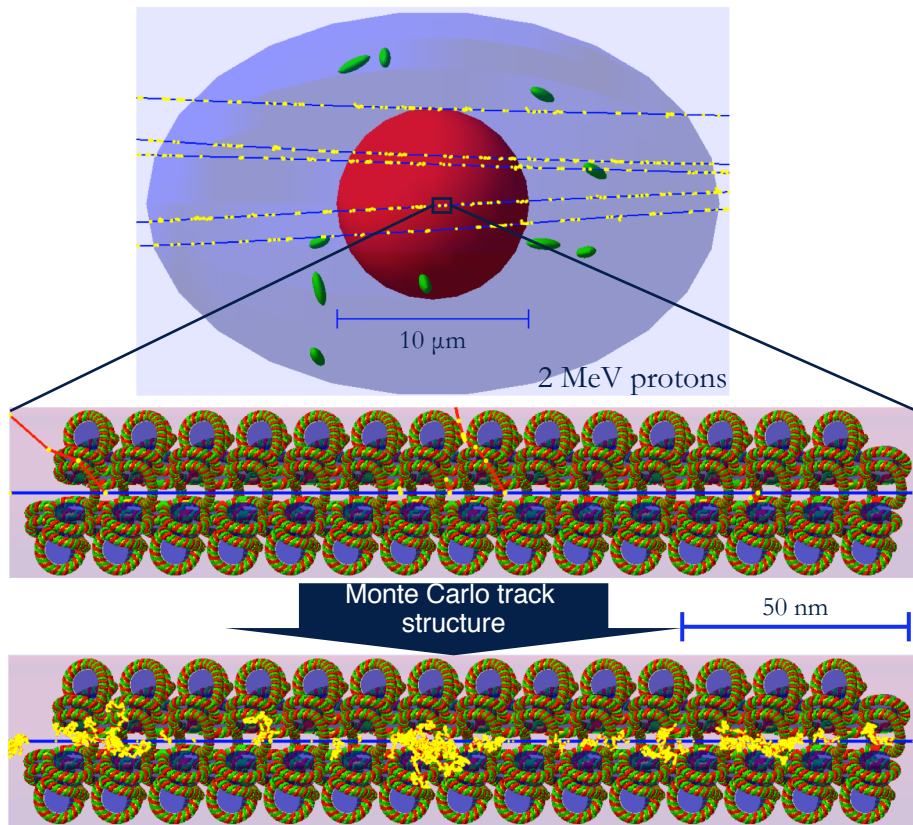


- DNA damage.
 - Formation of single-strand breaks.
 - Identification of clustered damage.
- Free radicals in radiation biology
 - Reactive oxygen
 - Radical repair by thiols
 - Cell signaling and bystander response
- Cancer treatment and diagnosis
 - Radiosensitizers –nitric oxide
 - Hypoxia
 - Radiation detector – Fricke dosimeter

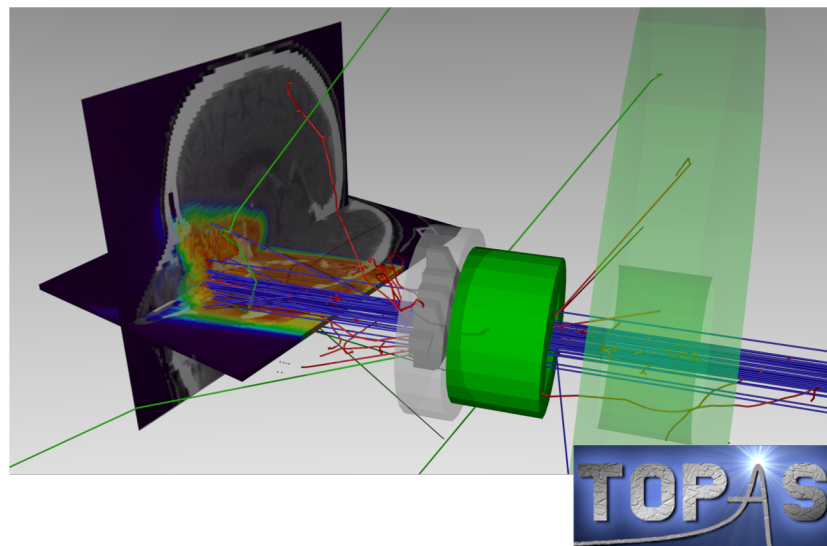


Formation of single-strand breaks by reaction of hydroxyl radical adducts of DNA bases with nitroimidazoles. From O'Neill and Wardman 2009, based on extensive search survey by von Sonntag (2006), cf. (Bamatraf et al. 1998, Wardman 2007a, 2009b)

Radiolysis of water



Condensed-history Monte Carlo



Radiolysis of water

- Track structure Monte Carlo codes are usually difficult to obtain.
- Geant4 Collaboration's¹ project: Geant4-DNA^{2,3}.
 - Steep learning curve.
- University of California San Francisco, Massachusetts General Hospital & SLAC National Laboratory developed TOPAS³ to facilitate MC for radiotherapy.
- TOPAS-nBio^{4,5,6} to facilitate MC track-structure simulations.



¹Agostinelli, S. *et al.* (2003) *Nucl. Inst. and Methods in Phys. Res. A*, 506(3), 250–303.

²Incerti, S. *et al.* (2010) *Med. Phys.*, 37(9), 4692–4708.

³Bernal, M. A. *et al.* (2015) *Physica Medica*, 31(8), 861–74.

⁴Perl, J. *et al.* (2012) *Med. Phys.*, 39, 6818–6837.

⁵McNamara, A. *et al.* (2017) "*Physica Medica*, 33, 207–215.

⁶Ramos-Mendez, J. *et al.* (2018) *Phys. Med. Biol.*. doi:10.1088/1361-6560/aac04c.

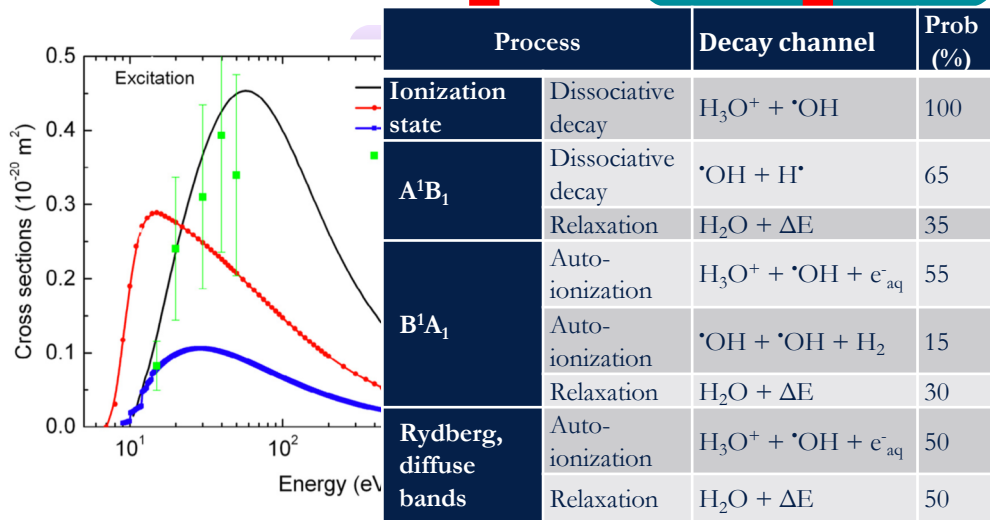
TOPAS-nBio Monte Carlo track structure simulations

Physics Stage
 step-by-step modeling of physics interactions of incoming & secondary radiation with biological medium

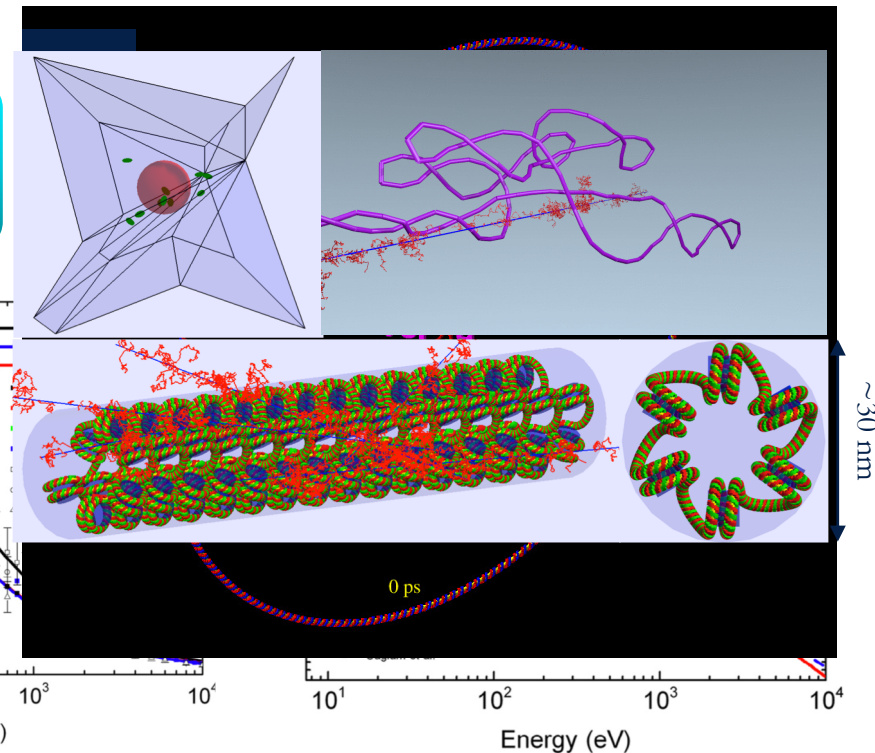
- Excited water molecules
- Ionized water molecules
- Solvated electrons

Physicochemical/chemistry Stage
 • Radical species production
 • Diffusion
 • Mutual interaction

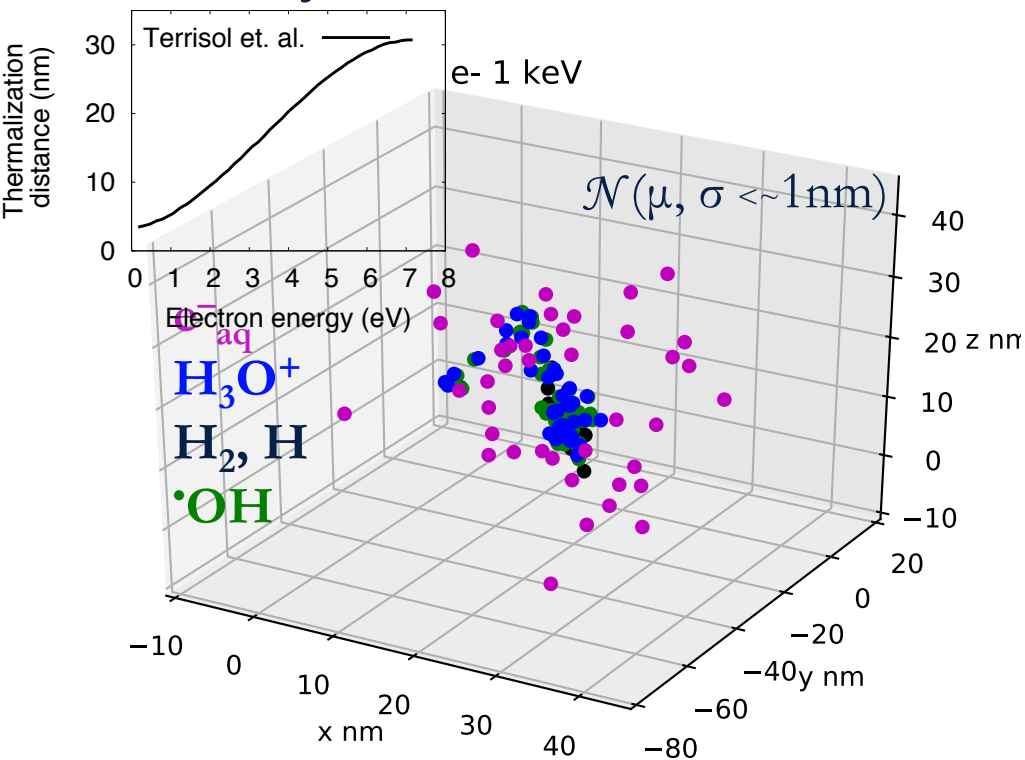
Kraath et al (2011) *J Phys Chem B* 115:11927-38
 Baraath et al (2005) *Physica Medica* 31:68-73



Mechanistic Biological modeling



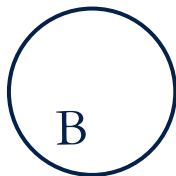
Radiolysis of water – Pre-chemical stage (10^{-15} - 10^{-12} s)



| Process | | Decay channel | Prob (%) |
|-----------------------------------|--------------------|--|----------|
| Ionization state | Dissociative decay | $\text{H}_3\text{O}^+ + \cdot\text{OH}$ | 100 |
| A¹B₁ | Dissociative decay | $\cdot\text{OH} + \text{H}\cdot$ | 65 |
| | Relaxation | $\text{H}_2\text{O} + \Delta\text{E}$ | 35 |
| B¹A₁ | Auto-ionization | $\text{H}_3\text{O}^+ + \cdot\text{OH} + \text{e}_{\text{aq}}^-$ | 55 |
| | Auto-ionization | $\cdot\text{OH} + \cdot\text{OH} + \text{H}_2$ | 15 |
| | Relaxation | $\text{H}_2\text{O} + \Delta\text{E}$ | 30 |
| Rydberg, diffuse bands | Auto-ionization | $\text{H}_3\text{O}^+ + \cdot\text{OH} + \text{e}_{\text{aq}}^-$ | 50 |
| | Relaxation | $\text{H}_2\text{O} + \Delta\text{E}$ | 50 |

Radiolysis of water – *Chemical stage* (10^{-12} - 10^{-6} s)

- Species move under Brownian motion
- Totally diffusion-controlled reactions



$$d < 2k/4\pi(D_A + D_B)$$

$$p(\mathbf{r}, t | \mathbf{r}_0) = \frac{1}{(4\pi Dt)^{n/2}} \exp\left[-\frac{(\mathbf{r} - \mathbf{r}_0)^2}{4Dt}\right]$$

Radiolysis of water – Chemical stage (10^{-12} - 10^{-6} s)

| Reaction | Reaction rate constant k_{obs} ($10^{10} \text{ M}^{-1}\text{s}^{-1}$) | |
|---|---|------------|
| | Geant4-DNA Default | TOPAS-nBio |
| $e_{\text{aq}}^- + e_{\text{aq}}^- \rightarrow \text{H}_2 + 2\text{OH}^-$ | 0.5 | 0.647 |
| $e_{\text{aq}}^- + \cdot\text{OH} \rightarrow \text{OH}^-$ | 2.95 | 2.95 |
| $e_{\text{aq}}^- + \text{H}^+ \rightarrow \text{H}_2 + \text{OH}^-$ | 2.65 | 2.65 |
| $e_{\text{aq}}^- + \text{H}_3\text{O}^+ \rightarrow \text{H}^+$ | 2.11 | 2.11 |
| $e_{\text{aq}}^- + \text{H}_2\text{O}_2 \rightarrow \text{OH}^- + \cdot\text{OH}$ | 1.41 | 1.41 |
| $\cdot\text{OH} + \cdot\text{OH} \rightarrow \text{H}_2\text{O}_2$ | 0.44 | 0.475 |
| $\cdot\text{OH} + \text{H}^+ \rightarrow \text{H}_2\text{O}$ | 1.44 | 1.44 |
| $\text{H}^+ + \text{H}^+ \rightarrow \text{H}_2$ | 1.2 | 0.503 |
| $\text{H}_3\text{O}^+ + \text{OH}^- \rightarrow \text{H}_2\text{O}$ | 14.3 | 11.0 |
| $\cdot\text{OH} + \text{H}_2 \rightarrow \text{H}$ | - | 0.0045 |
| $\cdot\text{OH} + \text{H}_2\text{O}_2 \rightarrow \text{HO}_2$ | - | 0.0023 |
| $\cdot\text{OH} + \text{HO}_2 \rightarrow \text{O}_2$ | - | 1.0 |
| $\cdot\text{OH} + \text{O}_2^- \rightarrow \text{O}_2 + \text{OH}^-$ | - | 0.9 |
| $\cdot\text{OH} + \text{HO}_2^- \rightarrow \text{HO}_2 + \text{OH}^-$ | - | 0.5 |
| $e_{\text{aq}}^- + \text{HO}_2 \rightarrow \text{HO}_2^-$ | - | 2.0 |
| $e_{\text{aq}}^- + \text{O}_2 \rightarrow \text{O}_2^-$ | - | 1.9 |
| $e_{\text{aq}}^- + \text{O}_2^- \rightarrow \text{OH}^- + \text{HO}_2^-$ | - | 1.3 |
| $\cdot\text{H} + \text{H}_2\text{O}_2 \rightarrow \cdot\text{OH}$ | - | 0.01 |
| $\cdot\text{H} + \text{HO}_2 \rightarrow \text{H}_2\text{O}_2$ | - | 2.0 |
| $\cdot\text{H} + \text{O}_2 \rightarrow \text{HO}_2$ | - | 2.0 |
| $\text{H} + \text{OH}^- \rightarrow e_{\text{aq}}^-$ | - | 0.002 |
| $\text{H} + \text{O}_2^- \rightarrow \text{HO}_2^-$ | - | 2.0 |
| $\text{H}_3\text{O}^+ + \text{O}_2^- \rightarrow \text{HO}_2$ | - | 3.0 |
| $\text{H}_3\text{O}^+ + \text{HO}_2^- \rightarrow \text{H}_2\text{O}_2$ | - | 2.0 |
| $\text{HO}_2 + \text{HO}_2 \rightarrow \text{H}_2\text{O}_2 + \text{O}_2$ | - | 0.000076 |
| $\text{HO}_2 + \text{O}_2^- \rightarrow \text{O}_2 + \text{HO}_2^-$ | - | 0.00274 |

| Chemical species | Name in TOPAS-nBio | Diffusion coefficients D ($10^{-9} \text{ m}^2 \text{ s}^{-1}$) at 25°C | | Reaction radius (nm) |
|------------------------|--------------------|---|------------|----------------------|
| | | Geant4-DNA Default | TOPAS-nBio | |
| e_{aq}^- | SolvatedElectron | 4.9 | 4.9 | 0.5 |
| $\cdot\text{OH}$ | Hydroxyl | 2.8 | 2.2 | 0.22 |
| H^+ | Hydrogen | 7.0 | 7.0 | 0.19 |
| H_3O^+ | Hydronium | 9.0 | 9.46 | 0.25 |
| H_2 | Dihydrogen | 5.0 | 4.8 | 0.14 |
| OH^- | Hydroxide | 5.0 | 5.3 | 0.33 |
| H_2O_2 | HydrogenPeroxide | 1.4 | 2.3 | 0.21 |
| O_2 | Oxygen | - | 2.1 | 0.17 |
| O_2^- | SuperoxideAnion | - | 2.1 | 0.22 |
| HO_2 | HydroPeroxide | - | 2.0 | 0.21 |
| HO_2^- | Dioxidanide | - | 2.0 | 0.25 |

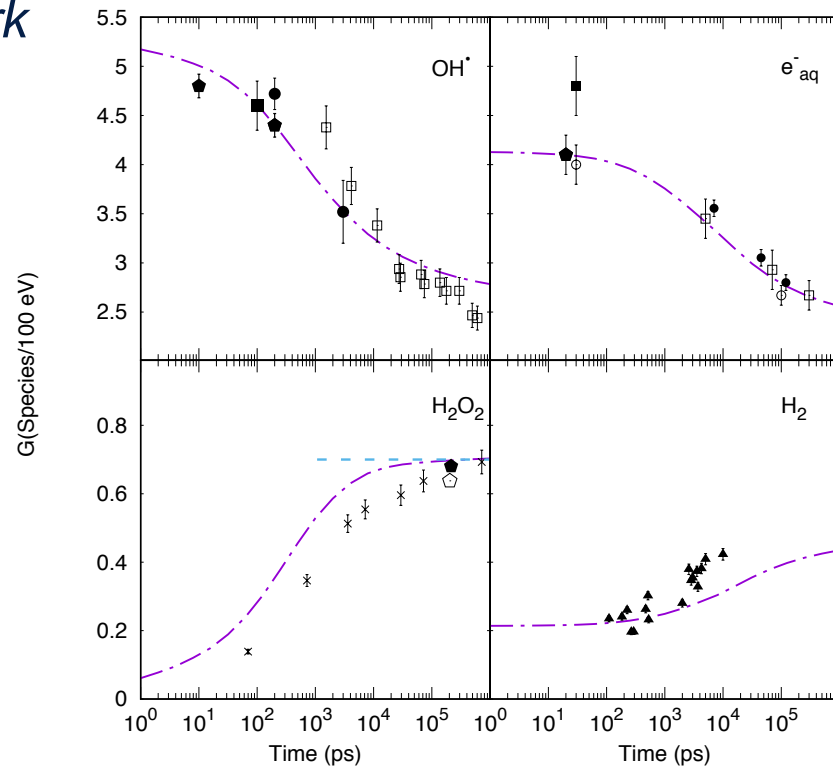
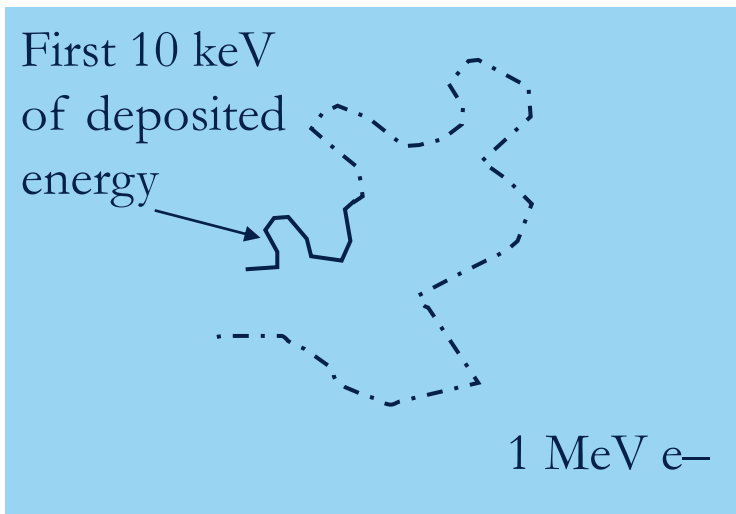
[1] M. S. Kreipl, W. Friedland, and H. G. Paretzke, “Time- and space-resolved Monte Carlo study of water radiolysis for photon, electron and ion irradiation,” *Radiat. Environ. Biophys.*, vol. 48, no. 1, pp. 11–20, 2009.

[2] J. Ramos-Mendez, J. Perl, J. Schuemann, A. McNamara, H. Paganetti and B. Faddegon “Monte Carlo simulation of chemistry following radiolysis with TOPAS-nBio” *Phys Med Biol.* 64, 105104-12pp (2018)

[3] S. Uehara and H. Nikjoo, “Monte Carlo simulation of water radiolysis for low-energy charged particles.,” *J. Radiat. Res.*, vol. 47, no. 1, pp. 69–81, 2006.

Radiolysis of water – Ongoing work

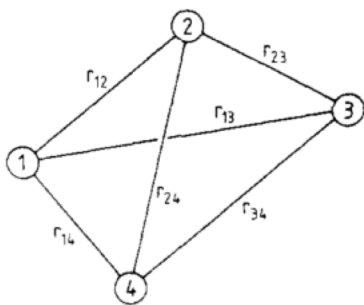
G-values: Yield of chemical species per 100 eV of deposited energy.



Ramos-Mendez J, et. al., Phys Med Biol. 64, 105104-12pp (2018)

Independent reaction times.

- Alternative to the step-by-step Monte Carlo for the chemistry stage.
- It is a stochastic method developed more than 30 years ago¹.



| Pair | r_{ij}/nm | t_{ij}/ps | Reaction |
|------|--------------------|--------------------|----------|
| 1+2 | 1.3 | 12 | x |
| 1+3 | 2.1 | 27 | x |
| 1+4 | 1.0 | 40 | 2 |
| 2+3 | 1.1 | 8 | 1 |
| 2+4 | 1.7 | 125 | x |
| 3+4 | 1.8 | ∞ | x |

From Green et. al. J. Phys. Chem. 1990

Totally diffusion-controlled reactions

$$W(r, t) = \frac{R_{eff}}{r_{eff}} = \text{erfc} \left(\frac{r_{eff} - R_{eff}}{\sqrt{4D't}} \right)$$

Partially diffusion-controlled reactions

$$W(r, t) = \frac{R_{eff}\alpha + 1}{r_{eff}\alpha} \left[-F \left(\frac{r_{eff} - R_{eff}}{\sqrt{4D't}}, -\alpha\sqrt{D't} \right) + \text{erfc} \left(\frac{r_{eff} - R_{eff}}{\sqrt{4D't}} \right) \right]$$

$$F(x, y) = \exp(2xy + y^2) \text{erfc}(x + y)$$

¹Clifford et. al. J. Chem. Soc., Faraday Trans 1, 1986

IRT

Independent Reaction Times
(71 reactions, 15 species)

Totally diffusion controlled
(10 reactions)

Partially diffusion controlled
(42 reactions)

First order reactions
(19 reactions)

Neutral species

$$P_I(t | r_0) = \frac{\sigma}{r_0} \operatorname{erfc} \left(\frac{r_0 - \sigma}{\sqrt{4Dt}} \right)$$

Charged species

$$P_{III} = \frac{\sigma_{eff}}{r_{eff}} \operatorname{erfc} \left(\frac{r_{eff} - \sigma_{eff}}{\sqrt{4Dt}} \right)$$

Neutral species

$$P_{II}(t | r_0) = \frac{\sigma\alpha + 1}{r_0\alpha} \left[\operatorname{erfc} \left(\frac{r_0 - \sigma}{\sqrt{4Dt}} \right) - W \left(\frac{r_0 - \sigma}{\sqrt{4Dt}}, -\alpha\sqrt{Dt} \right) \right]$$

Charged species

$$P_{IV} = \frac{\sigma''_{eff}}{r_{eff}} [\operatorname{erfc}(b) - W(a, b)]$$

$$W(x, y) = \exp(2xy + y^2) \operatorname{erfc}(x + y)$$

All species

$$P_{VI} = 1 - \exp(-k[B]\Delta t)$$

[B]: concentration of the specie
k: reaction rate
k[B]: scavenging power/capacity

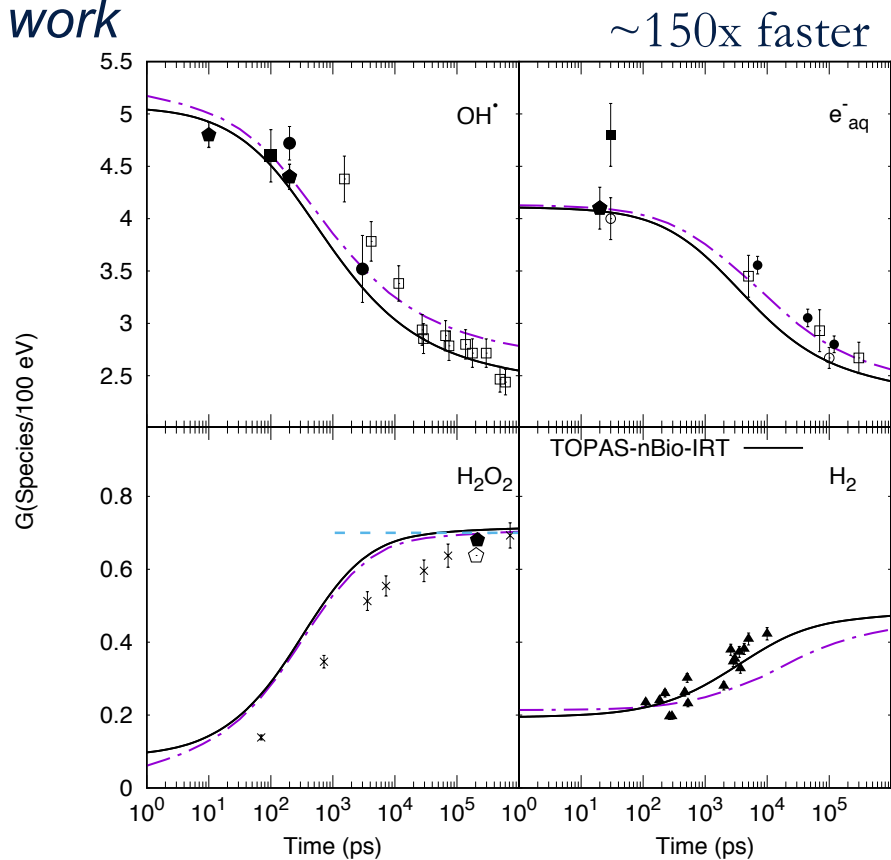
Radiolysis of water – Ongoing work

G-values: Yield of chemical species per 100 eV of deposited energy.

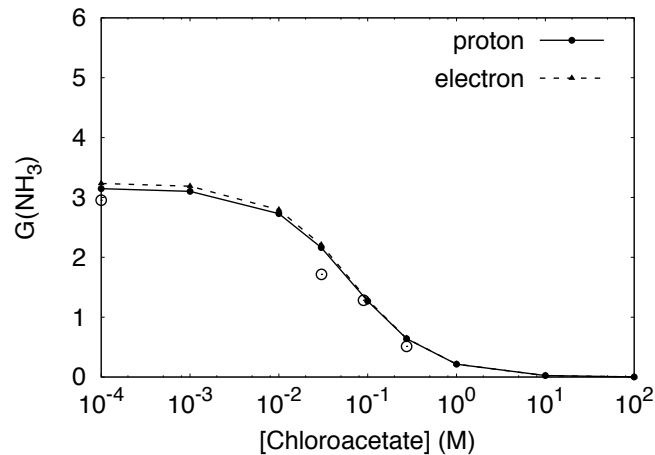
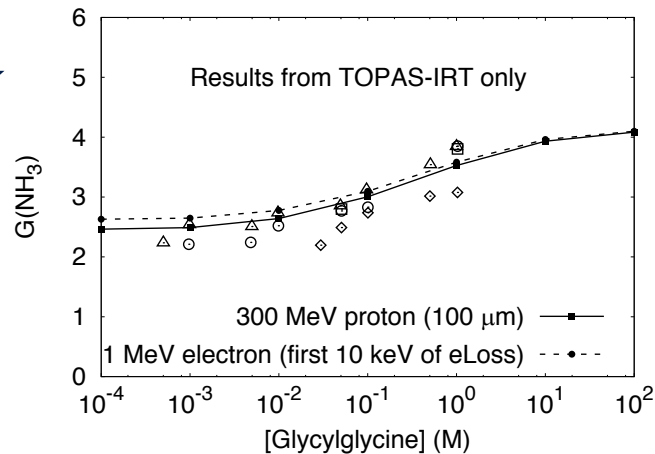
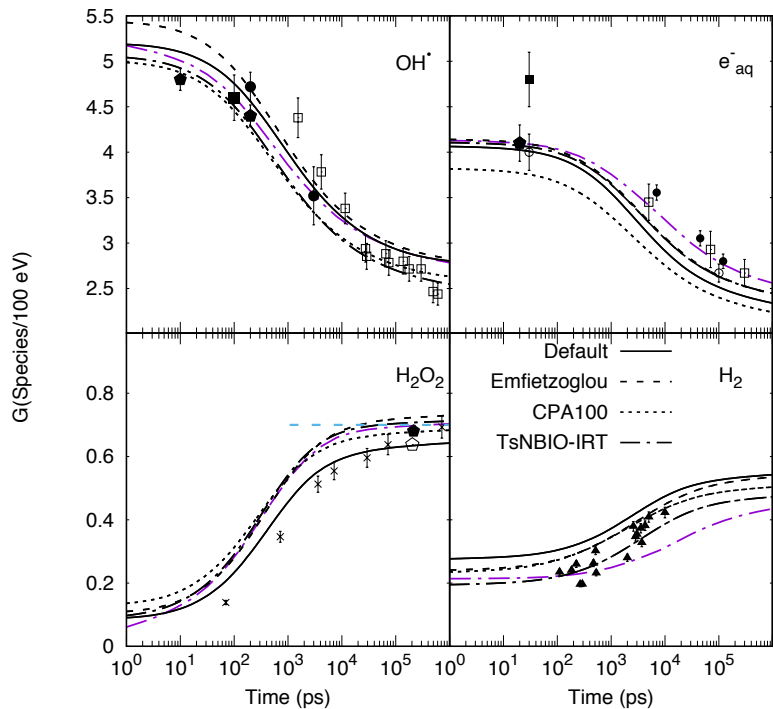
First 10 keV
of deposited
energy



1 MeV e⁻



Radiolysis of water – Ongoing work



Conclusion

- TOPAS-nBio wraps and extends Geant4-DNA for track-structure simulations.
- It provides with high flexibility to combine complex geometry volumes with sophisticated scoring routines.
- It facilitates the validation of Geant4-DNA.
- Still an alpha version, so new users are welcome.

Acknowledges

- NIH grants 1R01CA187003 and 1P20CA183640.
- TOPAS and TOPAS-nBio Collaboration.
- Naoki Dominguez from Benemérita Universidad Autónoma de Puebla, México.
- Dr. Aimee McNamara from Massachusetts General Hospital.
- Dr. Ianik Plante, NASA.
- Geant4-DNA Collaboration, Sebastien Incerti and Mathieu Karamitros.
- Dr. Eduardo Moreno from Benemérita Universidad Autónoma de Puebla, México.



University of California
San Francisco