



Radiation dosimetry at cellular level

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Introduction

- radiation therapy: kill cells selectively
- external radiation therapy
- 'internal' radiation therapy
 - targeted radiation therapy
 - boron neutron capture therapy (BNCT)

(LCE Friday seminar on BNCT in April)







Introduction

- energy deposition or dose D: energy/mass J/kg=Gy
 - external radiation therapy: broad, macroscopic
 - internal radiation therapy: inhomogeneous, microscopic
 - example: S-factors for a water sphere
 - activity inside a spherical volume
 - how much dose absorbed inside this volume







 $r(\mu m)$







Introduction

- this work
 - dosimetry of small metastases
 - activity entirely in the tumor cells
 - compare
 - three simple models to build cell cluster
 - three different isotopes: ¹¹¹In, ^{113m}In, ^{114m1}Inc







Radiation spectra





Radiation spectra



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Monte Carlo simulations

• EGS4 (Electron-Gamma Shower) simulation code

http://www.slac.stanford.edu/egs/

- EGS introduced in 1978
- free for non-commercial use
- applied in high-energy, nuclear and medical physics
 - electrons and photons from keV to 100 GeV
- written in MORTRAN a preprocessor for FORTRAN77 !!
 - conversion to C: EGSnova http://www.nemc.org/nova/







Monte Carlo simulations

- event-driven MC simulation:
 - follow particles from one scattering event to another
 - distance between events: random variable, calculated from total scattering cross section: Σ_t

MFP
$$\lambda = \frac{1}{\Sigma_t}$$
 $P\{\text{interact. in } dx\} = \frac{dx}{\lambda}$

$$N_{\lambda} = \int_{x_0}^{x} \frac{\mathrm{d}x}{\lambda(x)} \qquad P\{\hat{N}_{\lambda} < N_{\lambda}\} = 1 - \exp(-N_{\lambda})$$

number of MFP's sampled as $\ N_\lambda = -\ln\xi, \quad \xi \in [0,1[$



Monte Carlo simulations

- calculate many trajectories
 - spatial distribution of energy deposition
- dose kernel: fraction of the decay energy from a spherical source deposited at distance r from the center of the source

$$F(r) = \frac{1}{T_0} \frac{\delta E(r)}{\delta r}$$



• dose from a tumor cell to a healthy cell at a distance r

$$D_{\rm th}(r) = \frac{T_0}{m_j} \int_{r-r_j}^{r+r_j} F(r') A_j(r') dr'$$

• average dose to healthy cells

$$\overline{D}_{\text{th}} = \frac{1}{N_{\text{h}}} \sum_{i \in \text{tumor}} D_{\text{th}} (|\mathbf{r}_{i} - \mathbf{r}_{j}|)$$

$$\sum_{i \in \text{healthy}} D_{\text{th}} (|\mathbf{r}_{i} - \mathbf{r}_{j}|)$$

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Cell cluster models

- close-packed cubic geometry (i.e. fcc)
 - both tumor and healthy cells in fcc lattices
 - avoid cell overlap \rightarrow gap between tumor and healthy tissues
 - fast





Cell cluster models

- brute force packing
 - step spherical coordinates r, θ, φ
 - check for overlap (use cell method to calculate intercell distances)
 - no large gap between tumor and healthy tissues
 - non-optimum packing in bulk tissue
 - slow





Cell cluster models

- optimization through hard sphere collisions
 - initial configuration: two fcc lattices, sphere radii 60% and 93% of the final values
 - during the simulation radii are slowly increased towards the final values
 - event-driven MD:
 - calculate time to next collision:

$$|\mathbf{r}_{2} + \mathbf{v}_{2}t_{c} - \mathbf{r}_{1} - \mathbf{v}_{1}t_{c}|^{2} = (r_{10} + r_{c1}t_{c} + r_{20} + r_{c2}t_{c})^{2}$$

- advance spheres to time t_c
- collision: exchange normal components of the velocities







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• cluster geometry



close-packed

brute force

optimization





• cluster geometry







• dose distributions:

therapeutic effect

$$\frac{\overline{D}_{\text{tumor}}}{\overline{D}(r)} = \begin{vmatrix} \text{ratio of th} \\ \text{tumor ce} \\ \text{dose of th} \end{vmatrix}$$

ratio of the mean absorbed dose in tumor cells to the mean absorbed dose of the healthy cells inside radius *r*







 dose distributions







Conclusions

- cell model important
- future studies: model based on microscopy data of real tissue



- experimental data?
- BNCT: energy deposition also by ions (α , ⁷Li)
 - MD? (overkill)
 - binary collision approximation (BCA) codes (e.g. SRIM http://www.srim.org/)