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OXFORD

Radiation-induced unrepairable DSBs: their role in the late effects of radiation and possible applications to biodosimetry

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ABSTRACT

Although the vast majority of DNA damage induced by radiation exposure disappears rapidly, some lesions remain in the cell nucleus in very small quantities for days to months. These lesions may cause a considerable threat to an organism and include certain types of DNA double-strand breaks (DSBs) called 'unrepairable DSBs'. Unrepairable DSBs are thought to cause persistent malfunctioning of cells and tissues or cause late effects of radiation, especially the induction of delayed cell death, mutation, senescence, or carcinogenesis. Moreover, the measurement of unrepairable DSBs could potentially be used for retrospective biodosimetry or for identifying individuals at greater risk for developing the adverse effects associated with radiotherapy or chemotherapy. This review summarizes the concept of unrepairable DSBs in the context of persistent repair foci formed at DSBs.

Keywords: radiation; unrepairable DSBs; gamma H2AX; senescence; biodosimetry

INTRODUCTION

Among the genome damage induced by ionizing radiation, DNA double-strand breaks (DSBs) represent the most biologically deleterious type of lesion. To tackle these potentially lethally damaging lesions, cells have evolved orchestrated and conserved mechanisms known collectively as the DNA damage response (DDR) [1-4]; the DDR coordinates cellular DSB repair activities immediately after the damage is detected [4]. Indeed, within seconds, DSB repair proteins start to accumulate at the site of DSBs, and the DDR directs the cells to repair the breaks, undergo apoptosis, or become senescent [3]. DSB repair is a quick and efficient process whereby broken ends are rejoined. Using traditional and biochemical DNA size analyses [5-7] coupled with immunocytochemical staining [8-10], the kinetics of DSB repair have been shown to occur in two phases: a fast phase lasting up to a few hours, with a half-life of 30 min to 1 h, followed by a slower phase that may persist for a long time. A few persistent DSBs remain into the next day, and some of these DSBs persist for days, months, or even years [10-13]. These DSBs are more difficult to repair or remain unrepaired DSBs. Alternatively, these persistent DSBs are termed residual DSBs and are retained in the damaged cells unless the cells die and slip away [10, 14]. The number of unrepairable DSBs in a cell is measured by detecting and counting the number of persistent repair foci, which are composed of multiple proteins that accumulate at DSBs. Specifically, it is suggested that individual γ H2AX or 53BP1 foci represent a single DSB with a ratio of 1:1 [8, 15, 16]; hence, very few repair foci out of the many that form immediately after radiation exposure remain and become persistent in the cell nucleus. In this review, I summarize the observations obtained from the exposure of quiescent normal human cells to radiation, incorporate these observations into a discussion of the literature, and then discuss the biological implications of unrepairable DSBs with respect to the effects of radiation on cells and tissues. I also discuss the possible application of unrepairable DSB measurement for retrospective biodosimetry.

GENERATION OF UNREPAIRABLE DSBS AND PERSISTENT REPAIR FOCI

Radiation-induced DSBs are first detected by the MRE11/Rad50/ NBS1 (MRN) complex and the Ku70/Ku80 heterodimer [1, 3]. Accordingly, the DDR initiates many possible mediators and effectors to potentiate distinct signals within a cell. The MRN complex promotes the binding of a phosphorylated form of ATM and the phosphorylation of H2AX. The MRN complex also promotes the

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binding of MDC1 and a group of ubiquitination proteins (RNF8 and RNF168) and their effectors at the DSBs, thereby forming the fundamental structures of repair foci. Repair foci are a cluster of many kinds of proteins. They have different components depending on whether the cells are undergoing the NHEJ pathway (including microhomology-mediated NHEJ) or entering S phase, during which the cells undergo HR-mediated repair. In fact, in exponentially growing normal human fibroblasts, 100% of cell nuclei have many γ H2AX/53BP1 foci after radiation exposure, whereas only 30% of those exhibit BRCA1-positive foci. The repair foci containing BRCA1 are specifically formed for recombination repair in S phase (Fig. 1). This observation indicates that the accumulation of BRCA1 protein depends on cell progression through S phase. In somatic cells in living organisms, most of the cells are in quiescent (Go) phase; they are not dividing and are instead terminally differentiated, functioning to preserve their specified phenotype and exhibiting very long lifespans, up to years or decades. To create a model system for this in vivo cell status, we have adopted quiescent cultures of normal human diploid fibroblasts (NHDFs) in which the cells are maintained in MEM + 0.1% FCS. In this condition, the cells are permanently growth arrested (Go state) and can survive as long as weekly medium changes are maintained (any time after quiescence, cells can reenter the cell cycle in the presence of medium containing 10% FCS, which provides growth stimulation).

Analysis of the DSB repair kinetics of guiescent NHDFs showed that 95% of the initial DSBs disappeared within a couple of days after radiation exposure, followed by a very slow decline to ~1% DSBs remaining after 2 weeks. These DSBs persisted in the cell nucleus as long as the culture continued (up to 1 year in our study) [10], thus we designated them 'unrepaired DSBs'. The formation and resolution of yH2AX/53BP1 foci have been extensively studied in vitro [8, 17-20], and the kinetics have been shown to be biphasic [10-12, 21], which matches well with the kinetics of DSB repair [5]. Although the persistence of repair foci could arguably reflect insufficient H2AX dephosphorylation [22], it has been readily accepted that the foci represent unrepaired DSBs because the foci also include 53BP1 and all other previously reported DDR proteins. Moreover, the phosphorylated form of DNA-PKcs (ser-2056), which marks the initial recognition of a DSB, was also detected in the foci [10]. Treatment of cells with the ATM inhibitor KU55933 completely abolished the persistent foci, and the foci recovered at the DSB sites when the inhibitor was chased off. This effect was observed in cultures even 6 months post exposure to radiation. Treatment with the polyubiquitination inhibitor MG132 also produced the same effects. These results indicate that



Fig. 1. Staining of exponentially growing NHDFs immediately after radiation exposure (1 Gy/1 h) with antibodies against 53BP1 and BRCA1.

there is a continuous turnover of foci components at the unrepaired DSBs, which persists for a long time after the exposure.

Therefore, what is the function of the slow to very slow type of DSB repair that follows the fast phase? Is it truly a process of very slow repair, does this process establish unrepaired DSBs, or is it a mixture of both processes? At present, we do not know the biochemical difference between repairable and unrepairable DSB foci; however, it is almost certain that the size of the foci grows larger over time [10, 23–26]. We measured the foci enlargement over time and found that the unrepaired foci could be sorted from the small foci of the fast phase that would eventually disappear upon the completion of repair. The repairable foci displayed a normal size distribution immediately after radiation exposure, whereas after 1 or 2 months the unrepaired DSB foci formed a distinct distribution of foci larger in diameter and could be separately counted (Fig. 2).

PROSPECTIVE APPLICATIONS OF UNREPAIRABLE DSBS TO BIODOSIMETRY

We know that the effects of radiation stem from unrepairable damage and not from the repair process of repairable damage. It has been argued that the radiosensitivity of an individual cell can be attributed to the presence of unrepairable DSBs [27–29]. These unrepairable DSBs are responsible for the late effects of radiation, i.e. changes in the aging process, late onset mutations, and cancer. It is also argued that the measurement of unrepairable DSBs could be applied to radiation dosimetry on previously irradiated cells and tissues [30–33].

By counting the number of large-sized foci remaining 1 month after irradiation, we found that there was a linear relationship between the radiation dose and the number of unrepaired DSBs. Assuming that one unrepairable DSB is one lethal hit, then the actual average one-hit dose is 'D_o', which is the dose required for 37% cell survival in classic 'radiation hit theory'. A similar idea enables the calculation of the average two-hit dose required for 13.5% cell survival. Then, from the dose–response curves of the formation of unrepairable DSBs we made hypothetical survival



Fig. 2. Images of the size distribution of repair foci immediately after or a long time after radiation exposure. One month after exposure, transient/repairable DSB foci disappeared, and persistent DSB foci remained in the nucleus.

curves, which agreed well with actual cell survival phenotypes. The concept that unrepairable DSBs limit the radiosensitivity of cells has been validated by other studies [28, 29]. In fact, Menegakis *et al.* have also attempted to calculate hypothetical cell survival [34].

Based on observations that the number of unrepairable DSBs accumulates with repeated exposure [10, 20], retrospective radiation biodosimetry has been proposed [10, 20, 35, 36]. By measuring the number of unrepaired DSB foci per unit area of mouse skin, Bhogal et al. reported the possibility of biodosimetry [28]. In minipig, skin biopsies taken 70 days after a 50 Gy exposure showed typical unrepaired DSB foci patterns. Accordingly, minipig [37] and macaque models [38] for persistent DSB-mediated biodosimetry have been reported. Cells from plucked hair appeared to be especially useful [32]. In mouse spinal cord, the persistent foci were detectable 1 year after the exposure [39]. Overall, these reports may offer a new avenue for the use of unrepairable DSBs in radiation biodosimetry. We also observed the appearance of large repair foci one month after mouse pancreas was irradiated with 6 Gy. Similar observations have been reported in human buccal cells [40] and mouse germ cells [39, 41]. Moreover, increases in the number of unrepairable DSBs have been associated with aging in many studies [20, 36, 42–44].

Some pediatric cancer patients display increased unrepairable DSB production levels in their peripheral blood lymphocytes. These patients might represent groups at higher risk during radiation exposure [45]. A group of patients who demonstrated an excessive response to radio-therapy, which was assessed by normal tissue toxicity, exhibited increased levels of persistent repair foci [46]. The levels of persistent repair foci per cell in lymphocytes were examined after radiotherapy in breast cancer patients [29, 47]. This calculation will enable the fine-tuning of radiation doses for improved cancer treatments. In head and neck cancer cases, prescreening the levels of persistent foci in 2 Gy irradiated lymphocytes was applied to reduce the side effects of radio-therapy [48]. Thus, persistent repair foci are a useful measure for the detection and evaluation of previously irradiated cells.

However, it has been reported that repair kinetics and the formation of persistent repair foci vary extensively among mouse tissues [49, 50]. Therefore, tissue differences should be considered when evaluating unrepairable DSBs for biodosimetry.

THE REAL IDENTITY OF UNREPAIRABLE DSBS: ARE THEY TELOMERE DSBS OR ARE THEY LOCATED INSIDE THE CHROMOSOME?

Although the chemical structures of unrepairable DSBs have not yet been precisely determined, two types of 'complex' lesions can be postulated: those with non-ligatable termini due to crosslinks between bases and sugars (dirty DSBs) and those with damaged sites in which multiple DSBs/SSBs and/or base damage arise in close proximity (clustered damage) [51]. The complexity of the damage structure is evident by the slow rejoining kinetics of DSBs induced by high-LET radiation [52]. It is apparent that the damage induced by radiation is extremely heterogeneous and that damage occurs randomly throughout the chromosome.

Unrepaired DSBs induce permanent cell growth arrest or death. This fact is especially true in non-apoptotic terminally differentiated cells, which in general have very long lifespans. In these cells, cellular senescence is induced (radiation-induced senescence). It has been shown that many adult survivors of juvenile cancer develop symptoms of premature aging later in life [53], which may be related to the unrepaired DSBs that were induced after radiotherapy in normal tissues adjacent to the tumor. Their symptoms include advanced frailty with increased risk for heart failure, severe cognitive decline, coronary heart disease, and secondary neoplasms [53–55]. Meanwhile, neuronal stem cells bearing unrepairable DSBs undergo premature senescence or terminal differentiation to become astroglial cells [56]. This process also reduces stem cell number, leading to a premature aging of neuronal systems. Likewise, after radiation exposure, melanocyte stem cells in hair bulge undergo terminal differentiation in their niches, producing gray hairs [57].

Telomere-driven replicative senescence has been widely accepted as a primary mechanism of organismal aging and cellular senescence. Under this theory, eroded chromosome ends continuously activate the DDR, thereby permanently inducing growth arrest and senescence. Similarly, it has been hypothesized that radiation-induced unrepairable DSBs eventually accumulate in chromosome ends, which establishes senescence independent of the cell's telomere length [14, 35, 36, 58]. This idea is based on the specific structure of chromosome ends, known as Shelterin, which protect chromosomes from improper recombination and degradation. Indeed, TRF2 and RAP1, the components of Shelterin, were shown to inhibit DSB repair throughout the entire telomere [35, 59-61], not only at the very ends. Therefore, radiation-induced DSBs near the chromosome ends would be preferentially protected from repair systems, even though radiation exposure induces DSBs randomly throughout the chromosome. Such a mechanism may eventually provoke continuous DDRs, leading to cellular senescence in cells bearing radiation-induced unrepairable DSBs (Fig. 3A).



Fig. 3. (A) Possible mechanisms for the generation of unrepaired DSBs. Although radiation exposure induces DSBs randomly in the chromosome, those occurring proximal to the telomere or unrepairable DSBs inside the chromosome eventually become persistent. (B) Persistent repair foci tend to make pairs. In the lower photo, two pairs of foci of different sizes can be seen.

This idea has been confirmed via confocal microscope observations and telomere ChIP experiments [36]. Thus, the telomere hypothesis suggests that the location, not the chemical structure, is important for the establishment of unrepairable DSBs. This hypothesis may explain the radiation-induced senescence of young cells *in vitro* and nondividing quiescent cells *in vivo*. Both of these cell types have long telomeres and no chance of undergoing replicative senescence. Assuming that animals carrying longer telomere sequences have a wider target for the creation of unrepairable DSBs, telomere length may have a negative effect on lifespan. Indeed, inverse correlations between telomere length and lifespan have been reported in different animal species [62].

However, other mechanisms of radiation-induced senescence may exist. Most nuclei of 6 Gy-irradiated and 1-month-cultured young quiescent cells carried a few typical, large yH2AX/53BP1 foci, as mentioned above. Careful observations have revealed that these foci often appear in pairs (i.e. even numbers per nucleus), as if they had originated from a single DSB and then distantly separated. Taking a closer look, each partner of the foci pairs can be identified because each foci had a distinct size (Fig. 3B and reference photos provided in the supplemental data) [10]. In this case, the DSBs should have originated from the inside of the chromosome and not from the very end. Data in reports from Hewitt [36] and Fumagalli [35] indicates that, at most, 50% of unrepaired DSBs are located in telomeres; therefore, it is reasonable to assume that the causative mechanisms of unrepairable DSBs can be attributed to both their locations and chemical structures [63]. Collectively, while the fast phase of DSB repair constitutes rejoining of easy-to-repair 'clean DSBs', the slower phase might be composed of two distinct components, namely, telomere-silenced DSBs and unrepairable DSBs.

In yeast, several studies have indicated that persistent DNA lesions relocate to either the nuclear pore complex (NPC) or the nuclear envelope (NE) [64-66]. In such cases, specific membrane structures, including the components of nuclear pore proteins, appear to be crucial for the repair of persistent DSBs and eroded telomeres. In mammals, mutated forms of nuclear lamin A, referred to as progerin, induce deformation of the NE and impair certain processes of DSB repair. This process has been argued to contribute to the decrease in repair efficiency [67, 68] or to the enhanced production of unrepairable DSBs [69, 70]. In either case the expression of progerin has an adverse effect on DSB repair and induces premature senescence in cells. Likewise, NHDFs (normal cells) bearing radiation-induced unrepairable DSBs, which undergo premature senescence, show dysfunctional nuclear membrane structures [70]. These results indicate a possible link between unrepairable DSBs and premature senescence, which are both mediated by a dysfunctional nuclear membrane. The nuclear membrane and its periphery has been shown to associate with heterochromatin, where 'gene deserts' have also been shown to localize [71, 72]. The DSB repair of such regions is slow and dependent on both ATM and Artemis [51]. Overall, these data suggest that the final destination of unrepairable DSBs is the nuclear membrane. However, the nuclear periphery is also a place where telomeres locate [73] and is where telomere silencing occurs. Therefore, it is reasonable to speculate that both types of unrepairable DSBs, which primarily originated in either the telomere or heterochromatin, eventually colocalize in the nuclear membrane. New technologies [63, 74] and biomarkers will

be necessary to distinguish radiation-induced unrepairable DSBs from those found at telomere ends. A further application of unrepaired DSBs measurement would be its use as an early indicator for radiation risk.

New methods that artificially introduce unrepairable DSBs at specific chromatin sites have been developed. White *et al.* [75] delivered Sac I restriction enzyme to mouse liver using adenovirus. They observed a liver-specific pathology of aging and inflammation. However, lipofuscin accumulation was not observed. Kim *et al.* [76] generated a conditional I-PpoI restriction enzyme expression system in mouse. In this system, 19 persistent DSBs could be formed in each cell, and the mice appeared to exhibit premature aging phenotypes. However, the introduction of such unrepairable DSBs could not fully explain all of the normal aging phenomena. The application of recent gene editing technologies will enable the introduction of unrepairable DSBs at specific sites in the chromosome. Such systems will help us to understand the risks of unrepairable DSBs in specific cells and tissues in living organisms.

CONCLUSION: A VIEW OF RADIATION BIOLOGY USING A NEW BIOMARKER

Although considerable efforts have been made to analyze the repair of repairable damage, studies that measure and elucidate the biology of unrepairable DSBs have not come to the forefront until recently. To better measure unrepairable DSBs, especially *in vivo*, we need definitive criteria for distinguishing unrepairable DSBs from transient and repairable DSBs. These criteria may include characterizing the precise sequences and structures where these breaks originated or new biomarkers that can specifically detect and measure these lesions. Such new technologies may enable the further application of radiation-induced unrepairable DSB quantification, which could lead to a better understanding of the risks of radiation to the processes of organismal development, growth, and aging.

SUPPLEMENTARY DATA

Supplementary data are available at the *Journal of Radiation Research* online.

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CONFLICT OF INTEREST

The author has no conflict of interest to disclose.

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REFERENCES

- 1. Jackson SP, Bartek J. The DNA-damage response in human biology and disease. *Nature* 2009;461:1071–8.
- 2. Ciccia A, Elledge SJ. The DNA damage response: making it safe to play with knives. *Mol Cell* 2010;40:179–204.
- Polo SE, Jackson SP. Dynamics of DNA damage response proteins at DNA breaks: a focus on protein modifications. *Genes Dev* 2011;25:409–33.
- Ceccaldi R, Rondinelli B, D'Andrea AD. Repair pathway choices and consequences at the double-strand break. *Trends Cell Biol* 2016;26:52–64.
- Bradley MO, Kohn KW. X-ray induced DNA double strand break production and repair in mammalian cells as measured by neutral filter elution. *Nucleic Acids Res* 1979;7:793–804.
- Metzger L, Iliakis G. Kinetics of DNA double-strand break repair throughout the cell cycle as assayed by pulsed field gel electrophoresis in CHO cells. *Int J Radiat Biol* 1991;59:1325–39.
- Frankenberg-Schwager M, Frankenberg D. Survival curves with shoulders: damage interaction, unsaturated but dose-dependent rejoining kinetics or inducible repair of DNA double-strand breaks? *Radiat Res* 1994;138:S97–100.
- Rogakou EP, Pilch DR, Orr AH et al. DNA double-stranded breaks induce histone H2AX phosphorylation on serine 139. *J Biol Chem* 1998;273:5858–68.
- Nazarov IB, Smirnova AN, Krutilina RI et al. Dephosphorylation of histone gamma-H2AX during repair of DNA doublestrand breaks in mammalian cells and its inhibition by calyculin A. *Radiat Res* 2003;160:309–17.
- Noda A, Hirai Y, Hamasaki K et al. Unrepairable DNA doublestrand breaks that are generated by ionising radiation determine the fate of normal human cells. J Cell Sci 2012;125:5280–7.
- Riballo E, Kuhne M, Rief N et al. A pathway of double-strand break rejoining dependent upon ATM, Artemis, and proteins locating to gamma-H2AX foci. *Mol Cell* 2004;16:715–24.
- Lobrich M, Shibata A, Beucher A et al. γH2AX foci analysis for monitoring DNA double-strand break repair: strengths, limitations and optimization. *Cell Cycle* 2010;9:662–9.
- Sharma PM, Ponnaiya B, Taveras M et al. High throughput measurement of γH2AX DSB repair kinetics in a healthy human population. *PLoS One* 2015;10:e0121083.
- Rossiello F, Herbig U, Longhese MP et al. Irreparable telomeric DNA damage and persistent DDR signalling as a shared causative mechanism of cellular senescence and ageing. *Curr Opin Genet Dev* 2014;26:89–95.
- Rogakou EP, Boon C, Redon C et al. Megabase chromatin domains involved in DNA double-strand breaks *in vivo*. J Cell Biol 1999;146:905–16.
- Sedelnikova OA, Rogakou EP, Panyutin IG et al. Quantitative detection of ¹²⁵IdU-induced DNA double-strand breaks with gamma-H2AX antibody. *Radiat Res* 2002;158:486–92.

- 17. Stiff T, O'Driscoll M, Rief N et al. ATM and DNA-PK function redundantly to phosphorylate H2AX after exposure to ionizing radiation. *Cancer Res* 2004;64:2390–6.
- Madigan JP, Chotkowski HL, Glaser RL. DNA double-strand break-induced phosphorylation of *Drosophila* histone variant H2Av helps prevent radiation-induced apoptosis. *Nucleic Acids Res* 2002;30:3698–705.
- Roch-Lefevre S, Mandina T, Voisin P et al. Quantification of gamma-H2AX foci in human lymphocytes: a method for biological dosimetry after ionizing radiation exposure. *Radiat Res* 2010;174:185–94.
- Sedelnikova OA, Horikawa I, Zimonjic DB et al. Senescing human cells and ageing mice accumulate DNA lesions with unrepairable double-strand breaks. *Nat Cell Biol* 2004;6:168–70.
- 21. Jeggo PA, Geuting V, Lobrich M. The role of homologous recombination in radiation-induced double-strand break repair. *Radiother Oncol* 2011;101:7–12.
- 22. Mamouni K, Cristini A, Guirouilh-Barbat J et al. RhoB promotes gammaH2AX dephosphorylation and DNA doublestrand break repair. *Mol Cell Biol* 2014;34:3144–55.
- 23. Yamauchi M, Oka Y, Yamamoto M et al. Growth of persistent foci of DNA damage checkpoint factors is essential for amplification of G1 checkpoint signaling. DNA Repair 2008; 7:405–17.
- 24. Paris L, Cordelli E, Eleuteri P et al. Kinetics of gamma-H2AX induction and removal in bone marrow and testicular cells of mice after X-ray irradiation. *Mutagenesis* 2011;26:563–72.
- Markova E, Schultz N, Belyaev IY. Kinetics and dose-response of residual 53BP1/gamma-H2AX foci: co-localization, relationship with DSB repair and clonogenic survival. *Int J Radiat Biol* 2007;83:319–29.
- 26. Bracalente C, Ibanez IL, Molinari B et al. Induction and persistence of large gammaH2AX foci by high linear energy transfer radiation in DNA-dependent protein kinase-deficient cells. *Int J Radiat Oncol Biol Phys* 2013;87:785–94.
- Banath JP, Klokov D, MacPhail SH et al. Residual gammaH2AX foci as an indication of lethal DNA lesions. BMC Cancer 2010;10:4.
- Bhogal N, Kaspler P, Jalali F et al. Late residual gamma-H2AX foci in murine skin are dose responsive and predict radiosensitivity *in vivo*. *Radiat Res* 2010;173:1–9.
- Djuzenova CS, Elsner I, Katzer A et al. Radiosensitivity in breast cancer assessed by the histone gamma-H2AX and 53BP1 foci. *Radiat Oncol* 2013;8:98.
- Taneja N, Davis M, Choy JS et al. Histone H2AX phosphorylation as a predictor of radiosensitivity and target for radiotherapy. J Biol Chem 2004;279:2273–80.
- Dikomey E, Dahm-Daphi J, Brammer I et al. Correlation between cellular radiosensitivity and non-repaired double-strand breaks studied in nine mammalian cell lines. *Int J Radiat Biol* 1998;73:269–78.
- Redon CE, Nakamura AJ, Gouliaeva K et al. The use of gamma-H2AX as a biodosimeter for total-body radiation exposure in non-human primates. *PLoS One* 2010;5:e15544.
- Qvarnstrom OF, Simonsson M, Johansson KA et al. DNA double strand break quantification in skin biopsies. *Radiother Oncol* 2004;72:311–7.

- 34. Menegakis A, Yaromina A, Eicheler W et al. Prediction of clonogenic cell survival curves based on the number of residual DNA double strand breaks measured by gammaH2AX staining. *Int J Radiat Biol* 2009;85:1032–41.
- 35. Fumagalli M, Rossiello F, Clerici M et al. Telomeric DNA damage is irreparable and causes persistent DNA-damage-response activation. *Nat Cell Biol* 2012;14:355–65.
- 36. Hewitt G, Jurk D, Marques FD et al. Telomeres are favoured targets of a persistent DNA damage response in ageing and stress-induced senescence. *Nat Commun* 2012;3:708.
- 37. Ahmed EA, Agay D, Schrock G et al. Persistent DNA damage after high dose *in vivo* gamma exposure of minipig skin. *PLoS One* 2012;7:e39521.
- Moroni M, Maeda D, Whitnall MH et al. Redon CE, Evaluation of the gamma-H2AX assay for radiation biodosimetry in a swine model. *Int J Mol Sci* 2013;14:14119–35.
- 39. Andratschke N, Blau T, Schill S et al. Late residual gamma-H2AX foci in murine spinal cord might facilitate development of response-modifying strategies: a research hypothesis. *Anticancer Res* 2011;31:561–4.
- Siddiqui MS, Francois M, Fenech MF et al. gammaH2AX responses in human buccal cells exposed to ionizing radiation. *Cytometry A* 2015;87:296–308.
- Ahmed EA, van der Vaart A, Barten A et al. Differences in DNA double strand breaks repair in male germ cell types: lessons learned from a differential expression of Mdc1 and 53BP1. DNA Repair 2007;6:1243–54.
- Rube CE, Fricke A, Widmann TA et al. Accumulation of DNA damage in hematopoietic stem and progenitor cells during human aging. *PLoS One* 2011;6:e17487.
- 43. Titus S, Li F, Stobezki R et al. Impairment of BRCA1-related DNA double-strand break repair leads to ovarian aging in mice and humans. *Sci Transl Med* 2013;5:172ra21.
- 44. Wang C, Jurk D, Maddick M et al. DNA damage response and cellular senescence in tissues of aging mice. *Aging Cell* 2009;8:311-23.
- 45. Rube CE, Fricke A, Schneider R et al. DNA repair alterations in children with pediatric malignancies: novel opportunities to identify patients at risk for high-grade toxicities. *Int J Radiat Oncol Biol Phys* 2010;78:359–69.
- 46. Bourton EC, Plowman PN, Smith D et al. Prolonged expression of the gamma-H2AX DNA repair biomarker correlates with excess acute and chronic toxicity from radiotherapy treatment. *Int J Cancer* 2011;129:2928–34.
- 47. Chua ML, Somaiah N, A'Hern R et al. Residual DNA and chromosomal damage in *ex vivo* irradiated blood lymphocytes correlated with late normal tissue response to breast radiotherapy. *Radiother Oncol* 2011;99:362–6.
- 48. Goutham HV, Mumbrekar KD, Vadhiraja BM et al. DNA double-strand break analysis by gamma-H2AX foci: a useful method for determining the overreactors to radiation-induced acute reactions among head-and-neck cancer patients. *Int J Radiat Oncol Biol Phys* 2012;84:e607–12.
- Hudson D, Kovalchuk I, Koturbash I et al. Induction and persistence of radiation-induced DNA damage is more pronounced in young animals than in old animals. *Aging* 2011;3:609–20.

- Firsanov D, Vasilishina A, Kropotov A et al. Dynamics of gammaH2AX formation and elimination in mammalian cells after X-irradiation. *Biochimie* 2012;94:2416–22.
- 51. Woodbine L, Brunton H, Goodarzi AA et al. Endogenously induced DNA double strand breaks arise in heterochromatic DNA regions and require ataxia telangiectasia mutated and Artemis for their repair. *Nucleic Acids Res* 2011;39:6986–97.
- Schmid TE, Dollinger G, Beisker W et al. Differences in the kinetics of gamma-H2AX fluorescence decay after exposure to low and high LET radiation. *Int J Radiat Biol* 2010;86:682–91.
- Oeffinger KC, Mertens AC, Sklar CA et al. Chronic health conditions in adult survivors of childhood cancer. N Engl J Med 2006;355:1572–82.
- Inskip PD, Robison LL, Stovall M et al. Radiation dose and breast cancer risk in the childhood cancer survivor study. J Clin Oncol 2009;27:3901–7.
- Ness KK, Krull KR, Jones KE et al. Physiologic frailty as a sign of accelerated aging among adult survivors of childhood cancer: a report from the St Jude Lifetime cohort study. J Clin Oncol 2013;31:4496–503.
- Schneider L, Pellegatta S, Favaro R et al. DNA damage in mammalian neural stem cells leads to astrocytic differentiation mediated by BMP2 signaling through JAK-STAT. *Stem Cell Rep* 2013;1:123–38.
- 57. Inomata K, Aoto T, Binh NT et al. Genotoxic stress abrogates renewal of melanocyte stem cells by triggering their differentiation. *Cell* 2009;137:1088–99.
- Siddiqui MS, Francois M, Fenech MF et al. Persistent gammaH2AX: a promising molecular marker of DNA damage and aging. *Mutat Res Rev Mutat Res* 2015;766:1–19.
- Bae NS, Baumann P. A RAP1/TRF2 complex inhibits nonhomologous end-joining at human telomeric DNA ends. *Mol Cell* 2007;26:323–34.
- Sarthy J, Bae NS, Scrafford J et al. Human RAP1 inhibits nonhomologous end joining at telomeres. EMBO J 2009;28:3390–9.
- Bombarde O, Boby C, Gomez D et al. TRF2/RAP1 and DNA-PK mediate a double protection against joining at telomeric ends. *EMBO J* 2010;29:1573–84.
- 62. Gomes NM, Ryder OA, Houck ML et al. Comparative biology of mammalian telomeres: hypotheses on ancestral states and the roles of telomeres in longevity determination. *Aging Cell* 2011; 10:761–8.
- Crosetto N, Mitra A, Silva MJ et al. Nucleotide-resolution DNA double-strand break mapping by next-generation sequencing. *Nat Methods* 2013;10:361–5.
- 64. Therizols P, Fairhead C, Cabal GG et al. Telomere tethering at the nuclear periphery is essential for efficient DNA double strand break repair in subtelomeric region. J Cell Biol 2006;172:189–99.
- Nagai S, Dubrana K, Tsai-Pflugfelder M et al. Functional targeting of DNA damage to a nuclear pore-associated SUMOdependent ubiquitin ligase. *Science* 2008;322:597–602.
- Freudenreich CH, Su XA. (1 December 2016) Relocalization of DNA lesions to the nuclear pore complex. *FEMS Yeast Res*, 10. 1093/femsyr/fox095.
- 67. Liu B, Wang J, Chan KM et al. Genomic instability in laminopathy-based premature aging. *Nat Med* 2005;11:780–5.

- Constantinescu D, Csoka AB, Navara CS et al. Defective DSB repair correlates with abnormal nuclear morphology and is improved with FTI treatment in Hutchinson-Gilford progeria syndrome fibroblasts. *Exp Cell Res* 2010;316:2747–59.
- 69. Richards SA, Muter J, Ritchie P et al. The accumulation of unrepairable DNA damage in laminopathy progeria fibroblasts is caused by ROS generation and is prevented by treatment with N-acetyl cysteine. *Hum Mol Genet* 2011;20:3997–4004.
- 70. Noda A, Mishima S, Hirai Y et al. Progerin, the protein responsible for the Hutchinson-Gilford progeria syndrome, increases the unrepaired DNA damages following exposure to ionizing radiation. *Genes Environ* 2015;37:13.
- 71. Redon CE, Bonner WM. High salt and DNA double-strand breaks. *Proc Nat Acad Sci U S A* 2011;108:20281–2.

- Dmitrieva NI, Cui K, Kitchaev DA et al. DNA double-strand breaks induced by high NaCl occur predominantly in gene deserts. *Proc Nat Acad Sci U S A* 2011;108:20796–801.
- Burla R, La Torre M, Saggio I. Mammalian telomeres and their partnership with lamins. *Nucleus* 2016;7:187–202.
- 74. Canela A, Sridharan S, Sciascia N et al. DNA breaks and end resection measured genome-wide by end sequencing. *Mol Cell* 2016;63:898–911.
- 75. White RR, Milholland B, de Bruin A et al. Controlled induction of DNA double-strand breaks in the mouse liver induces features of tissue ageing. *Nat Commun* 2015;6:6790.
- 76. Kim J, Sturgill D, Tran AD et al. Controlled DNA doublestrand break induction in mice reveals post-damage transcriptome stability. *Nucleic Acids Res* 2016;44:e64.