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Bystander signalling: Exploring clinical relevance through new approaches and new models

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Abstract

Classical radiation biology research has centred on nuclear DNA as the main target of radiation induced damage. Over the past two decades, this has been challenged by a significant amount of scientific evidence clearly demonstrating radiation induced cell signalling effects to have important roles in mediating overall radiobiological response. These effects, generally termed radiation induced bystander effects (RIBEs) have challenged the traditional DNA targeted theory in radiation biology and highlighted an important role for cells not directly traversed by radiation. The multiplicity of experimental systems and exposure conditions in which RIBEs have been observed has hindered precise definitions of these effects. However, RIBEs have recently been classified for different relevant human radiation exposure scenarios in attempt to clarify their role in vivo. Despite significant research effort in this area, there is little direct evidence for their role for in clinically relevant exposure scenarios. In this review, we explore the clinical relevance of RIBEs from classical experimental approaches through to novel models that have been used to further determine their potential implications in vivo.

Key words

bystander effect, in vivo, ionising radiation, microbeam

Searches

Articles cited in this manuscript were sourced through a literature search on the Medline/Pubmed database using the search terms ‘radiation’, bystander’ ‘in vivo’ and ‘microbeam’. Full articles were retrieved when the abstract was deemed relevant. The bibliographies of retrieved papers were also searched and relevant articles included.

Conflict of Interest

The authors have no conflict of interest to declare.
**Introduction**

Ionising radiation (IR) is an effective cancer therapy due to its ability to induce cell death as a consequence of DNA damage resulting from energy deposition in the cellular environment. Most cellular responses to IR are mediated through direct energy deposition in the DNA or indirectly through reactive oxygen species (ROS) and other free radicals formed due to the radiolysis of water (1).

Classically, radiation biology research has focussed on nuclear DNA as the sole target of radiation induced damage. However, over the last 25 years a large body of scientific evidence has challenged the view that radiobiological responses occur only in cells directly targeted by radiation as biological effects have been shown to occur outside of the radiation target. These “Non-targeted effects” include genomic instability and several radiation induced signalling effects (2), generally termed radiation induced bystander effects (RIBEs). RIBEs were first identified by Nagasawa and Little (3) who observed chromosome damage in the form of sister chromatid exchanges in more than 30% of a cell population under conditions in which only 1% of cell nuclei had been targeted using α-particles. Since then, RIBEs have been demonstrated using a range of experimental systems with multiple biological endpoints. Despite increasing evidence in a growing number of model systems, the implications of RIBEs for radiotherapy and cancer risk remain to be fully determined. In this review, we describe RIBEs in the context of current experimental and clinical exposure scenarios and consider potential implications for cancer risk and radiotherapy.

**Discussion**

**Defining RIBEs**

In general terms, RIBEs may be defined as radiobiological responses observed in cellular systems which have not been directly traversed by ionising radiation but are in close proximity to irradiated cells. These effects are cell signal mediated either through direct physical cell contact via gap junction intercellular communication (GJIC) (4) or through secreted, diffusible signalling molecules into the surrounding media (5-7). A number of candidate signalling molecules have been identified in
mediating RIBEs such as reactive oxygen and nitrogen species (ROS/NOS) including nitric oxide (NO), and cytokines such as transforming growth factor-β (TGF-β) and interleukin-8 (IL-8). These have been shown to initiate multiple downstream signalling pathways including the mitogen activated protein kinases (MAPKs) and nuclear factor-κB (NF-κB) pathways (8).

Although RIBEs can be considered primarily as signalling mediated effects, precise definitions have remained difficult as effects are often dependent on the experimental system or exposure conditions being measured. The caveats associated with these different effects observed under different experimental and exposure conditions were the subject of recent review by Blyth and Sykes (9) who stated that ‘most reports in the literature are accompanied by the authors own definition, usually framed in the context of the data presented in that report’. This is an important consideration which the authors addressed by attempting to establish a general framework for the classification radiation induced signalling effects based on human radiation exposure scenarios. They define three different categories; bystander effects, cohort effects and abscopal effects.

The most well-established of these classes are abscopal effects. These are defined as radiation induced effects in unirradiated tissues occurring outside of an irradiated volume. Radiation induced abscopal effects were observed more than 60 years ago in some patients following radiotherapy and do not appear to be dose dependent dose making them particularly relevant to the partial body exposures typically delivered during conformal radiotherapy. Abscopal effects are rarely recognised in the clinic and so their importance in radiotherapy response remains controversial (10).

More recently, bystander effects have been defined for human exposure scenarios as radiation induced, signal mediated effects in unirradiated cells within an irradiated volume exposed to a sufficiently low dose that a portion of cells within the exposed volume remain unexposed and survive. These effects are relevant for whole and partial body exposures to very low doses, such as those from background radiation, high altitude flights and ingested radioactive potassium.

A third classification of effects, termed cohort effects describe the component of overall radiobiological response in irradiated cells which is not a consequence of direct energy deposition in
the target cell but rather due to communication between cells within an irradiated volume. Cohort effects are relevant for any exposures where the majority of a cell population is exposed to significant dose and whilst this interpretation is relatively uncommon in the literature, there is increasing evidence that intercellular signalling plays a role in overall radiation response.

These classifications of radiation induced signalling are shown schematically in figure 1. Distinction between the classifications is difficult in modern radiotherapy as patients are exposed to complex field geometries with steep dose gradients resulting in delivery of differential doses to cells in close proximity which can freely signal to one another. Whilst these effects are typically classified and studied separately, they have many common characteristics in that they all occur in response to radiation exposure, are mediated by extracellular signalling factors and cause negative radiobiological effects in neighbouring cells.

Consequently, many of these experimentally and clinically observed phenomena which are often classified as different effects actually stem from the same or similar cellular signalling origin and may be interpreted as different consequences of the same generalised RIBEs. As a result, there is significant opportunity for novel approaches to investigate these effects in more clinically relevant scenarios.

**Classical experimental approaches for studying RIBEs**

A diverse range of experimental approaches have been used to investigate RIBEs at the single cell, multicellular and whole organism levels. A selection of these studies in single cells is summarised in Table 1 and the various approaches used shown schematically in figure 2. Classical *in vitro* studies used polonium needles (11) or low fluence α-particle exposures (12) to target a small number of cells within a population. Technological developments have driven more sophisticated approaches using radiation microbeams allowing the delivery of highly focussed low energy micron sized radiation beams to single cells or subcellular targets. They have been successfully used as mechanistic probes to investigate biological processes including kinetics of DNA damage repair and subcellular
signalling processes involved in RIBEs (13). Microbeam approaches have utilised not only ion beams, including protons and helium ions, but focussed soft X-ray microbeams and electron microbeams.

In addition to microbeam studies, several cell culture methods have been applied to study RIBEs. These have involved the transfer of culture medium from irradiated cells onto unirradiated recipient cells known as media transfer (14,15) or co-culturing methods where irradiated and unirradiated cell populations are physically separated but free to signal to one another (6). These techniques have been used to demonstrate RIBEs manifested in a range of biological endpoints including DNA damage (4,16), cellular transformation (17), changes in gene expression (18), chromosomal aberrations (3,19,20) and mutations (21).

Whilst these early experimental approaches have provided significant understanding of the signalling mechanisms and kinetics of RIBEs, in most cases, they do not accurately represent the physiological multicellular environment or radiation dose distributions delivered during clinical treatment protocols. This has led to the development of new approaches to investigate RIBEs under more clinically relevant exposure conditions and at the multicellular and whole organism levels.

**New experimental approaches for studying RIBEs**

Early approaches to investigate RIBEs fail to accurately replicate clinically relevant exposure scenarios in respect to beam energy, delivery time and dose distributions. This has recently been addressed in several studies which have delivered modulated 6 MV radiation fields using clinical linear accelerators to more accurately replicate exposure conditions in vivo (7,23,24). Work from our laboratory determined cell survival responses occurring in- and out-of-field using a modulated beam profile generated using a multi leaf collimator (MLC) and demonstrated significantly reduced out-of-field cell survival following irradiation compared to the level of response predicted on scattered dose alone (7). In addition, the observed out-of-field responses were shown to be dependent on cell signalling between the differentially irradiated cell populations in part mediated by nitric oxide. Whilst these studies have provided evidence for RIBEs in response to more clinically relevant
radiation exposures at high dose using MV energy, cell culture models are limited to two dimensions, lacking cellular architecture and physiological context.

Several multicellular tissue models summarised in table 2, have been used to investigate RIBEs ex vivo using sections of porcine ureter (26,27) and reconstructed 3D skin models (28,29). Recently, Sheridan et al, (31) observed RIBEs manifest as elevated DNA damage in nonadjacent colon tissue obtained from patients receiving post neoadjuvant radiotherapy.

At the whole organism level, RIBEs have been observed in a number of systems for different exposure scenarios (32). These are summarised in table 3. The first in vivo evidence of RIBEs was provided by the identification of cell damaging or clastogenic factors in the serum of irradiated patients which when transferred onto cultures of unirradiated lymphocytes showed cell damaging activity (33,34). As in more simplistic cell models, it is difficult to distinguish between types of radiation induced signalling in vivo. Most experimental models have involved partial body exposures and are therefore classified as abscopal effects. These have been observed clinically for many years and were originally defined as systemic radiation effects following local radiotherapy (35).

Experimentally, abscopal effects have been demonstrated in a number of whole organisms, summarised in table 3. Using the C57BL/6 mouse, Camphausen et al, (36) showed significant reduction in the growth of tumours implanted to the dorsal midline when the legs of the animals were irradiated. In the same model, Koturbash et al, (37) showed induction of DNA damage in skin tissue up to 7 mm away from the irradiated site following partial body exposure.

An important model which has been used to demonstrate the tumourigenic potential of abscopal effects is the Patched-1 (Ptch1+/−) mouse (41,42). PTCH is a Sonic Hedgehog (SHH) receptor and negative regulator of the pathway causing predisposition to childhood medulloblastoma (43). Ptch1+/− mutant mice develop cerebellar tumours resembling human medulloblastoma which is accelerated when irradiated as neonates (44). Partial body irradiation of neonatal Ptch1+/− mice was shown to significantly increase the occurrence of medulloblastoma compared to control animals, accompanied with increased DNA damage and apoptosis in the cerebellum.
In addition to partial body exposures, other approaches have involved bone marrow transplantations in which irradiated bone marrow cells differentiated from unirradiated bone marrow cells using cytogenetic markers, are capable of inducing genetic instability in the progeny of unirradiated bone marrow cells (38).

*In vivo* demonstration of RIBEs has not been exclusively limited to mouse models. In Sprague Dawley rats, partial irradiation of the lower lung has been shown to cause increase damage in the upper out-of-field regions of the lung (45,46). Mothersill and colleagues have used several species of fish including rainbow trout (47), zebrafish (48) and medaka (49) to demonstrate RIBEs *in vivo*.

Although RIBEs have clearly been demonstrated *in vivo*, the systems in which they have been investigated do not accurately replicate typical exposure conditions during radiotherapy. Whilst next generation higher energy microbeams will provide improved subcellular targeting accompanied with advanced imaging, there is a need to determine RIBEs under conditions analogous to clinical protocols. This is potentially possible through the application of tumour bearing animals models in combination with advanced small animal radiation research platforms (52) and presents an exciting opportunity to determine the precise implications of RIBEs for radiotherapy and cancer risk following clinically relevant exposures.

**Predictive models of RIBEs**

To fully determine the implications of RIBEs in the clinical context, several attempts to describe predictive frameworks of radiation induced cell signalling have been proposed (53-61). Generally, these models tend to describe cellular responses to ionising radiation as resulting from two distinct components – a “direct” component, due to radiation interactions within a particular cell, and an “indirect” component resulting from intercellular communication, which is dependent on the exposure across the entire cellular population.

One of the primary areas for disagreement between these different models is the case where cells are subject to both direct and indirect effects, determining how they interact and which effects are dominant under which conditions. Many older models, based on media transfer or co-culture
experiments suggested that indirect components are small compared to the direct component, or even mutually exclusive with direct exposures. However, more recent work such as that of Ebert et al. (60) and McMahon et al. (61) suggest that indirect effects may play a significant role not only in out-of-field bystander cells, but also on the survival of cells within an irradiated population at high doses, reflecting the significance of intercellular communication in cohort effects, as outlined above. These results may have significant implications for the interpretation of a large portion of radiobiology data as they suggest the traditional paradigm of independent cellular responses may not be valid, with further consequences for the evaluation of many clinical conditions.

**Implications for radiotherapy and cancer risk**

From our understanding of RIBEs and increasing evidence of their role *in vivo*, it is clear that radiation induced signalling is of significant importance in responses to radiotherapy. From the clinical perspective RIBEs have a number of important potential implications particularly relating to tumour control, normal tissue response and risk of secondary cancer.

Whilst classically defined bystander effects were interpreted as a low dose phenomenon, recent findings have highlighted the potential significance of radiation induced signalling in cohort and bystander effects following modulated dose distributions suggesting these may have a role in directly irradiated regions (7,23,24). Signalling from cells irradiated to high doses within the planning target volume (PTV) may contribute to increased tumour cell killing and improved outcomes. Validation of such effects is likely to have significant impact on clinical decisions particularly at tumour margins as it is currently suggested that the optimal planning solution should incorporate a ‘biological’ margin to ensure the irradiation of some healthy tissue.

In the absence of an accurate estimate of the spatial component of radiation induced signalling, it is difficult to define the role of RIBEs in normal tissue. Potentially, RIBEs may have a negative impact in normal tissue due to the signalling extending beyond the physical dose distributions of conventional treatment plans. This would mitigate the anticipated reduction in radiation toxicity achieved with more conformal dose distributions to the tumour target.
As a result of these effects, definitions of PTV based on physical dose constraints may not accurately describe biologically effective dose mediated through cell signalling. Current tumour control probability (TCP) and normal tissue complication probability (NTCP) models could be significantly enhanced by the input of biological parameters to account for spatial dose distributions and biological response. An example of this would be in dose-painting scenarios e.g. where dose is boosted into discrete regions of the tumour based on biological activity (62).

The role of RIBEs on secondary cancer risk has been considered for some time in relation to low dose environmental exposures and out-of-field regions in radiotherapy (63, 64). Attempts to attribute additional cancer risk to these effects have remained challenging as current empirical models of secondary cancer risk do not address specific mechanisms such as in vivo signalling effects.

In addition, further understanding of the underlying molecular mechanisms mediating RIBEs may identify novel therapeutic targets enabling the exploitation of cell signalling mechanisms to improve the specificity of radiotherapy treatment by sparing healthy tissues which lie close to radiation field.

Taken together it is clear that the radiation induced signalling effects demonstrated at the single-, multi-cellular and whole organism level are likely to have important radiobiological consequences for clinical exposures. Further understanding of these effects gained through the application of new models and new technologies including precision image-guided in vivo radiobiology is likely to provide opportunities to improve the efficacy of radiotherapy through biologically optimised treatment planning and novel therapeutic targets.

Acknowledgements

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**Table 1**
Summary of experimental approaches to study RIBEs using *in vitro* single cell models

<table>
<thead>
<tr>
<th>Experimental system</th>
<th>Observed effects</th>
<th>Biological endpoint</th>
<th>Reference(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Low fluence α-particle</td>
<td>Bystander</td>
<td>Genomic instability</td>
<td>(3)</td>
</tr>
<tr>
<td>Radiation microbeam</td>
<td></td>
<td>Clonogenic survival</td>
<td>(22)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Cell survival</td>
<td>(17)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Transformation</td>
<td></td>
</tr>
<tr>
<td>Media transfer</td>
<td></td>
<td>Clonogenic survival</td>
<td>(5, 14, 15)</td>
</tr>
<tr>
<td>Co-culture</td>
<td></td>
<td>Micronuclei</td>
<td>(6)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>DNA damage</td>
<td></td>
</tr>
<tr>
<td>Modulated field</td>
<td>Bystander &amp;</td>
<td>Clonogenic survival</td>
<td>(7, 23, 24)</td>
</tr>
<tr>
<td></td>
<td>cohort</td>
<td>DNA damage</td>
<td>(25)</td>
</tr>
</tbody>
</table>

**Table 2**
Summary of experimental approaches to study RIBEs using *ex vivo* tissue models

<table>
<thead>
<tr>
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<th>Observed effects</th>
<th>Biological endpoint</th>
<th>Reference(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Urothelial explant model</td>
<td>Bystander</td>
<td>Differentiation</td>
<td>(26, 27)</td>
</tr>
<tr>
<td>3D skin model</td>
<td></td>
<td>MN, apoptosis</td>
<td>(28, 29)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>DSBs</td>
<td></td>
</tr>
<tr>
<td>Fish explant model</td>
<td></td>
<td>Clonogenic survival</td>
<td>(30)</td>
</tr>
<tr>
<td>Human colon explant</td>
<td></td>
<td>DSBs</td>
<td>(31)</td>
</tr>
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</table>

Table 3
## Summary of experimental approaches to study RIBEs using *in vivo* models

<table>
<thead>
<tr>
<th>Organism</th>
<th>Model system</th>
<th>Observed effects</th>
<th>Biological endpoint</th>
<th>Reference(s)</th>
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</thead>
<tbody>
<tr>
<td>Mouse</td>
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<td>Bystander</td>
<td>Genomic instability</td>
<td>(38)</td>
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<td></td>
<td>NCr nude mice</td>
<td>Bystander</td>
<td>Tumour volume</td>
<td>(39)</td>
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<tr>
<td></td>
<td>C57BL/6</td>
<td>Abscopal</td>
<td>Tumour growth</td>
<td>(36)</td>
</tr>
<tr>
<td></td>
<td>C57BL/6 &amp;</td>
<td>Abscopal</td>
<td>DNA damage</td>
<td>(37)</td>
</tr>
<tr>
<td></td>
<td>BALB/c mice</td>
<td></td>
<td>Proliferation</td>
<td></td>
</tr>
<tr>
<td></td>
<td>C57BL/6</td>
<td>Bystander</td>
<td>DNA methylation</td>
<td>(40)</td>
</tr>
<tr>
<td></td>
<td><em>Ptch1</em>^-/-^ mouse</td>
<td>Abscopal</td>
<td>Tumourigenesis</td>
<td>(41, 42)</td>
</tr>
<tr>
<td>Fish</td>
<td>Rainbow trout</td>
<td>Bystander</td>
<td>Reporter assay</td>
<td>(47)</td>
</tr>
<tr>
<td></td>
<td>Zebrafish</td>
<td>Bystander</td>
<td>Reporter assay</td>
<td>(48)</td>
</tr>
<tr>
<td></td>
<td>Medaka</td>
<td>Bystander</td>
<td>Apoptosis</td>
<td>(49)</td>
</tr>
<tr>
<td>Other</td>
<td>Rat</td>
<td>Abscopal</td>
<td>DNA damage</td>
<td>(45, 46)</td>
</tr>
<tr>
<td></td>
<td><em>C. elegans</em></td>
<td>Abscopal</td>
<td>hsp GFP reporter</td>
<td>(50)</td>
</tr>
<tr>
<td></td>
<td><em>A. thaliana</em></td>
<td>Abscopal</td>
<td>DNA damage</td>
<td>(51)</td>
</tr>
</tbody>
</table>
Figure 1
Schematic representation of radiation induced signalling effects classified by Blyth and Sykes (8). Irradiated cells are shown in red; unirradiated cells in blue. Bystander effects occur in unirradiated cells within an irradiated volume; within the same volume, radiation induced signalling contributes to the overall response through cohort effects. Abscopal effects occur in unirradiated tissue at a distant site outside of an irradiated volume.
Figure 2
Schematic representation of experimental approaches for investigating radiation induced signalling effects in vitro using single cell models. a) Radiation microbeams are used to target single cells or subcellular components within cell populations. b) Irradiated cell conditioned medium is transferred to unirradiated recipient cells from irradiated donor cells. c) Irradiated cells are co-cultured with unirradiated using a cell insert system. d) Modulated fields are created using a multi-leaf collimator (MLC) to place cells out-of-field.