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# Applying radiobiology to clinical molecular radiotherapy

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### ARTICLE INFO

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The aim of radiation therapy, including molecular radiotherapy (MRT), is to maximise the biological damage to a target tissue, most commonly malignant cancer cells, whilst minimising the damage to healthy organs. Physics, chemistry and biology all have a role to play in achieving this aim. Appropriate selection and development of the radiopharmaceutical will result in preferential uptake of radioactivity within a specific biological target, but little retention in healthy organs. Imaging enables *in-vivo* verification that this has been achieved at the pre-clinical development stage or for individual patients in the clinic. Quantification of those images is the first step in calculating radiation absorbed doses delivered to both target cells and normal organs. The second step is to model the deposition of energy from distributions of radioactivity, either by bespoke Monte Carlo simulation, convolution of a dose point kernel or application of reference dose-factors which describe radiation transport within a reference geometry [1–4].

Estimates of radiation absorbed doses are key to understanding the likely impact of treatments for which radiation is the primary driver of biological response. The biological response to radiation can be modulated by physical and biological factors. Pre-clinical investigations are critical to the understanding of various factors that impact clinical efficacy of a new radiotherapeutic and can inform initial treatment protocols, saving unnecessary and costly clinical studies.

It has long been understood that the rate at which radiation is delivered has a differential effect on cancer tissues and normal tissues [5]. As a result, a total prescribed dose of external beam radiotherapy (EBRT) is usually delivered in a series of daily fractions over a period of weeks. Development of the linear quadratic (LQ) model of cell death alongside the frameworks of the biologically effective dose (BED) and equivalent 2 Gy fraction dose (EQD2) has allowed the comparison of different fractionation schemes [6,7]. It is worth noting that the LQ model does not describe all aspects of EBRT radiobiology, for example the FLASH phenomenon where delivery of EBRT doses over milliseconds rather than minutes appears to lead to normal tissue sparing [8]. Nonetheless as MRT results in the continuous delivery of exponentially decreasing levels of radiation, it can be considered within the LQ framework as the delivery of infinitely small fractions of radiation and infinitely small intervals. Consequently, extensions of the LQ model have been developed specifically for MRT in order to allow comparison of doses delivered at differing dose-rates [9,10]. This is particularly useful in for MRT since inter-patient pharmacokinetics will lead to inter-patient differences in both absorbed doses and dose-rate within the same normal organs. For example, in the case of 18 patients treated with [90Y]Y-DOTA-TOC, the kidney BED was shown to have a higher correlation with renal toxicity than the absorbed dose alone [11]. These data were further analysed within MIRD pamphlet 20 alongside data from an additional 25 patients who also received [90Y]Y-DOTA-TOC [12]. The authors compared normal tissue complication probabilities (NTCP) in these cohorts as a function of BED and found them to match those derived from patients undergoing EBRT. Hence it could be concluded that the same radiobiological model predicted the effect of both EBRT and MRT in relation to renal toxicity. However, the same analysis applied to patients treated with [177Lu]Lu-DOTA-TATE has demonstrated an increased renal tolerance to radiation [13]. An explanation for this was provided by ex-vivo autoradiography studies which demonstrated significant heterogeneity of [<sup>111</sup>In]In-DTPA-octreotide within kidney tissue [14]. Simulated absorbed dose distributions of <sup>90</sup>Y from these data were relatively uniform due to the longer length of <sup>90</sup>Y beta particles. Simulated <sup>177</sup>Lu absorbed dose distributions matched the underlying heterogeneity of the radiopharmaceutical distribution due to much smaller beta particle path lengths. However, SPECT imaging cannot measure such heterogeneities due to their limited spatial resolution. Therefore it could be concluded that the effect of absorbed dose heterogeneity was to modulate the apparent response to radiation doses that were measured on a macroscopic scale. Equally, analysis of hepatocarcinoma and metastatic colorectal cancer patients treated with 90Y microspheres have demonstrated apparent tumour radiosensitivies lower than those observed in response to EBRT [15-17]. Again, heterogeneity at a microscopic level has been proposed as an explanation

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for this observation. Although models have been proposed to incorporate the effect of dose non-uniformity, [18,19] quantitative measurements may be beyond the spatial resolution of gamma camera imaging.

Therefore, the development of both alpha particle [20] and Auger electron [21] emitting radiotherapeutics provides new challenges with respect to establishing and understanding a dose-response relationship. Both alpha particles and Auger electrons have a high linear energy transfer (LET) and consequently deposit their energy over micrometre and nanometre ranges respectively, compared to the millimetre range associated with beta particles. Hence, heterogeneity of uptake will result in heterogeneous radiation dose distributions. For Auger electron emitters in particular, knowledge of their localisation within the cell is critical to prediction of the absorbed dose delivered to the cell nucleus. Models describing the deposition of energy to particular cell compartments have been published [22,23].

Heterogeneity is not always taken into account in clinical dosimetry. For example, in the case of the alpha emitter [<sup>223</sup>Ra]RaCl<sub>2</sub> the distribution of activity within bone tissue is assumed to be concentrated within the endosteal layer rather than uniformly throughout the entire bone volume [24]. However, recent publications describing the clinical dosimetry of  $^{\rm 223}{\rm Ra}$  have assumed a uniform distribution of activity within prostate cancer bone metastases and have reported the mean absorbed dose to the entire lesion volume [25,26]. Such an approach also potentially overlooks the stochastic nature of energy deposition at the microscopic scale for high LET particles emitted by highly localised radionuclides. In such scenarios, energy deposition in target cells is better described by probability density functions rather than a single discrete value [27,28]. However, application of these stochastic models demonstrate that as the number of emitted particles increases, the relative deviation around the average specific energy will decrease, such that a macroscopic description using absorbed dose will suffice [27]. This has been demonstrated at the in-vitro level [29] and thus may indicate why the <sup>223</sup>Ra macroscopic absorbed doses to bone metastases correlate with reported lesion response [25].

Classical microdosimetry models only describe the deposition of energy to a biological target. The radiobiology of short range radiation such as alpha particles or Auger electrons is further complicated by the knowledge that cellular response to radiation is not limited to direct effects [30]. The bystander effect is an umbrella term for the phenomenon whereby irradiation of a single cell can induce cell death in neighbouring cells, either via direct cell to cell communication or via the release of soluble factors [31] and is not exclusive to alpha particles and Auger electrons. The effect is distinct from the crossfire effect which describes the irradiation from targeted cells to non-targeted cells. Bystander effects have been demonstrated in response to EBRT as well as alpha, beta an Auger electron therapy [31,32]. However, for high LET radiation it is possible that this biological effect will extend beyond the range of radiation with a path length of the order of a cell diameter (or less) and may therefore compensate for heterogeneity of uptake in the absence of a crossfire effect. A number of mathematical models of the bystander effect have been developed [33-35] but overall the clinical relevance is not well understood. Furthermore the immune response is thought to be the biological mechanism behind the abscopal effect whereby EBRT is observed to elicit a response beyond the irradiated volume [36,37]. Whilst mathematical models describing the synergy between immunotherapy and radiotherapy have been developed [38] the relevance of this phenomenon to systemically administered MRT is also unclear.

High LET particles also have a higher relative biological effectiveness (RBE), defined as the ratio of low LET absorbed dose compared to the high LET absorbed dose required to achieve a defined biological endpoint [39]. RBE values for alpha particles have typically been found to have a value between 3 and 5 *in-vitro* [40]. In the absence of human *in-vivo* data a value of 5 is normally assumed [41]. The LQ framework has been extended to alpha particles in order to incorporate RBE into the calculation of both the BED [9,42] and the EQ2D [43]. Thus predictions of efficacy or toxicity can be made using tissue radiosensitivity

parameters derived from EBRT data. Such calculations were recently applied in a study by Belli et al. of dose to the salivary glands and the probability of xerostomia in metastatic castrate resistant prostate cancer patients treated with the alpha emitter [225Ac]Ac-PSMA-617 [44]. (Salivary glands express PSMA and are therefore known to be organs at risk in these treatments.) The authors simulated potential [<sup>225</sup>Ac]Ac-PSMA-617 absorbed doses by converting dosimetry data acquired in a cohort of patients undergoing [177Lu]Lu-PSMA-617 treatment. Absorbed doses were converted to both BED and EQ2D and used to predict NTCP as a function of the administered activity of [<sup>225</sup>Ac]Ac-PSMA-617. The authors then compared these predictions to recorded observed instances of xerostomia in a separate cohort of patients treated with [<sup>225</sup>Ac]Ac-PSMA-617 (but lacking dosimetry data). They noted a significant difference between predictions of xerostomia based on the LQ formalism and those observed clinically. As the authors themselves suggested, this discrepancy could be due to a lack of consideration of absorbed dose heterogeneity within the salivary glands as well as uncertainty around the value of the RBE.

Finally it should be noted that the LQ model is a mechanistic model of radiation damage and does not take into account the tumour microenvironment or systemic reactions. Over the past twenty years the field of radiobiology has been revitalised by a succession of new discoveries and observations [45]. The biochemical response of cells exposed to radiation damage is increasingly well understood, resulting in a new generation of radiation sensitizers. For example, Olaparib is an inhibitor of PARP-1, a DNA damage sensor protein associated with DNA repair [46]. Preclinical investigations of radiation sensitizers in combination with molecular radiotherapy have demonstrated encouraging efficacy regarding the treatment of cancer cells [47]. When combined with EBRT the potential for increased radiation sensitivity and toxicity in normal organs has been noted [48]. It would seem prudent to assume the same could be true in MRT. Therefore as clinical trials of radiation sensitizers in combination with molecular radiotherapy are developed, further modulations to existing radiobiological models are likely to be required.

In summary, initial applications of radiobiological modelling to [90Y]Y-DOTA-TOC therapy suggested that models developed for EBRT could be applied to MRT. This is an attractive proposition due to the considerable body of literature describing the response of both normal organs and tumour tissue in EBRT [49,50]. As new therapeutic modalities such as targeted alpha therapies are introduced, a common radiobiological framework would allow fair comparison of alternative strategies. However, it has been demonstrated that even for lower energy beta emitters such as <sup>177</sup>Lu, deviations from uniform dose deposition within tissue will result in apparently different radiosensitivities to those predicted. Strategies to model heterogeneity beyond the spatial resolution of gamma camera imaging have been proposed. Digital autoradiography performed on ex-vivo samples can be used to relate the mean absorbed dose within an organ to the absorbed dose within a particular sub-volume of interest [41]. A drawback of such an approach is that the same model would need to be applicable to a population and would result in a systematic shift in measured doses. Alternatively, kinetic modelling of radiopharmaceuticals has been proposed as a means of inferring time activity curves within sub volumes of tissue [41,51].

As clinical trials of new MRT strategies are developed, it is important that adequate dosimetry is conducted. In order to understand and inform the modulations to current radiobiological models that are required to address the issues raised in this commentary, dosimetry metrics relating to individual outcomes need to be collected. In the first instance they can be used to test dose-response relationships predicted by pre-clinical radiobiology. When discrepancies between predicted and observed outcomes occur, the same data can be used to refine the models and further the development of clinical protocols that fully realise the potential of this treatment modality.

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