Henry Ford Health [Henry Ford Health Scholarly Commons](https://scholarlycommons.henryford.com/)

[Radiation Oncology Articles](https://scholarlycommons.henryford.com/radiationoncology_articles) **Radiation Oncology** Articles **Radiation Oncology**

2-1-2023

Molecular targets that sensitize cancer to radiation killing: From the bench to the bedside

Sushmitha Sriramulu

Shivani Thoidingjam

Stephen L. Brown

Farzan Siddiqui

Benjamin Movsas

See next page for additional authors

Follow this and additional works at: [https://scholarlycommons.henryford.com/radiationoncology_articles](https://scholarlycommons.henryford.com/radiationoncology_articles?utm_source=scholarlycommons.henryford.com%2Fradiationoncology_articles%2F389&utm_medium=PDF&utm_campaign=PDFCoverPages)

Authors

Sushmitha Sriramulu, Shivani Thoidingjam, Stephen L. Brown, Farzan Siddiqui, Benjamin Movsas, and Shyam Nyati

Review

Contents lists available at [ScienceDirect](www.sciencedirect.com/science/journal/07533322)

Biomedicine & Pharmacotherapy

journal homepage: www.elsevier.com/locate/biopha

Molecular targets that sensitize cancer to radiation killing: From the bench to the bedside

Sushmitha Sriramulu , Shivani Thoidingjam , Stephen L. Brown , Farzan Siddiqui , Benjamin Movsas, Shyam Nyati *

Department of Radiation Oncology, Henry Ford Health, Detroit, MI 48202, USA

1. Introduction

Preservation of genomic sequence information in biological systems is crucial for the survival of life. On one hand, mutagenesis is critical for genetic variation necessary for natural selection, on the other hand it could predispose for the development of cancer and other diseases. DNA is an innately reactive molecule that is sensitive to chemical alterations from both endogenous and exogenous sources [\[1\]](#page-10-0). Endogenous DNA damage may be caused by cellular factors such as hydrolysis, oxidation, alkylation or reactive chemical species (e.g., reactive oxygen species, ROS, and reactive nitrogen species, RNS) that are generated during physiochemical reactions. Exogenous DNA damage are caused by environmental (ionizing radiation, UV radiation), physical or chemical agents [\[2\].](#page-10-0) DNA lesions, such as single- and double-strand breaks, mismatches, chemical modifications of the bases, and inter- or intra-strand cross-links can be caused by ionizing radiation, UV, and chemotherapeutic agents. [Fig. 1](#page-3-0) shows the most common sources of DNA damage. A variety of clinical abnormalities, including dementia, infertility, immunodeficiencies, and cancer susceptibility, are, at least in part, caused by cells' inability to properly repair DNA damage [\[3\].](#page-10-0) Additionally, the chance of developing cancer is increased by the development of DNA

damage in cells upon exposure to carcinogens. Genomic instability and mutation, one of the hallmarks of cancer will result if DNA damage is not repaired [\[4\].](#page-10-0)

1.1. DNA damage response mechanism

The DNA damage response (DDR) is an evolutionarily-conserved defensive mechanism against various DNA lesions that may recognize, signal, and repair the damage through a series of enzymatic activities controlled by multiple proteins [\[3,5\]](#page-10-0). Proteins that are involved in the DDR identify and process different forms of DNA damage [\[5\].](#page-10-0) A majority of cancer cells have DDR pathway defects that promote malignant growth and increase the chance of tumor cell survival through natural selection [\[3\].](#page-10-0) Cancer cells become more reliant on alternative DDR pathways to preserve the integrity of their chromosomes when one or more DDR pathways are disrupted [\[6\].](#page-10-0) As a result, there is interest in the therapeutic development of inhibitors that target a specific subset of DDR network components; several are approved while many are undergoing clinical trials [\[7\].](#page-10-0)

Nucleus and mitochondria are the two primary organelles that contain DNA in mammalian cells. Major pathways for nuclear DNA

<https://doi.org/10.1016/j.biopha.2022.114126>

Available online 13 December 2022 Received 19 October 2022; Received in revised form 5 December 2022; Accepted 9 December 2022

^{*} Correspondence to: Department of Radiation Oncology, Henry Ford Health, 1 Ford Place, OFP-5D, Detroit, MI 48202, USA. *E-mail address:* snyati1@hfhs.org (S. Nyati).

^{0753-3322/© 2022} Published by Elsevier Masson SAS. This is an open access article under the CC BY-NC-ND license [\(http://creativecommons.org/licenses/by-nc](http://creativecommons.org/licenses/by-nc-nd/4.0/) $nd/4.0/$).

repair **(**[Fig. 2](#page-4-0)**)** include:

a) Direct reversal which rectifies damage caused by alkylating chemicals,

b) Base excision repair (BER) which targets non-bulky defective DNA bases and DNA single strand breaks (SSBs),

c) Nucleotide excision repair (NER) which reconstitutes large DNA damages that deform the DNA helix,

d) Base-base mismatch repair, insertion or deletion loop repair, and mismatch repair (MMR),

e) Recombination repair which is categorized into homologous recombination repair (HR) and non-homologous end joining (NHEJ),

f) Repair of DSBs is achieved via alternative non-homologous end joining (alt-NHEJ),

g) Translesion synthesis (TLS) which is a DNA damage tolerance process.

In contrast, the mitochondrial DNA (mtDNA) repair pathways that repair damaged DNA retain mitochondrial integrity, safeguard mtDNA from oxidative damage, and enhance survival are typically those that result in direct reversal: MMR, BER, TLS, and DSB repair [\[8\],](#page-10-0) although mutations to the mitochondrial DNA still occur.

SSBs and minor alterations in DNA are repaired via the BER [\[9\]](#page-10-0) components XRCC1, PARP-1, apurinic apyrimidinic endonuclease (APE1), and DNA ligase IIIa. The excision repair cross-complementing protein 1 (ERCC1)-dependent NER mechanism [\[10\]](#page-10-0) handles bulky DNA lesions including pyrimidine dimers induced by UV-irradiation. MMR pathway can fix base mismatches that occur as a consequence of replication errors [\[11\].](#page-10-0)

The MRE11–RAD50–NBS1 (MRN) complex identifies and binds to DSB sites. It subsequently autophosphorylates and triggers ataxia telangiectasia mutated (ATM) kinase $[12,13]$. When ATM is activated, it phosphorylates many downstream proteins [\[14\].](#page-10-0) Phosphorylation of CHK2 causes CDC25A, a protein phosphatase, to phosphorylate, causing cell cycle arrest. BRCA1 phosphorylation results in DSB repair and S phase cell cycle arrest, whereas p53 activation results in G1 phase cell cycle arrest or apoptosis. The ataxia telangiectasia and Rad3-related (ATR) kinase is triggered and recruited to the DNA damage sites during the primary response to SSBs/DNA replication fork collapse [\[15\]](#page-10-0). CHK1 [\[16\],](#page-11-0) which controls the activation of CDC25 phosphatases and consequently plays a role in the S and G2/M cell checkpoints, is phosphorylated and activated by ATR. To repair several forms of DNA damage, the DNA repair processes can function alone or in tandem.

Radiation mediated base damages and SSBs are quickly and efficiently repaired by cells. Most cytotoxic effects occur due to cells inability to repair DSBs. The two most prominent strategies to repair DSBs are NHEJ and HR [\[17\].](#page-11-0) NHEJ is an error-prone repair process that involves directly re-joining two broken ends and it is the most efficient DSB repair process. NHEJ is mediated by a group of proteins that includes the Ku70/Ku80 complex, DNA-PK catalytic subunit (DNA-PKcs), Artemis nuclease, XRCC4-like factor (XLF), DNA ligase IV, and X-ray repair cross complementing 4 (XRCC4). Binding of heterodimer Ku70/Ku80 at DSB ends [\[18\]](#page-11-0) is the initial stage of NHEJ, and it facilitates the recruitment of catalytic component DNAPKcs to create an active holoenzyme complex [\[19\]](#page-11-0). H2AX [\[20\]](#page-11-0), Artemis [\[21\],](#page-11-0) XRCC4, ligase IV complex [\[22\]](#page-11-0), and XLF [\[23\]](#page-11-0) are attracted to the DSB site and partake in its repair. NHEJ is probably much more precise than previously thought due to the flexibility of the NHEJ components, but if the DSB ends are incompatible, NHEJ-mediated repair can cause minor deletions, insertions, or indels. Given these factors, NHEJ seems to be a reliable repair mechanism that might help a cell increase its chances of surviving [\[24\].](#page-11-0) It is also the main pathway through which radiation damage is repaired and impairing this pathway may lead to radiosensitization.

HR, on the other hand, is an error-free repair strategy that uses a nondamaged complementary chromatid as a template [\[17\].](#page-11-0) HR is mediated by BRCA1 and 2, replication protein A (RPA), MRN complex, CtBP-interacting protein (CtIP), RAD51, and PALB2. The DSB site binds the MRN complex, DNA2-BLM (Bloom syndrome), CtIP, BRCA1, and exonuclease 1 (EXO1), which play a role in DNA reprocessing and the production of 3′ single-strand DNA, which is consequently encapsulated by RPA protein [\[25\].](#page-11-0) The missing sections can be synthesized by DNA polymerases. Holliday junctions, branched nucleic acid structure that impair function, are eventually resolved by cleaving and ligating neighboring ends [\[26\].](#page-11-0) Although HR is considered "error-free" repair, the process can be error prone when templates are swapped, as in repeat sequences [\[27\].](#page-11-0) Cell cycle stage, chromatin context, and availability of critical players such as Ku complex, RAD51, and 53BP1 may impact the selection between NHEJ and HR [\[28\]](#page-11-0).

1.2. Human cancers with aberrations in DDR

Enhanced autophosphorylation of ATM as well as ATM-dependent phosphorylation of CHK2 have been seen in early-stage malignancies, implying that the DDR may prevent progression to malignancy [\[29,30\]](#page-11-0). In radiation-resistant malignancies, DNA-PKcs have been found to be increased [\[31,32\]](#page-11-0). Upregulation of PARP1, BRCA1, APE1, RAD51, and ERCC1 have been observed in a variety of malignancies and have been linked to chemotherapy resistance [\[33\]](#page-11-0). In human sporadic malignancies, p53 is one of the most commonly altered genes. The reported frequency of p53 mutations vary by cancer type with more than half of all malignancies having inactivated p53 as a result of mutations, deletions, loss of heterozygosity, or reduced expression [\[34,35\]](#page-11-0). While BRCA1 and 2, as well as ATM inactivating mutations are less prevalent than p53 inactivating mutations [\[36](#page-11-0)–40], reduced expression of the MRN complex, ATM, BRCA1 and 2, CHK2, RAD51, and ERCC1 is widespread in sporadic malignancies, suggesting that DNA damage response aberration is common [\[41](#page-11-0)–49]. Interestingly, functional inactivation BRCA2 was observed in cancer cells that abnormally expressed SYCP3 [\[3\]](#page-10-0) a cancer/testis antigen that is known to regulate strand

Fig. 1. Most common sources of DNA damage. There are possibly other mechanisms for causing DNA damage but they are omitted here for simplicity.

Fig. 2. Major pathways for DNA repair (Adapted from "DNA Repair Mechanisms", by BioRender.com (2022).

invasion activities of Rad51.

1.3. Radiation-induced DDR

DDR is a complex protein network that coordinates cell cycle regulation and DNA repair. It disrupts the cell cycle, stopping DNA damage from propagating to daughter cells by allowing time for repair. DDR signaling is crucial for the initiation of apoptosis [\[50\]](#page-11-0). It is estimated that 1 Gray (Gy) of radiation results in approximately 10,000 damaged bases, 1000 SSBs, and 40 DSBs per cell [\[51,52\].](#page-11-0) Most base damages and SSBs are promptly repaired [\[53\]](#page-11-0). DSBs are the most damaging to cells despite their low frequency since even a single unrepaired DSB could result in cell death. Radiation therapy causes DSBs directly by ionizing molecules on DNA and indirectly by hydrolyzing molecules it encounters, i.e. predominantly water, to produce free radicals such as a hydroxyl free radical that interacts with and inactivates DNA indirectly. If the resultant DNA damage is not repaired (particularly DSBs), it leads to cell death during replication [\[54,55\]](#page-11-0).

1.4. Repair of radiation-induced DNA damage

As described, following exposure to ionizing radiation, multiple types of damages can occur to DNA resulting in activation of repair processes. The BER repairs damaged bases caused by oxidative stress [56–[61\]](#page-11-0). DNA glycosylases excise damaged bases during BER, resulting in apurinic (AP) sites. Following that, apurinic endonuclease 1 (APE1) cleaves these AP sites, resulting in SSBs. SSB repair is a component of the BER pathway that repairs SSBs [\[62\].](#page-11-0) Depending on the lesion type and cell cycle phase, either short- or long-patch SSB repair is utilized. PARP binds to SSB, which stimulates auto-PARylation and causes BER/SSBR proteins to be recruited. PARP-1 has been implicated as a DNA repair gene regulator in the E2F1 pathway [\[63\].](#page-11-0) While most of the radiation-mediated oxidative damage is repaired by BER, damage that occurs under hypoxic conditions is repaired through NER [\[64\].](#page-11-0) ERCC2, a DNA helicase and part of the NER system, reconstitutes intrastrand crosslinks induced by genotoxins like UV irradiation and cisplatin. Mutation of ERCC2 has been linked to the risk of breast cancer caused by ionizing radiation [\[65\]](#page-12-0). The activation of three important PIKK family enzymes, ATM, ATR, and DNA-PK is triggered by the creation of DSBs [\[66\]](#page-12-0) which initiates downstream signaling cascades to access DNA damage and trigger DNA repair [\[67\].](#page-12-0) Phosphorylation of γ-H2AX indicates the existence of DSB and directs proteins to repair irradiation-induced foci (IRIF) in initial stages [\[68\].](#page-12-0)

1.5. Radiotherapy sensitization targets in cancer

Cancer cells' natural ability to repair DNA damage may cause cellular resistance and restrict the effectiveness of treatment, despite the fact that radiation-induced DNA DSBs are the most efficient molecular events for eliminating cancer cells [\[69\].](#page-12-0) It is expected that target-based radiosensitization approaches would increase the effectiveness of radiotherapy by selectively sensitizing tumor tissue to ionizing radiation [\[70\]](#page-12-0). Recently, a variety of approaches have been used to develop radiosensitizers that are highly effective and have low toxicity [\[71\]](#page-12-0).

Targeting DDR signaling pathways has emerged as a promising approach to overcome tumor radioresistance, and significant advancements and discoveries have already been made in this area in recent years. Utilizing DDR to sensitize the cancer cells to ionizing radiation is considered as a viable therapeutic alternative for treating cancer patients effectively. [Table 1](#page-5-0) summarizes the types of DNA damageinducing therapies (cytotoxic chemotherapies, targeted therapies and combination approaches for radiation sensitization) employed in cancer treatment with few examples.

Table 1

• Gemcitabine with Radiotherapy • Carboplatin with Radiotherapy, and so on.

1.5.1. ATM/ATR

ATM was found during a clinical case observation when Gotoff et al. [\[98\]](#page-12-0) identified immunodeficiency in a patient with a rare inherited autosomal-recessive genetic A–T syndrome in 1967 [\[99\]](#page-12-0). Bentley et al. [\[100\]](#page-12-0) discovered in 1996 that ATR could improve esr1-1 radiosensitivity in *S. cerevisiae*. ATM is triggered and attracted to DSB sites by the MRN complex, which functions as a DNA damage sensor, whilst ATR is triggered and attracted to DSB sites by its binding partner ATR-interacting protein [\[12\].](#page-10-0) AZD1390, a brain penetrant ATM kinase inhibitor is highly potent and exhibits powerful activity in combination with radiation. This orally accessible drug is under Phase 1 clinical trial in patients with glioblastoma multiforme, brain metastases or leptomeningeal metastases. By preventing ATM from repairing tumor DNA, AZD1390 increases the likelihood that radiation will be able to kill cancer cells (NCT03423628) [\[101\].](#page-12-0) XRD-0394, an oral dual kinase inhibitor of ATM and DNA-PK is also under Phase 1 trial to evaluate in combination with palliative radiotherapy for the treatment of solid cancers that have spread or that are recurrent or locally progressed. This drug may enhance the effectiveness of radiotherapy by increasing the sensitivity and responsiveness of cancer tumors by inhibiting proteins that allow cells to respond to DNA damage caused by radiation (NCT05002140) [\[102\].](#page-12-0) AZD6738 (Ceralasertib), Berzosertib, BAY1895344, Schisandrin B, NU6027, and NVP-BEZ235 are all reported ATR inhibitors. A Phase I trial of AZD6738 was done to assess the biological effects, tolerance, and safety of palliative radiotherapy in cancer patients. To examine the impact of a fractionation schedule appropriate for radical therapy, the radiation dosage in this experiment was increased from 20 to 30 Gy over the course of 2 Gy fractions. If the combination treatment is well tolerated, the study is intended to move

on with a randomised trial that compares the addition of AZD6738 to standard-of-care radiotherapy or chemoradiotherapy in a patient group receiving treatment with radical intent (status of this trial is still unknown) [\[103\].](#page-12-0) Berzosertib, formerly named as (M6620, VX-970), a highly potent and selective inhibitor which has an IC_{50} of 19 nM is used in Phase 1 clinical trial to test the combination of M6620 with palliative radiotherapy in oesophageal cancer. Since this combination is well-tolerated, the combination of M6620 with chemoradiotherapy was evaluated to improve the current standard of care and provide a targeted, efficient method of treating oesophageal cancer and squamous cell carcinoma (NCT03641547) [\[104\].](#page-12-0) An oral inhibitor BAY1895344 in combination with stereotactic body radiation and pembrolizumab is also under Phase 1 clinical trial to study the possible benefits in recurrent and unresectable head and neck cancer (NCT04576091) [\[105\]](#page-12-0). As of Aug 30, 2022, there are 12 active clinical studies (clinicaltrials.gov) with ATM/ATR inhibitors alone or in combination with RT in malignancies such as ovarian and SCLC.

1.5.2. CDK1

CDK1 regulates cell cycle progression and the G1/S transition by controlling the centrosome cycle and mitotic initiation, regulating G1 advancement, and boosting the G2/M transition [\[106\]](#page-12-0). When ionizing radiation causes DNA damage, CDK1 is blocked, which causes the cell cycle to stop at G2 checkpoint, allowing DSB repair [\[107\].](#page-12-0) A non-specific CDK1 inhibitor MEK162, inhibits and dephosphorylates CDK1, CDK2, and Wee1 [\[108\]](#page-12-0), and is shown to delay DDR in glioblastoma cells after ionizing radiation. Raghavan et al., [\[109\]](#page-12-0) investigated AZD5438, a new-generation CDK1 inhibitor in lung cancer cell lines in conjunction with radiation. Radiation sensitivity of lung cancer cells was significantly boosted by AZD5438. In lymphoma, JNJ-7706621, a CDK1 and AURKA/B dual inhibitor has been demonstrated to reverse radioimmunotherapy resistance [\[110\].](#page-12-0) Dinaciclib, a CDK1/2/5/9 inhibitor, decreased tumor growth in ten out of ten subcutaneous pancreatic ductal adenocarcinoma mice models studied, with considerable growth reduction (*>*40%) in eight of ten [\[111\]](#page-13-0). Rohitukine, from which P276–00 is generated, is found to be a selective CDK4-D1 and CDK1-B inhibitor in preclinical studies. Cell cycle events are prevented at initial stage of development when these CDKs are inhibited, which results in cell cycle arrest between the G1-S transition. As a result, it may be effective while having fewer negative effects (NCT00408018) [\[112\]](#page-13-0). EM-1421, (also terameprocol) is used under Phase 1/2 clinical trials for patients with refractory solid cancers [\[113\].](#page-13-0) This inhibitor is also used in combination with survivin (inhibitor of apoptosis protein) in patients with hematological malignancies under Phase 1 trials $[114]$. With IC_{50} values ranging from 30 to 200 nM, AG-024322, a potent, multi-targeted inhibitor showed strong antiproliferative effect in various human cancer cell lines [\[115\]](#page-13-0). There are currently 16 clinical trials utilizing CDK1 inhibitors, with three active and the rest terminated. Although based on strong preclinical data, the results from several clinical trials showed only weak to moderate benefits. Moreover, no radiotherapy studies using this inhibitor are currently in clinical trials [116–[118\].](#page-13-0)

1.5.3. CDK4/6

CDK4 and CDK6, directly regulate cell-cycle transitions and cell division [\[119\].](#page-13-0) Cyclin-CDK complexes, which are largely formed by the association of CDK4 and CDK6, with D-type cyclins (Cyclin D1, D2, and D3), control the course of the cell cycle by phosphorylating the tumor suppressor protein retinoblastoma [[120](#page-13-0)]. The G1-S checkpoint, which controls genome replication in the cell cycle, is crucial for CDK4/6 [\[121\].](#page-13-0) [Fig. 3](#page-6-0) depicts where various DDR targeting drugs act in cell cycle. Radiosensitivity of multiple cell lines is increased by CDK4/6 inhibitors [\[122\].](#page-13-0) Three CDK4/6 inhibitors (palbociclib, ribociclib, and abemaciclib) were approved as combination treatments for HR+ , HER2-, and metastatic breast cancers in recent years [\[123\].](#page-13-0) Palbociclib, the first

Fig. 3. Cell cycle phases and DDR targeting in cancer $(G1 = \text{Gap/growth Phase})$ 1, S = DNA replication phase, G2 = Gap/growth phase 2, M = Cell division phase; ATM - Ataxia-Telangiesctasia Mutated, ATR - ataxia telangiectasia and Rad3-related, CDK1 - Cyclin-dependent kinase 1, CDK4/6 - Cyclin-dependent kinase 4/6, CHK1 - Checkpoint kinase 1, DNA-PK - DNA-dependent protein kinase, MPS1 - Monopolar spindle1, PARP - Poly (ADP-ribose) polymerases).

CDK4/6 inhibitor, received approval for women with untreated metastatic HR+ , HER2- breast cancer in 2015 on the basis of the outcomes of a randomized Phase 2 research carried out in a front-line setting (PAL-OMA-1 trial) [\[124\].](#page-13-0) According to Huang et al., [\[125\]](#page-13-0) combination of radiation and palbociclib significantly suppressed radiation-induced cell proliferation and decreased radiation resistance in hepatocellular carcinoma. As shown by Naz et al., [\[126\]](#page-13-0) palbociclib and ribociclib did not increase radiation sensitivity in non-small cell lung cancer cell lines when administered either pre- or post-IR. On the other hand, abemaciclib, when administered post-IR, it increased radiosensitivity in the non-small cell lung cancer cell lines [\[122\].](#page-13-0) *In vivo* studies showed that palbociclib with IR improved median survival time in brain cancer xenografts [\[127\].](#page-13-0) DeWire et al., [\[128\]](#page-13-0) reported the ribociclib administration following radiation therapy in diffuse intrinsic pontine gliomas and diffuse midline glioma. Increased tumor necrosis was observed as a side effect of this therapy (NCT02607124). In children with diffuse intrinsic pontine gliomas and high-grade glioma, a combination of ribociclib and everolimus following radiation therapy was well-tolerated in Phase 1 trials (NCT03355794) [\[129\].](#page-13-0) There are now six active clinical trials, the majority of which are directed at patients with HR+ /HER2- metastatic breast cancer. Beyond those included here, a number of ongoing trials continue to investigate these inhibitors in various patient populations and with various treatment regimens.

1.5.4. CHK1

CHK1 is essential for DNA repair activation and checkpoint-mediated cell-cycle arrest in response to DNA damage [\[130\].](#page-13-0) As cellular proliferation and survival are greatly controlled by CHK1, increasing evidence suggests that targeting them as prospective techniques for sensitizing cancer cells to radiation is a viable option [\[131\]](#page-13-0). The first selective CHK1 inhibitor identified was LY2606368 (Prexasertib). This inhibitor is used in Phase 1 clinical trials in combination with anti-cancer drugs (Cetuximab and Cisplatin) and radiation therapy for patients with head and neck cancer (NCT02555644) [\[132\].](#page-13-0) PF-00477736, a selective CHK1 inhibitor, significantly increased the radiosensitivity of multiple cancer cell lines. [\[133\].](#page-13-0) This inhibitor entered in Phase 1 clinical trials in combination with gemcitabine for advanced solid tumors, but the study was terminated due to business reasons (NCT00437203) [\[134\]](#page-13-0). CCT244747, the first orally accessible CHK1 inhibitor, sensitized head and neck cancer and bladder cancer cells to radiation through modulating G2/M checkpoint, implying that CCT244747 could be acceptable for oral administration [\[135,136\].](#page-13-0) Another inhibitor, SRA737, was discovered to have the ability to restrict cell proliferation when combined with Niraparib [\[137\]](#page-13-0). Several CHK1 inhibitors have entered into clinical trials which include AZD7762, GDC-0575, MK-8776, Prexasertib, PF-00477736, Rabusertib, and SRA737 [\[138\].](#page-13-0)

1.5.5. DNA-PKcs

DNA-PKcs was initially linked to repairing DSBs via the NHEJ pathway; however, later studies demonstrated that DNAPKcs has additional activities including the choice of NHEJ and HR repair pathways [139–[142\]](#page-13-0), cell cycle checkpoint regulation [\[143,144\],](#page-13-0) and telomere maintenance [\[145](#page-13-0)–147]. DNA end processing is made possible by autophosphorylation, which occurs when DNAPKcs autophosphorylates at Thr2609, Thr2647, and Ser2056 [148–[150\].](#page-13-0) DNA-PKcs depletion is known to increase cancer cells sensitivity to ionizing radiation and genotoxic chemotherapy [\[151,152\]](#page-14-0). Several DNA-PKCs inhibitors with varying selectivity and radiosensitization potential have been described including LY294002 [\[153,154\],](#page-14-0) NU7026 [\[155\],](#page-14-0) Vanillin, [\[156\]](#page-14-0), IC87102, IC87361, & IC86621 [\[157\]](#page-14-0), VX-984 [\[158\]](#page-14-0) and AZD7648 [\[159\]](#page-14-0) among others. Doxycycline was the first FDA approved DNA-PK inhibitor and is a good radiosensitizer of breast cancer cells [\[160\].](#page-14-0) Zenke et al., [\[161\]](#page-14-0) demonstrated the radiosensitizing efficacy of orally bioavailable M3814 (Peposertib), a DNAPK inhibitor in preclinical models. Currently, M3814 and XRD-0394 (ATM and DNA-PK dual inhibitor) are under Phase 1 trials in combination with radiotherapy to examine the tolerability and safety.

1.5.6. PARP-1

PARP-1 is a widely studied DNA DSB repair protein [\[162\].](#page-14-0) PARP-1 can catalyze the heterodimer produced by XPC-RAD23B and free PAR, suggesting that PARP-1 is involved in the radiation-induced DNA damage [\[163\]](#page-14-0). Inhibiting PARP-1 could make cancer cells more sensitive to radiation [\[164\].](#page-14-0) Several PARP inhibitors with varying degree of specificity and radiosensitization potential have been developed including KJ-28d [\[165\]](#page-14-0), ABT-888 [\[166\]](#page-14-0), Mk-4827 [\[167\],](#page-14-0) AZD2281 (Olaparib) [\[168\],](#page-14-0) Niraparib [\[169,170\],](#page-14-0) Veliparib, Rucaparib, Talazoparib, etc. Olaparib was the first PARP inhibitor to receive approval from the FDA and the European Union for treating advanced ovarian cancer patients with BRCA mutations. Olaparib received Breakthrough Therapy Designation from the FDA in 2016 for patients with metastatic castration-resistant prostate cancer who have mutations in BRCA1/2 or ATM. There are *>* 600 clinical trials at clinicaltrial.gov with search term "PARP" and *>* 60 when co-searched with "radiation" (as of Nov 30, 2022). These results indicate the importance of targeting PARP for anti-cancer therapies. There are 19 clinical trials on PARP-1 inhibitors in combination with radiation out of which 5 completed, 7 enrolling and rest not yet enrolled/withdrawn. Several of these trials will be game changer for patient management. For example, Olaparib is currently under Phase 1/2a trial combined with intensity modulated radiotherapy (IMRT) and temozolomide in first-line treatment of glioblastoma (NCT03212742) [\[171\]](#page-14-0). Similarly, Niraparib is being evaluated with Dostarlimab and radiation in either PD-L1 negative TNBC or in patients who failed to respond to prior immunotherapy (NCT04837209) [\[172\]](#page-14-0). Likewise, Veliparib is undergoing Phase 2 trial in combination with temozolomide and radiotherapy to assess its efficacy in treating individuals with malignant glioma (NCT03581292) [\[173\]](#page-14-0).

1.5.7. Wee1

Wee1 acts as a negative regulator of cell cycle during the G2 to M transition [\[174\]](#page-14-0) and Wee1 inhibition enhances cell death in response to chemotherapy and ionizing radiation by affecting both cell cycle progression and DNA damage repair [\[175\].](#page-14-0) Several Wee1 inhibitors have shown radiosensitization in preclinical models including Wee 1 inhibitor

II (681641) [\[176\],](#page-14-0) R1–1 [\[176\]](#page-14-0), PD0166285 [\[177\]](#page-14-0), AZD1775 (also MK-1775, Adavosertib) [\[178](#page-14-0)–180]. Four Wee 1 inhibitors IMP7068, AZD1775, Zn-c3, and SY-4835 have moved into the clinical trials. There are more than 50 clinical trials for AZD1775 listed on the clinical trial website (as of Nov 30, 2022). Majority of these clinical studies have used a combination of chemotherapy and radiation to treat different cancers and identified that p53 mutation may be a cancer- or chemotherapy-specific prognostic biomarker for responsiveness to Wee1 inhibition. Early results from Wee1 clinical studies have been encouraging. For example, AZD1775 improved 1-year survival to 90% in head and neck cancer when combined with radiation therapy and cisplatin [\[180\].](#page-14-0) AZD1775 also improved overall survival in locally advanced pancreatic adenocarcinoma (Phase 2 trial) when combined with radiotherapy and gemcitabine (NCT02037230) [\[181\].](#page-14-0)

1.5.8. MPS1/TTK

Monopolar spindle 1 (MPS1) is an important spindle assembly checkpoint kinase. It is overexpressed in several cancers including breast, bronchogenic, lung, and thyroid papillary carcinomas [\[182\]](#page-14-0). Increased MPS1 levels correlate with a higher histological grade [\[183\]](#page-14-0). It is implicated in the genotoxic stress response, such as stress produced by DNA damage. Inhibition of MPS1 causes cell death by polyploidization and mitotic collapse [\[184\]](#page-14-0) and prevents DNA repair after RT, allowing DNA damage to accumulate leading to mitotic catastrophe. Genetic and pharmacological MPS1/TTK inhibition is shown to radiosensitize basal-like breast [\[185\]](#page-14-0) and glioblastoma [\[186\]](#page-14-0) cell lines. Inhibitors of MPS1 (Mps-BAY2b and MPS-IN-3) have been demonstrated in human colon cancer and glioblastoma cells to improve sensitivity to microtubule poisons (Paclitaxel and Vincristine) [\[187\].](#page-14-0) Currently, there are three MPS1 inhibitors (BAY1217389, BAY1161909, and BOS172722) in clinical trials. BAY1217389, an oral selective inhibitor led to increased toxicity (nausea, fatigue and diarrhea) when given in combination with paclitaxel in patients with solid tumors (NCT02366949) [\[188\]](#page-14-0). BAY1161909 has demonstrated enhanced efficacy in xenograft models when combined with paclitaxel and found to be well-tolerated with manageable adverse events (Phase 1) in patients with advanced malignancies (NCT02138812) [\[189\]](#page-14-0). BOS172722, another orally bioavailable inhibitor is currently under clinical trials in combination with paclitaxel for patients with advanced non-hematologic malignancies (NCT03328494) [\[190\].](#page-14-0) MPS1 inhibitors have not been evaluated in combination with radiation in human patients. The [Table 2](#page-8-0) below lists molecular targets evaluated in clinical trials for radiosensitizing different cancers.

1.6. Other novel targets

1.6.1. DNA LIG4

In radiation-induced NHEJ pathway, the DNA LIG4 is an important DNA repair factor [\[191\]](#page-14-0). XRCC4, LIG4, and Cernunos-XLF are commonly recruited to the break site and form temporary connections with the DNA ends to guarantee that the break is ligated [\[192\]](#page-15-0). Patients with LIG4 syndrome have higher radiation sensitivity, higher chance of neurological problems, bone marrow dysfunction, and cancer predisposition [\[193\].](#page-15-0) LIG4 mutations have been linked to clinical radiosensitization in a number of investigations. According to Riballo et al., [\[194\]](#page-15-0) LIG4 mutation impairs synthesis of adenylate complex while also lowering rejoining activity. Furthermore, healthy patients with LIG4 rs1805388 polymorphism were more vulnerable to radiation than healthy individuals by H2AX foci analysis [\[195\]](#page-15-0). DNA-binding protein-1 suppresses the production of LIG4 and hence negatively influences DNA repair mechanisms. In a screen of 5280 compounds, Tseng et al., [\[196\]](#page-15-0) showed that rabeprazole and U73122 could selectively disrupt adenylate transfer phase and DNA rejoining to impede radiation-induced DNA repair by targeting LIG4. NU7026 affects the radiosensitivity of LIG4 wild-type mouse embryonic fibroblasts [\[197\].](#page-15-0) A specific DNA LIG4

inhibitor SCR7 blocks NHEJ [\[198\]](#page-15-0) and enhances HR-mediated gene editing in mammalian cells [\[199\].](#page-15-0) It is found to effectively inhibit the cell proliferation of various breast, lung and cervical cancer cell lines with IC₅₀ values ranging from 5 to 50 μ M [\[200\].](#page-15-0) Treatment with SCR7 in vivo prolongs lifespan by four times and lowers breast adenocarcinoma-induced tumor. It greatly increases the cytotoxic effects of radiation, etoposide, and 3-Aminobenzamide on tumors generated from Dalton's lymphoma cells [\[200\]](#page-15-0). Further research into LIG4 activities in the radiation-induced DDR will be useful in characterizing it as a molecular target for radiosensitization.

1.6.2. BUB1

The spindle assembly checkpoint ensures faithful chromosome segregation during cell division [\[201\]](#page-15-0). Budding Uninhibited by Benzimidazole 1 (BUB1) is a key component of spindle assembly checkpoint and it also facilitates chromosome alignment and mitotic duration [\[202\].](#page-15-0) Upregulation of mitotic factors is a significantly more common event in human tumors, and increased BUB1 levels have been reported in various cancers including breast, gastric, lung, brain and lymphoma [\[203\].](#page-15-0) Furthermore, BUB1 upregulation is linked to a poor clinical outcome in different tumor types [\[204\].](#page-15-0) In transgenic mice, BUB1 overexpression was observed to promote spontaneous carcinogenesis and promote the development of Myc-induced lymphoma [\[205\]](#page-15-0). Reduced BUB1 mRNA levels in colon carcinomas were associated with shorter relapse-free survival after surgery [\[206\].](#page-15-0) BUB1 has been identified as a possible radio-enhancing target and the lack of mitotic spindle checkpoints may actually elicit fatal mitotic events during RT, as BUB1 suppression was correlated with an increased production of micronuclei [\[207\].](#page-15-0) A focused human kinome screen identified BUB1 as one of the potential radiosensitizing target in breast cancer [\[208\]](#page-15-0). BUB1 is known to localize near DSB sites where Rad53 and H2AX are also recruited [\[209\]](#page-15-0) and it also co-localizes with 53BP1 suggesting a role in NHEJ pathway. Knockdown of BUB1 results in prolonged H2AX foci and comet tail formation as well as hypersensitivity in response to ionizing radiation [\[210\].](#page-15-0) BUB1 expression is known to associate with poor outcome in a study analyzing DNA repair gene expression patterns as prognostic and predictive factors in breast cancer subtypes [\[211\]](#page-15-0). To enhance the efficacy of cancer therapies, further research into the underlying mechanisms of BUB1 mediated DDR is needed. Our lab is actively seeking to decipher the mechanism of BUB1 mediated radioresistance by combining genomic (siRNA, CRISPR) and biochemical (BAY1816032) ablation of BUB1 with biochemical, molecular, and radio-biological techniques.

2. Conclusion and future perspectives

Deficits in the DNA damage repair pathway may sensitize cancer cells to cytotoxic agents. Drugs that target the DDR have been clinically validated in small groups of patients, and various combination approaches to block multiple pathways that cancer cells rely on for survival are now being investigated. Indeed, better understanding the fundamental mechanisms of radiation-induced DDR and specific functions of major genes and proteins in DDR pathways, as well as their interaction partners, are critical for clinical identification of novel intervention targets and the development of efficacious cancer treatments. A variety of tumors are being targeted by the therapeutic landscape of anti-tumor drugs that target the DDR, which has rapidly extended to encompass inhibitors of other crucial mediators of DNA repair [\[212\]](#page-15-0). While increasing the effectiveness of killing tumor cells is the main goal of radiation-drug combinations, it is also critical to reduce the toxicities to normal tissue because the therapeutic benefit depends on the difference between efficacy and toxicity. Recently, MPS1/TTK has entered clinical trials but only with Taxols (chemotherapy). It will be interesting to see if a combination of TTK inhibitor with radiation provides any clinical benefit. Additionally, novel molecular targets such as (BUB1, LIG4 etc.) warrant further clinical exploration for evaluating

Table 2

Molecular targets evaluated in clinical trials in combination with DNA-damaging agents or radiation.

Table 2 (*continued*)

(*continued on next page*)

* Targets that exclude combination with radiation therapy as there are no radiation therapy studies in clinical trials (as on Nov 30, 2022)

their usability in enhancing radiation-induced cell killing and improving clinical outcomes. A greater knowledge is needed into how these new targets and their inhibitors contribute to radiation-induced DNA damage leading to radiation sensitization. We believe that a better knowledge of the molecular processes behind the DDR, as well as the genetic interactions between distinct DDR pathways and other cellular pathways such as epithelial to mesenchymal transition (EMT), cell-cycle regulation, cancer stem cells (CSCs) resilience, tumor-immune microenvironment will lead to new treatment options for a variety of human diseases including cancer.

CRediT authorship contribution statement

Shyam Nyati, Sushmitha Sriramulu: Conceptualization. **Shyam Nyati:** Supervision. **Sushmitha Sriramulu:** Writing – original draft. **Shyam Nyati, Stephen L. Brown, Farzan Siddiqui, Benjamin Movsas, Shivani Thoidingjam, Sushmitha Sriramulu:** Writing − review & editing. All authors read and approved the final manuscript.

Conflict of interest statement

Benjamin Movsas receives research support from Varian Medical Systems, ViewRay, and Philips. Farzan Siddiqui has received honoraria from Varian Medical Systems for lectures and presentations, reimbursement for travel, food and lodging expenses; Honoraria from American College of Radiology for services provided as Radiation Oncology Practice Accreditation site surveyor, reimbursement for travel, food and lodging expenses; Honoraria from Varian Noona Medical Advisory Board for services as a board member. All other authors have no conflicts of interest related to this publication.

Data availability

No data was used for the research described in the article.

Acknowledgments

This work was supported by NCI R21 (1R21CA252010–01A1), Henry Ford Health System (HFHS) Research Administration Start up, Henry Ford Health System (HFHS) Proposal Development Award and Henry Ford Health System (HFHS)-Radiation Oncology Start up to Shyam Nyati. We also thank Henry Ford Cancer Institute (HFCI) for

providing Translational Oncology Postdoctoral Fellowship to Sushmitha Sriramulu.

Ethics approval and consent to participate

Not Applicable.

References

- [1] N. Chatterjee, G.C. Walker, Mechanisms of DNA damage, repair, and mutagenesis, Environ. Mol. Mutagen 58 (5) (2017) 235–263, [https://doi.org/](https://doi.org/10.1002/em.22087) [10.1002/em.22087](https://doi.org/10.1002/em.22087).
- [2] R. Visconti, D. Grieco, New insights on oxidative stress in cancer, Curr. Opin. Drug Disco Devel 12 (2) (2009) 240–245. 〈[https://www.ncbi.nlm.nih.gov/](https://www.ncbi.nlm.nih.gov/pubmed/19333869) d/19333869
- [3] S.P. Jackson, J. Bartek, The DNA-damage response in human biology and disease, Nature 461 (7267) (2009) 1071–1078, <https://doi.org/10.1038/nature08467>.
- [4] D. Hanahan, R.A. Weinberg, Hallmarks of cancer: the next generation, Cell 144 (5) (2011) 646–674, [https://doi.org/10.1016/j.cell.2011.02.013.](https://doi.org/10.1016/j.cell.2011.02.013)
- [5] N. Hosoya, K. Miyagawa, Targeting DNA damage response in cancer therapy, Cancer Sci. 105 (4) (2014) 370–388, [https://doi.org/10.1111/cas.12366.](https://doi.org/10.1111/cas.12366)
- [6] M.J. O'Connor, Targeting the DNA damage response in cancer, Mol. Cell 60 (4) (2015) 547–560,<https://doi.org/10.1016/j.molcel.2015.10.040>.
- [7] S.P. Jackson, T. Helleday, DNA REPAIR. Drugging DNA repair, Science 352 (6290) (2016) 1178-1179, https://doi.org/10.1126/science.aab095
- [8] S. Ohta, Contribution of somatic mutations in the mitochondrial genome to the development of cancer and tolerance against anticancer drugs, Oncogene 25 (34) (2006) 4768–4776, [https://doi.org/10.1038/sj.onc.1209602.](https://doi.org/10.1038/sj.onc.1209602)
- I. Kamileri, I. Karakasilioti, G.A. Garinis, Nucleotide excision repair: new tricks with old bricks, Trends Genet 28 (11) (2012) 566–573, [https://doi.org/10.1016/](https://doi.org/10.1016/j.tig.2012.06.004) tig.2012.06.004.
- [10] P. Hsieh, K. Yamane, DNA mismatch repair: molecular mechanism, cancer, and ageing, Mech. Ageing Dev. 129 (7–8) (2008) 391–407, [https://doi.org/10.1016/](https://doi.org/10.1016/j.mad.2008.02.012) [j.mad.2008.02.012.](https://doi.org/10.1016/j.mad.2008.02.012)
- [11] J. Bartkova, Z. Horejsi, K. Koed, A. Kramer, F. Tort, K. Zieger, P. Guldberg, M. Sehested, J.M. Nesland, C. Lukas, T. Orntoft, J. Lukas, J. Bartek, DNA damage response as a candidate anti-cancer barrier in early human tumorigenesis, Nature 434 (7035) (2005) 864–870, <https://doi.org/10.1038/nature03482>.
- [12] J.H. Lee, T.T. Paull, ATM activation by DNA double-strand breaks through the Mre11-Rad50-Nbs1 complex, Science 308 (5721) (2005) 551–554, [https://doi.](https://doi.org/10.1126/science.1108297) [org/10.1126/science.1108297](https://doi.org/10.1126/science.1108297).
- [13] C.J. Bakkenist, M.B. Kastan, DNA damage activates ATM through intermolecular autophosphorylation and dimer dissociation, Nature 421 (6922) (2003) 499–506, <https://doi.org/10.1038/nature01368>.
- [14] S. Matsuoka, B.A. Ballif, A. Smogorzewska, E.R. McDonald 3rd, K.E. Hurov, J. Luo, C.E. Bakalarski, Z. Zhao, N. Solimini, Y. Lerenthal, Y. Shiloh, S.P. Gygi, S. J. Elledge, ATM and ATR substrate analysis reveals extensive protein networks responsive to DNA damage, Science 316 (5828) (2007) 1160–1166, [https://doi.](https://doi.org/10.1126/science.1140321) [org/10.1126/science.1140321](https://doi.org/10.1126/science.1140321).
- [15] L. Zou, S.J. Elledge, Sensing DNA damage through ATRIP recognition of RPAssDNA complexes, Science 300 (5625) (2003) 1542–1548, [https://doi.org/](https://doi.org/10.1126/science.1083430) [10.1126/science.1083430.](https://doi.org/10.1126/science.1083430)
- [16] H. Zhao, H. Piwnica-Worms, ATR-mediated checkpoint pathways regulate phosphorylation and activation of human Chk1, Mol. Cell Biol. 21 (13) (2001) 4129–4139, [https://doi.org/10.1128/MCB.21.13.4129-4139.2001.](https://doi.org/10.1128/MCB.21.13.4129-4139.2001)
- [17] A.J. Hartlerode, R. Scully, Mechanisms of double-strand break repair in somatic mammalian cells, Biochem J. 423 (2) (2009) 157–168, [https://doi.org/10.1042/](https://doi.org/10.1042/BJ20090942) [BJ20090942.](https://doi.org/10.1042/BJ20090942)
- [18] S. Britton, J. Coates, S.P. Jackson, A new method for high-resolution imaging of Ku foci to decipher mechanisms of DNA double-strand break repair, J. Cell Biol. 202 (3) (2013) 579-595, https://doi.org/10.1083/jcb.20130307
- [19] T.M. Gottlieb, S.P. Jackson, The DNA-dependent protein kinase: requirement for DNA ends and association with Ku antigen, Cell 72 (1) (1993) 131-142, https:/ [doi.org/10.1016/0092-8674\(93\)90057-w.](https://doi.org/10.1016/0092-8674(93)90057-w)
- [20] E.J. Park, D.W. Chan, J.H. Park, M.A. Oettinger, J. Kwon, DNA-PK is activated by nucleosomes and phosphorylates H2AX within the nucleosomes in an acetylationdependent manner, Nucleic Acids Res 31 (23) (2003) 6819–6827, [https://doi.](https://doi.org/10.1093/nar/gkg921) [org/10.1093/nar/gkg921](https://doi.org/10.1093/nar/gkg921).
- [21] Y. Ma, U. Pannicke, K. Schwarz, M.R. Lieber, Hairpin opening and overhang processing by an Artemis/DNA-dependent protein kinase complex in nonhomologous end joining and V(D)J recombination, Cell 108 (6) (2002) 781–794, [https://doi.org/10.1016/s0092-8674\(02\)00671-2](https://doi.org/10.1016/s0092-8674(02)00671-2).
- [22] S.A. Nick McElhinny, C.M. Snowden, J. McCarville, D.A. Ramsden, Ku recruits the XRCC4-ligase IV complex to DNA ends, Mol. Cell Biol. 20 (9) (2000) 2996–3003, s://doi.org/10.1128/MCB.20.9.2996-3003.2000.
- [23] P. Ahnesorg, P. Smith, S.P. Jackson, XLF interacts with the XRCC4-DNA ligase IV complex to promote DNA nonhomologous end-joining, Cell 124 (2) (2006) 301–313, [https://doi.org/10.1016/j.cell.2005.12.031.](https://doi.org/10.1016/j.cell.2005.12.031)
- [24] M.T. Hemann, From breaking bad to worse: exploiting homologous DNA repair deficiency in cancer, Cancer Disco 4 (5) (2014) 516–518, [https://doi.org/](https://doi.org/10.1158/2159-8290.CD-14-0316) [10.1158/2159-8290.CD-14-0316](https://doi.org/10.1158/2159-8290.CD-14-0316).
- [25] A.A. Sartori, C. Lukas, J. Coates, M. Mistrik, S. Fu, J. Bartek, R. Baer, J. Lukas, S. P. Jackson, Human CtIP promotes DNA end resection, Nature 450 (7169) (2007) 509–514, [https://doi.org/10.1038/nature06337.](https://doi.org/10.1038/nature06337)
- [26] J.A. Solinger, W.D. Heyer, Rad54 protein stimulates the postsynaptic phase of Rad51 protein-mediated DNA strand exchange, Proc. Natl. Acad. Sci. USA 98 (15) (2001) 8447-8453, https://doi.org/10.1073/pnas.121009
- [27] K. Rodgers, M. McVey, Error-Prone Repair of DNA Double-Strand Breaks, J. Cell Physiol. 231 (1) (2016) 15–24, [https://doi.org/10.1002/jcp.25053.](https://doi.org/10.1002/jcp.25053)
- [28] R. Scully, A. Panday, R. Elango, N.A. Willis, DNA double-strand break repairpathway choice in somatic mammalian cells, Nat. Rev. Mol. Cell Biol. 20 (11) (2019) 698–714, <https://doi.org/10.1038/s41580-019-0152-0>.
- [29] V.G. Gorgoulis, L.V. Vassiliou, P. Karakaidos, P. Zacharatos, A. Kotsinas, T. Liloglou, M. Venere, R.A. Ditullio Jr., N.G. Kastrinakis, B. Levy, D. Kletsas, A. Yoneta, M. Herlyn, C. Kittas, T.D. Halazonetis, Activation of the DNA damage checkpoint and genomic instability in human precancerous lesions, Nature 434 (7035) (2005) 907–913,<https://doi.org/10.1038/nature03485>.
- [30] M. Kase, M. Vardja, A. Lipping, T. Asser, J. Jaal, Impact of PARP-1 and DNA-PK expression on survival in patients with glioblastoma multiforme, Radio. Oncol. 101 (1) (2011) 127–131, [https://doi.org/10.1016/j.radonc.2011.06.024.](https://doi.org/10.1016/j.radonc.2011.06.024)
- [31] Y. Li, W.Y. Wang, J.H. Xiao, F. Xu, D.Y. Liao, L. Xie, J. Wang, F. Luo, Overexpression of Rad51 Predicts Poor Prognosis in Colorectal Cancer: Our Experience with 54 Patients, PLoS One 12 (1) (2017), e0167868, https://doi.org/ [10.1371/journal.pone.0167868](https://doi.org/10.1371/journal.pone.0167868).
- [32] P. Bouchaert, S. Guerif, C. Debiais, J. Irani, G. Fromont, DNA-PKcs expression predicts response to radiotherapy in prostate cancer, Int J. Radiat. Oncol. Biol. Phys. 84 (5) (2012) 1179–1185, <https://doi.org/10.1016/j.ijrobp.2012.02.014>.
- [33] R. Brosh, V. Rotter, When mutants gain new powers: news from the mutant p53 field, Nat. Rev. Cancer 9 (10) (2009) 701–713, [https://doi.org/10.1038/nrc2693.](https://doi.org/10.1038/nrc2693)
- [34] A.S. Coutts, C.J. Adams, N.B. La Thangue, p53 ubiquitination by Mdm2: a never ending tail? DNA Repair (Amst.) 8 (4) (2009) 483–490, [https://doi.org/](https://doi.org/10.1016/j.dnarep.2009.01.008) [10.1016/j.dnarep.2009.01.008.](https://doi.org/10.1016/j.dnarep.2009.01.008)
- [35] C.A. Cremona, A. Behrens, ATM signalling and cancer, Oncogene 33 (26) (2014) 3351–3360, <https://doi.org/10.1038/onc.2013.275>.
- [36] A.V. Biankin, N. Waddell, K.S. Kassahn, M.C. Gingras, L.B. Muthuswamy, A. L. Johns, D.K. Miller, P.J. Wilson, A.M. Patch, J. Wu, D.K. Chang, M.J. Cowley, B. B. Gardiner, S. Song, I. Harliwong, S. Idrisoglu, C. Nourse, E. Nourbakhsh, S. Manning, S.M. Grimmond, Pancreatic cancer genomes reveal aberrations in axon guidance pathway genes, Nature 491 (7424) (2012) 399–405, [https://doi.](https://doi.org/10.1038/nature11547) $re/10.1038/nature11$
- [37] C.S. Grasso, Y.M. Wu, D.R. Robinson, X. Cao, S.M. Dhanasekaran, A.P. Khan, M. J. Quist, X. Jing, R.J. Lonigro, J.C. Brenner, I.A. Asangani, B. Ateeq, S.Y. Chun, J. Siddiqui, L. Sam, M. Anstett, R. Mehra, J.R. Prensner, N. Palanisamy, S. A. Tomlins, The mutational landscape of lethal castration-resistant prostate cancer, Nature 487 (7406) (2012) 239–243, [https://doi.org/10.1038/](https://doi.org/10.1038/nature11125) [nature11125](https://doi.org/10.1038/nature11125).
- [38] N. Cancer Genome Atlas, Comprehensive molecular portraits of human breast tumours, Nature 490 (7418) (2012) 61–70, [https://doi.org/10.1038/](https://doi.org/10.1038/nature11412) [nature11412](https://doi.org/10.1038/nature11412).
- [39] S. Banerjee, S. Kaye, PARP inhibitors in BRCA gene-mutated ovarian cancer and beyond, Curr. Oncol. Rep. 13 (6) (2011) 442–449, [https://doi.org/10.1007/](https://doi.org/10.1007/s11912-011-0193-9) [s11912-011-0193-9](https://doi.org/10.1007/s11912-011-0193-9).
- [40] S. Angele, I. Treilleux, A. Bremond, P. Taniere, J. Hall, Altered expression of DNA double-strand break detection and repair proteins in breast carcinomas, Histopathology 43 (4) (2003) 347–353, [https://doi.org/10.1046/j.1365-](https://doi.org/10.1046/j.1365-2559.2003.01713.x) $2003.01713.x$
- [41] L. Ai, Q.N. Vo, C. Zuo, L. Li, W. Ling, J.Y. Suen, E. Hanna, K.D. Brown, C.Y. Fan, Ataxia-telangiectasia-mutated (ATM) gene in head and neck squamous cell

carcinoma: promoter hypermethylation with clinical correlation in 100 cases, Cancer Epidemiol. Biomark. Prev. 13 (1) (2004) 150–156, [https://doi.org/](https://doi.org/10.1158/1055-9965.epi-082-3) [10.1158/1055-9965.epi-082-3](https://doi.org/10.1158/1055-9965.epi-082-3).

- [42] J. Bartkova, J. Tommiska, L. Oplustilova, K. Aaltonen, A. Tamminen, T. Heikkinen, M. Mistrik, K. Aittomaki, C. Blomqvist, P. Heikkila, J. Lukas, H. Nevanlinna, J. Bartek, Aberrations of the MRE11-RAD50-NBS1 DNA damage sensor complex in human breast cancer: MRE11 as a candidate familial cancerpredisposing gene, Mol. Oncol. 2 (4) (2008) 296–316, [https://doi.org/10.1016/j.](https://doi.org/10.1016/j.molonc.2008.09.007) $.2008$
- [43] D.S. Kim, M.J. Kim, J.Y. Lee, S.M. Lee, J.E. Choi, S.Y. Lee, J.Y. Park, Epigenetic inactivation of checkpoint kinase 2 gene in non-small cell lung cancer and its relationship with clinicopathological features, Lung Cancer 65 (2) (2009) 247–250, <https://doi.org/10.1016/j.lungcan.2009.03.011>.
- [44] A. Sullivan, M. Yuille, C. Repellin, A. Reddy, O. Reelfs, A. Bell, B. Dunne, B. A. Gusterson, P. Osin, P.J. Farrell, I. Yulug, A. Evans, T. Ozcelik, M. Gasco, T. Crook, Concomitant inactivation of p53 and Chk2 in breast cancer, Oncogene 21 (9) (2002) 1316–1324, [https://doi.org/10.1038/sj.onc.1205207.](https://doi.org/10.1038/sj.onc.1205207)
- [45] O. Galamb, F. Sipos, E. Dinya, S. Spisak, Z. Tulassay, B. Molnar, mRNA expression, functional profiling and multivariate classification of colon biopsy specimen by cDNA overall glass microarray, World J. Gastroenterol. 12 (43) (2006) 6998–7006, [https://doi.org/10.3748/wjg.v12.i43.6998.](https://doi.org/10.3748/wjg.v12.i43.6998)
- [46] K. Yoshikawa, T. Ogawa, R. Baer, H. Hemmi, K. Honda, A. Yamauchi, T. Inamoto, K. Ko, S. Yazumi, H. Motoda, H. Kodama, S. Noguchi, A.F. Gazdar, Y. Yamaoka, R. Takahashi, Abnormal expression of BRCA1 and BRCA1-interactive DNA-repair proteins in breast carcinomas, Int J. Cancer 88 (1) (2000) 28–36. 〈[https://www.](https://www.ncbi.nlm.nih.gov/pubmed/10962436) [ncbi.nlm.nih.gov/pubmed/10962436](https://www.ncbi.nlm.nih.gov/pubmed/10962436)〉.
- [47] J.L. Hilton, J.P. Geisler, J.A. Rathe, M.A. Hattermann-Zogg, B. DeYoung, R. E. Buller, Inactivation of BRCA1 and BRCA2 in ovarian cancer, J. Natl. Cancer Inst. 94 (18) (2002) 1396–1406, [https://doi.org/10.1093/jnci/94.18.1396.](https://doi.org/10.1093/jnci/94.18.1396)
- [48] A. Catteau, J.R. Morris, BRCA1 methylation: a significant role in tumour development? Semin Cancer Biol. 12 (5) (2002) 359–371, [https://doi.org/](https://doi.org/10.1016/s1044-579x(02)00056-1) [10.1016/s1044-579x\(02\)00056-1](https://doi.org/10.1016/s1044-579x(02)00056-1).
- [49] N. Hosoya, M. Okajima, A. Kinomura, Y. Fujii, T. Hiyama, J. Sun, S. Tashiro, K. Miyagawa, Synaptonemal complex protein SYCP3 impairs mitotic recombination by interfering with BRCA2, EMBO Rep. 13 (1) (2011) 44–51, [https://doi.org/10.1038/embor.2011.221.](https://doi.org/10.1038/embor.2011.221)
- [50] M.B. Kastan, J. Bartek, Cell-cycle checkpoints and cancer, Nature 432 (7015) (2004) 316–323, <https://doi.org/10.1038/nature03097>.
- [51] W. Burkart, T. Jung, G. Frasch, Damage pattern as a function of radiation quality and other factors, C. R. Acad. Sci. III 322 (2–3) (1999) 89–101, https://doi.org [10.1016/s0764-4469\(99\)80029-8](https://doi.org/10.1016/s0764-4469(99)80029-8).
- [52] I.R. Radford, The level of induced DNA double-strand breakage correlates with cell killing after X-irradiation, Int J. Radiat. Biol. Relat. Stud. Phys. Chem. Med 48 (1) (1985) 45–54,<https://doi.org/10.1080/09553008514551051>.
- [53] A. Kuzminov, Single-strand interruptions in replicating chromosomes cause double-strand breaks, Proc. Natl. Acad. Sci. USA 98 (15) (2001) 8241–8246, [https://doi.org/10.1073/pnas.131009198.](https://doi.org/10.1073/pnas.131009198)
- [54] J. Biau, F. Devun, W. Jdey, E. Kotula, M. Quanz, E. Chautard, M. Sayarath, J. S. Sun, P. Verrelle, M. Dutreix, A preclinical study combining the DNA repair inhibitor Dbait with radiotherapy for the treatment of melanoma, Neoplasia 16 (10) (2014) 835–844, <https://doi.org/10.1016/j.neo.2014.08.008>.
- [55] H.E. Krokan, M. Bjoras, Base excision repair, Cold Spring Harb. Perspect. Biol. 5 (4) (2013), a012583,<https://doi.org/10.1101/cshperspect.a012583>.
- [56] B. van Loon, E. Markkanen, U. Hubscher, Oxygen as a friend and enemy: How to combat the mutational potential of 8-oxo-guanine, DNA Repair (Amst.) 9 (6) (2010) 604–616, [https://doi.org/10.1016/j.dnarep.2010.03.004.](https://doi.org/10.1016/j.dnarep.2010.03.004)
- [57] T. Visnes, M. Grube, B.M.F. Hanna, C. Benitez-Buelga, A. Cazares-Korner, T. Helleday, Targeting BER enzymes in cancer therapy, DNA Repair (Amst.) 71 (2018) 118–126, [https://doi.org/10.1016/j.dnarep.2018.08.015.](https://doi.org/10.1016/j.dnarep.2018.08.015)
- [58] C. Vens, A.C. Begg, Targeting base excision repair as a sensitization strategy in radiotherapy, Semin Radiat. Oncol. 20 (4) (2010) 241–249, [https://doi.org/](https://doi.org/10.1016/j.semradonc.2010.05.005) [10.1016/j.semradonc.2010.05.005](https://doi.org/10.1016/j.semradonc.2010.05.005).
- [59] J.W. Hyun, G.J. Cheon, H.S. Kim, Y.S. Lee, E.Y. Choi, B.H. Yoon, J.S. Kim, M. H. Chung, Radiation sensitivity depends on OGG1 activity status in human leukemia cell lines, Free Radic. Biol. Med 32 (3) (2002) 212–220, [https://doi.](https://doi.org/10.1016/s0891-5849(01)00793-6) [org/10.1016/s0891-5849\(01\)00793-6.](https://doi.org/10.1016/s0891-5849(01)00793-6)
- [60] N. Yang, H. Galick, S.S. Wallace, Attempted base excision repair of ionizing radiation damage in human lymphoblastoid cells produces lethal and mutagenic double strand breaks, DNA Repair (Amst.) 3 (10) (2004) 1323–1334, [https://doi.](https://doi.org/10.1016/j.dnarep.2004.04.014) [org/10.1016/j.dnarep.2004.04.014.](https://doi.org/10.1016/j.dnarep.2004.04.014)
- [61] P. Fortini, E. Dogliotti, Base damage and single-strand break repair: mechanisms and functional significance of short- and long-patch repair subpathways, DNA Repair (Amst.) 6 (4) (2007) 398–409, [https://doi.org/10.1016/j.](https://doi.org/10.1016/j.dnarep.2006.10.008) [dnarep.2006.10.008](https://doi.org/10.1016/j.dnarep.2006.10.008).
- [62] M.J. Schiewer, A.C. Mandigo, N. Gordon, F. Huang, S. Gaur, R. de Leeuw, S. G. Zhao, J. Evans, S. Han, T. Parsons, R. Birbe, P. McCue, C. McNair, S.N. Chand, Y. Cendon-Florez, P. Gallagher, J.J. McCann, N. Poudel Neupane, A.A. Shafi, K. E. Knudsen, PARP-1 regulates DNA repair factor availability, EMBO Mol. Med 10 (12) (2018),<https://doi.org/10.15252/emmm.201708816>.
- [63] P. Rajaraman, P. Bhatti, M.M. Doody, S.L. Simon, R.M. Weinstock, M.S. Linet, M. Rosenstein, M. Stovall, B.H. Alexander, D.L. Preston, A.J. Sigurdson, Nucleotide excision repair polymorphisms may modify ionizing radiation-related breast cancer risk in US radiologic technologists, Int J. Cancer 123 (11) (2008) 2713-2716, https://doi.org/10.1002/ijc.237
- [64] M. Hosseini, K. Ezzedine, A. Taieb, H.R. Rezvani, Oxidative and energy metabolism as potential clues for clinical heterogeneity in nucleotide excision

repair disorders, J. Invest Dermatol. 135 (2) (2015) 341–351, [https://doi.org/](https://doi.org/10.1038/jid.2014.365) [10.1038/jid.2014.365.](https://doi.org/10.1038/jid.2014.365)

- [65] Y. Zhang, L.H. Rohde, H. Wu, Involvement of nucleotide excision and mismatch repair mechanisms in double strand break repair, Curr. Genom. 10 (4) (2009) 250–258, <https://doi.org/10.2174/138920209788488544>.
- [66] C.A. Lovejoy, D. Cortez, Common mechanisms of PIKK regulation, DNA Repair (Amst.) 8 (9) (2009) 1004–1008, [https://doi.org/10.1016/j.](https://doi.org/10.1016/j.dnarep.2009.04.006) harep.2009.04.00
- [67] A.N. Blackford, S.P. Jackson, ATM, ATR, and DNA-PK: The Trinity at the Heart of the DNA Damage Response, Mol. Cell 66 (6) (2017) 801-817, https://doi.org/ [10.1016/j.molcel.2017.05.015.](https://doi.org/10.1016/j.molcel.2017.05.015)
- [68] D. Cortez, Replication-Coupled DNA Repair, Mol. Cell 74 (5) (2019) 866–876, [https://doi.org/10.1016/j.molcel.2019.04.027.](https://doi.org/10.1016/j.molcel.2019.04.027)
- [69] R.X. Huang, P.K. Zhou, DNA damage response signaling pathways and targets for radiotherapy sensitization in cancer, Signal Transduct. Target Ther. 5 (1) (2020) 60, [https://doi.org/10.1038/s41392-020-0150-x.](https://doi.org/10.1038/s41392-020-0150-x)
- [70] M.R. Berry, T.M. Fan, Target-Based Radiosensitization Strategies: Concepts and Companion Animal Model Outlook, Front Oncol. 11 (2021), 768692, [https://doi.](https://doi.org/10.3389/fonc.2021.768692) [org/10.3389/fonc.2021.768692.](https://doi.org/10.3389/fonc.2021.768692)
- [71] L.K. Kvols, Radiation sensitizers: a selective review of molecules targeting DNA and non-DNA targets, J. Nucl. Med 46 (Suppl 1) (2005), 187S-190S, 〈[https://](https://www.ncbi.nlm.nih.gov/pubmed/15653668) www.ncbi.nlm.nih.gov/pubmed/15653668〉.
- [72] Jonsson Comprehensive Cancer Center. Trastuzumab Deruxtecan Alone or in Combination With Anastrozole for the Treatment of Early Stage HER2 Low, Hormone Receptor Positive Breast Cancer. ClinicalTrials.gov Identifier: NCT04553770. https://clinicaltrials.gov/ct2/show/NCT04553770.
- [73] US Oncology Research. Trial of Ibrutinib Plus Trastuzumab in HER2-amplified Metastatic Breast Cancer. ClinicalTrials.gov Identifier: NCT03379428. 〈https:// clinicaltrials.gov/ct2/show/NCT03379428〉.
- [74] National Cancer Institute (NCI). Testing the Drug Atezolizumab or Placebo With Usual Therapy in First-Line HER2-Positive Metastatic Breast Cancer. ClinicalTrials.gov Identifier: NCT03199885. https://clinicaltrials.gov/ct2/show/ NCT03199885.
- [75] Sidney Kimmel Comprehensive Cancer Center at Johns Hopkins. Pertuzumab and Trastuzumab as Neoadjuvant Treatment in Patients With HER2-Positive Breast Cancer. ClinicalTrials.gov Identifier: NCT01937117. https://clinicaltrials.gov/ ct2/show/NCT01937117.
- [76] ETOP IBCSG Partners Foundation, BELIEF (Bevacizumab and ErLotinib In EGFR Mut+ NSCLC) (BELIEF). ClinicalTrials.gov Identifier: NCT01562028. https:// clinicaltrials.gov/ct2/show/NCT01562028.
- [77] Academic and Community Cancer Research United. Erlotinib Hydrochloride With or Without Bevacizumab in Treating Patients With Stage IV Non-small Cell Lung Cancer With Epidermal Growth Factor Receptor Mutations. ClinicalTrials.gov Identifier: NCT01532089. https://clinicaltrials.gov/ct2/show/NCT01532089.
- [78] Dana-Farber Cancer Institute, A Phase 2 Study of Osimertinib in Combination With Selumetinib in EGFR Inhibitor naïve Advanced EGFR Mutant Lung Cancer. ClinicalTrials.gov Identifier: NCT03392246. https://clinicaltrials.gov/ct2/show/ NCT03392246.
- [79] Memorial Sloan Kettering Cancer Center, Osimertinib Alone or With Chemotherapy for EGFR-Mutant Lung Cancers. ClinicalTrials.gov Identifier: NCT04410796. https://clinicaltrials.gov/ct2/show/NCT04410796.
- [80] Eli Lilly and Company. Study of Erbitux (Cetuximab) in Patients With Metastatic Colorectal Carcinoma. ClinicalTrials.gov Identifier: NCT00044863. https:// clinicaltrials.gov/ct2/show/NCT00044863.
- [81] NCIC Clinical Trials Group. Cetuximab + Best Supportive Care Compared With Best Supportive Care Alone in Metastatic Epidermal Growth Factor Receptor-Positive Colorectal Cancer. ClinicalTrials.gov Identifier: NCT00079066. https:// www.clinicaltrials.gov/ct2/show/NCT00079066.
- [82] M.D. Anderson Cancer Center. Panitumumab With or Without Trametinib in Treating Patients With Stage IV Colorectal Cancer. ClinicalTrials.gov Identifier: NCT03087071. https://clinicaltrials.gov/ct2/show/NCT0308707.
- [83] Northwestern University. Dual Epidermal Growth Factor Receptor Inhibition With Erlotinib and Panitumumab With or Without Chemotherapy for Advanced Colorectal Cancer. ClinicalTrials.gov Identifier: NCT00940316. https:// clinicaltrials.gov/ct2/show/NCT00940316.
- [84] AstraZeneca. Efficacy, Safety, Tolerability of Gefitinib as 1st Line in Caucasian Patients With EGFR Mutation Positive Advanced NSCLC (IFUM). ClinicalTrials. gov Identifier: NCT01203917. https://clinicaltrials.gov/ct2/show/ NCT01203917.
- [85] Dana-Farber Cancer Institute. Osimertinib and Gefitinib in EGFR Inhibitor naïve Advanced EGFR Mutant Lung Cancer. ClinicalTrials.gov Identifier: NCT03122717. https://www.clinicaltrials.gov/ct2/show/NCT03122717.
- [86] Bayer. A. Phase III Study of Sorafenib in Patients With Advanced Hepatocellular Carcinoma (SHARP). ClinicalTrials.gov Identifier: NCT00105443. https://www. clinicaltrials.gov/ct2/show/NCT00105443.
- [87] Bristol-Myers Squibb. An Investigational Immuno-therapy Study of Nivolumab Compared to Sorafenib as a First Treatment in Patients With Advanced Hepatocellular Carcinoma. ClinicalTrials.gov Identifier: NCT02576509. https:// clinicaltrials.gov/ct2/show/NCT02576509.
- [88] Pfizer. A. Clinical Trial Comparing Efficacy And Safety Of Sunitinib Versus Placebo For The Treatment Of Patients At High Risk Of Recurrent Renal Cell Cancer (S-TRAC). ClinicalTrials.gov Identifier: NCT00375674. https:// clinicaltrials.gov/ct2/show/NCT00375674.
- [89] Pfizer. Treatment Use Study With Sunitinib (SU011248) For Patients With Cytokine-Refractory Metastatic Renal Cell Carcinoma. ClinicalTrials.gov

Identifier: NCT00130897. https://www.clinicaltrials.gov/ct2/show/ NCT00130897.

- [90] GlaxoSmithKline. Safety and Efficacy of GW786034 (Pazopanib) In Metastatic Renal Cell Carcinoma. ClinicalTrials.gov Identifier: NCT00334282. https:// clinicaltrials.gov/ct2/show/NCT00334282.
- [91] Novartis Pharmaceuticals. Study of Efficacy, Safety, and Quality of Life of Pazopanib in Patients With Advanced and/or Metastatic Renal Cell Carcinoma After Prior Checkpoint Inhibitor Treatment (IO-PAZ). ClinicalTrials.gov Identifier: NCT03200717. https://clinicaltrials.gov/ct2/show/NCT03200717.
- [92] Hoffmann-La Roche. A Study of Bevacizumab in Combination With Standard of Care Treatment in Participants With Advanced Non-squamous Non-small Cell Lung Cancer (NSCLC). ClinicalTrials.gov Identifier: NCT01351415. https:// clinicaltrials.gov/ct2/show/NCT01351415.
- [93] Hoffmann-La Roche. A Study of the Addition of Avastin (Bevacizumab) to Carboplatin and Paclitaxel Therapy in Patients With Ovarian Cancer. ClinicalTrials.gov Identifier: NCT01239732. https://clinicaltrials.gov/ct2/show/ NCT01239732.
- [94] Pfizer. Temsirolimus Versus Sorafenib As Second-Line Therapy In Patients With Advanced RCC Who Have Failed First-Line Sunitinib (INTORSECT). ClinicalTrials.gov Identifier: NCT00474786. https://www.clinicaltrials.gov/ct2/ show/NCT00474786.
- [95] National Cancer Institute (NCI). Temsirolimus in Treating Patients With Metastatic or Locally Advanced Recurrent Endometrial Cancer. ClinicalTrials.gov Identifier: NCT00072176. https://clinicaltrials.gov/ct2/show/NCT00072176.
- [96] Novartis Pharmaceuticals. A Phase II Study of Everolimus in Combination With Exemestane Versus Everolimus Alone Versus Capecitabine in Advance Breast Cancer. (BOLERO-6). ClinicalTrials.gov Identifier: NCT01783444. https://www. clinicaltrials.gov/ct2/show/NCT01783444.
- [97] Novartis Pharmaceuticals. Efficacy and Safety of Everolimus (RAD001) Compared to Placebo in Patients With Advanced Neuroendocrine Tumors (RADIANT-3). ClinicalTrials.gov Identifier: NCT00510068. https://www.clinicaltrials.gov/ct2/ show/NCT00510068.
- [98] S.P. Gotoff, E. Amirmokri, E.J. Liebner, Ataxia telangiectasia. Neoplasia, untoward response to x-irradiation, and tuberous sclerosis, Am. J. Dis. Child 114 (6) (1967) 617-625, https://doi.org/10.1001/archpedi.1967.02090270073
- [99] F.P. Imray, C. Kidson, Perturbations of cell-cycle progression in gamma-irradiated ataxia telangiectasia and Huntington's disease cells detected by DNA flow cytometric analysis, Mutat. Res 112 (6) (1983) 369–382, [https://doi.org/](https://doi.org/10.1016/0167-8817(83)90030-5) [10.1016/0167-8817\(83\)90030-5.](https://doi.org/10.1016/0167-8817(83)90030-5)
- [100] N.J. Bentley, D.A. Holtzman, G. Flaggs, K.S. Keegan, A. DeMaggio, J.C. Ford, M. Hoekstra, A.M. Carr, The Schizosaccharomyces pombe rad3 checkpoint gene, EMBO J. 15 (23) (1996) 6641–6651. 〈[https://www.ncbi.nlm.nih.gov/pubmed/](https://www.ncbi.nlm.nih.gov/pubmed/8978690) [8978690](https://www.ncbi.nlm.nih.gov/pubmed/8978690)〉.
- [101] AstraZeneca. A. Study to Assess the Safety and Tolerability of AZD1390 Given With Radiation Therapy in Patients With Brain Cancer. ClinicalTrials.gov identifier: NCT03423628. 〈https://clinicaltrials.gov/ct2/show/NCT03423628〉.
- [102] XRad Therapeutics Inc. Trial of XRD-0394, a Kinase Inhibitor, in Combination With Palliative Radiotherapy in Advanced Cancer Patients. ClinicalTrials.gov identifier: NCT05002140. https://clinicaltrials.gov/ct2/show/NCT05002140.
- [103] M.T. Dillon, Z. Boylan, D. Smith, J. Guevara, K. Mohammed, C. Peckitt, M. Saunders, U. Banerji, G. Clack, S.A. Smith, J.F. Spicer, M.D. Forster, K. J. Harrington, PATRIOT: A phase I study to assess the tolerability, safety and biological effects of a specific ataxia telangiectasia and Rad3-related (ATR) inhibitor (AZD6738) as a single agent and in combination with palliative radiation therapy in patients with solid tumours, Clin. Transl. Radiat. Oncol. 12 (2018) 16–20, [https://doi.org/10.1016/j.ctro.2018.06.001.](https://doi.org/10.1016/j.ctro.2018.06.001)
- [104] University of Oxford. M6620 Plus Standard Treatment in Oesophageal and Other Cancer (CHARIOT).ClinicalTrials.govIdentifier:NCT03641547.https:// clinicaltrials.gov/ct2/show/NCT03641547.
- [105] National Cancer Institute (NCI). Testing the Addition of an Anti-cancer Drug, BAY 1895344, With Radiation Therapy to the Usual Pembrolizumab Treatment for Recurrent Head and Neck Cancer. ClinicalTrials.gov Identifier: NCT04576091. https://clinicaltrials.gov/ct2/show/NCT04576091.
- [106] Y. Wang, L. Yang, J. Zhang, M. Zhou, L. Shen, W. Deng, L. Liang, R. Hu, W. Yang, Y. Yao, H. Zhang, Z. Zhang, Radiosensitization by irinotecan is attributed to G2/M phase arrest, followed by enhanced apoptosis, probably through the ATM/Chk/ Cdc25C/Cdc2 pathway in p53-mutant colorectal cancer cells, Int J. Oncol. 53 (4) (2018) 1667–1680. https://doi.org/10.3892/ijo.2018.4514. /doi.org/10.3892/ijo.2018.451
- [107] Z. Wang, S.T. Lai, N.Y. Ma, Y. Deng, Y. Liu, D.P. Wei, J.D. Zhao, G.L. Jiang, Radiosensitization of metformin in pancreatic cancer cells via abrogating the G2 checkpoint and inhibiting DNA damage repair, Cancer Lett. 369 (1) (2015) 192–201, https://doi.org/10.1016/j.canlet.2015.08.015. (10.1016/j.canlet.2015.08.015.
- [108] R.S. Narayan, A. Gasol, P.L.G. Slangen, F.M.G. Cornelissen, T. Lagerweij, H. Veldman, R. Dik, J. van den Berg, B.J. Slotman, T. Wurdinger, D.A. Haas-Kogan, L.J.A. Stalpers, B.G. Baumert, B.A. Westerman, J. Theys, P. Sminia, Identification of MEK162 as a Radiosensitizer for the Treatment of Glioblastoma, Mol. Cancer Ther. 17 (2) (2018) 347–354, [https://doi.org/10.1158/1535-7163.](https://doi.org/10.1158/1535-7163.MCT-17-0480) MCT-17-0480
- [109] P. Raghavan, V. Tumati, L. Yu, N. Chan, N. Tomimatsu, S. Burma, R.G. Bristow, D. Saha, AZD5438, an inhibitor of Cdk1, 2, and 9, enhances the radiosensitivity of non-small cell lung carcinoma cells, Int J. Radiat. Oncol. Biol. Phys. 84 (4) (2012) e507–e514, [https://doi.org/10.1016/j.ijrobp.2012.05.035.](https://doi.org/10.1016/j.ijrobp.2012.05.035)
- [110] G.E. Rodland, K. Melhus, R. Generalov, S. Gilani, F. Bertoni, J. Dahle, R. G. Syljuasen, S. Patzke, The Dual Cell Cycle Kinase Inhibitor JNJ-7706621 Reverses Resistance to CD37-Targeted Radioimmunotherapy in Activated B Cell

S. Sriramulu et al.

Like Diffuse Large B Cell Lymphoma Cell Lines, Front Oncol. 9 (2019) 1301, [https://doi.org/10.3389/fonc.2019.01301.](https://doi.org/10.3389/fonc.2019.01301)

- [111] G. Feldmann, A. Mishra, S. Bisht, C. Karikari, I. Garrido-Laguna, Z. Rasheed, N. A. Ottenhof, T. Dadon, H. Alvarez, V. Fendrich, N.V. Rajeshkumar, W. Matsui, P. Brossart, M. Hidalgo, R. Bannerji, A. Maitra, B.D. Nelkin, Cyclin-dependent kinase inhibitor Dinaciclib (SCH727965) inhibits pancreatic cancer growth and progression in murine xenograft models, Cancer Biol. Ther. 12 (7) (2011) 598–609, [https://doi.org/10.4161/cbt.12.7.16475.](https://doi.org/10.4161/cbt.12.7.16475)
- [112] Piramal Enterprises Limited, Phase I Study of P276–00 in Patients With Advanced Refractory Neoplasms. ClinicalTrials.gov Identifier: NCT00408018. https:// clinicaltrials.gov/ct2/show/NCT00408018.
- [113] H. Galons, N. Oumata, O. Gloulou, L. Meijer, Cyclin-dependent kinase inhibitors closer to market launch, Expert Opin. Ther. Pat. 23 (8) (2013) 945–963, [https://](https://doi.org/10.1517/13543776.2013.789861) [doi.org/10.1517/13543776.2013.789861.](https://doi.org/10.1517/13543776.2013.789861)
- [114] R. Tibes, K.T. McDonagh, L. Lekakis, N. Frazer, S. Mohrland, B. Dawn, R. Garcia, K. Schroeder, V. Shanmugam, J. Carpten, D. Von Hoff, T.C. Shea, Phase I Study of the Novel Surivin and cdc2/CDK1 Inhibitor Terameprocol in Patients with Advanced Leukemias, Blood 114 (22) (2009), [https://doi.org/10.1182/blood.](https://doi.org/10.1182/blood.v114.22.1039.1039) [v114.22.1039.1039](https://doi.org/10.1182/blood.v114.22.1039.1039).
- [115] [C.C. Zhang, S. Kephart, I. McAlpine, J. Nonomiya, J. Higgins, M.E. Arango,](http://refhub.elsevier.com/S0753-3322(22)01515-3/sbref84) Z. Yan, D. Knighton, R.A. Ferre, J. Tikhe, G. Verkhivker, M. Xu, W. Romin [C. Palmer, J. Park, S. Reich, W. Chong, L. Li, G. Los, C. Lewis, AG-024322 is a](http://refhub.elsevier.com/S0753-3322(22)01515-3/sbref84) [potent and selective multi-targeted CDK inhibitor with broad spectrum anti](http://refhub.elsevier.com/S0753-3322(22)01515-3/sbref84)[proliferative activity, Cancer Res. 65 \(9_Supplement\) \(2005\), 1045-1045](http://refhub.elsevier.com/S0753-3322(22)01515-3/sbref84).
- [116] A.G. Murphy, M. Zahurak, M. Shah, C.D. Weekes, A. Hansen, L.L. Siu, A. Spreafico, N. LoConte, N.M. Anders, T. Miles, M.A. Rudek, L.A. Doyle, B. Nelkin, A. Maitra, N.S. Azad, E.-S. Team, A Phase I Study of Dinaciclib in Combination With MK-2206 in Patients With Advanced Pancreatic Cancer, Clin. Transl. Sci. 13 (6) (2020) 1178–1188, [https://doi.org/10.1111/cts.12802.](https://doi.org/10.1111/cts.12802)
- [117] R.D. Carvajal, A. Tse, M.A. Shah, R.A. Lefkowitz, M. Gonen, L. Gilman-Rosen, J. Kortmansky, D.P. Kelsen, G.K. Schwartz, E.M. O'Reilly, A phase II study of flavopiridol (Alvocidib) in combination with docetaxel in refractory, metastatic pancreatic cancer, Pancreatology 9 (4) (2009) 404–409, [https://doi.org/](https://doi.org/10.1159/000187135) [10.1159/000187135.](https://doi.org/10.1159/000187135)
- [118] S. Aspeslagh, K. Shailubhai, R. Bahleda, A. Gazzah, A. Varga, A. Hollebecque, C. Massard, A. Spreafico, M. Reni, J.C. Soria, Phase I dose-escalation study of milciclib in combination with gemcitabine in patients with refractory solid tumors, Cancer Chemother. Pharm. 79 (6) (2017) 1257–1265, [https://doi.org/](https://doi.org/10.1007/s00280-017-3303-z) [10.1007/s00280-017-3303-z.](https://doi.org/10.1007/s00280-017-3303-z)
- [119] J. Cicenas, M. Valius, The CDK inhibitors in cancer research and therapy, J. Cancer Res Clin. Oncol. 137 (10) (2011) 1409–1418, [https://doi.org/10.1007/](https://doi.org/10.1007/s00432-011-1039-4) [s00432-011-1039-4](https://doi.org/10.1007/s00432-011-1039-4).
- [120] T. VanArsdale, C. Boshoff, K.T. Arndt, R.T. Abraham, Molecular Pathways: Targeting the Cyclin D-CDK4/6 Axis for Cancer Treatment, Clin. Cancer Res 21 (13) (2015) 2905–2910, <https://doi.org/10.1158/1078-0432.CCR-14-0816>.
- [121] S. Goel, M.J. DeCristo, S.S. McAllister, J.J. Zhao, CDK4/6 Inhibition in Cancer: Beyond Cell Cycle Arrest, Trends Cell Biol. 28 (11) (2018) 911–925, [https://doi.](https://doi.org/10.1016/j.tcb.2018.07.002) [org/10.1016/j.tcb.2018.07.002.](https://doi.org/10.1016/j.tcb.2018.07.002)
- [122] Y. Yang, J. Luo, X. Chen, Z. Yang, X. Mei, J. Ma, Z. Zhang, X. Guo, X. Yu, CDK4/6 inhibitors: a novel strategy for tumor radiosensitization, J. Exp. Clin. Cancer Res 39 (1) (2020) 188, [https://doi.org/10.1186/s13046-020-01693-w.](https://doi.org/10.1186/s13046-020-01693-w)
- [123] N. Sobhani, A. D'Angelo, M. Pittacolo, G. Roviello, A. Miccoli, S.P. Corona, O. Bernocchi, D. Generali, T. Otto, Updates on the CDK4/6 Inhibitory Strategy and Combinations in Breast Cancer, Cells 8 (4) (2019), [https://doi.org/10.3390/](https://doi.org/10.3390/cells8040321) [cells8040321](https://doi.org/10.3390/cells8040321).
- [124] R.S. Finn, J.P. Crown, I. Lang, K. Boer, I.M. Bondarenko, S.O. Kulyk, J. Ettl, R. Patel, T. Pinter, M. Schmidt, Y. Shparyk, A.R. Thummala, N.L. Voytko, C. Fowst, X. Huang, S.T. Kim, S. Randolph, D.J. Slamon, The cyclin-dependent kinase 4/6 inhibitor palbociclib in combination with letrozole versus letrozole alone as first-line treatment of oestrogen receptor-positive, HER2-negative, advanced breast cancer (PALOMA-1/TRIO-18): a randomised phase 2 study, Lancet Oncol. 16 (1) (2015) 25–35, [https://doi.org/10.1016/S1470-2045\(14\)](https://doi.org/10.1016/S1470-2045(14)71159-3) [71159-3](https://doi.org/10.1016/S1470-2045(14)71159-3).
- [125] C.Y. Huang, F.S. Hsieh, C.Y. Wang, L.J. Chen, S.S. Chang, M.H. Tsai, M.H. Hung, C.W. Kuo, C.T. Shih, T.I. Chao, K.F. Chen, Palbociclib enhances radiosensitivity of hepatocellular carcinoma and cholangiocarcinoma via inhibiting ataxia telangiectasia-mutated kinase-mediated DNA damage response, Eur. J. Cancer 102 (2018) 10–22, [https://doi.org/10.1016/j.ejca.2018.07.010.](https://doi.org/10.1016/j.ejca.2018.07.010)
- [126] S. Naz, A. Sowers, R. Choudhuri, M. Wissler, J. Gamson, A. Mathias, J.A. Cook, J. B. Mitchell, Abemaciclib, a Selective CDK4/6 Inhibitor, Enhances the Radiosensitivity of Non-Small Cell Lung Cancer In Vitro and In Vivo, Clin. Cancer Res 24 (16) (2018) 3994–4005, [https://doi.org/10.1158/1078-0432.CCR-17-](https://doi.org/10.1158/1078-0432.CCR-17-3575) [3575](https://doi.org/10.1158/1078-0432.CCR-17-3575).
- [127] R. Hashizume, A. Zhang, S. Mueller, M.D. Prados, R.R. Lulla, S. Goldman, A. M. Saratsis, A.P. Mazar, A.H. Stegh, S.Y. Cheng, C. Horbinski, D.A. Haas-Kogan, J. N. Sarkaria, T. Waldman, C.D. James, Inhibition of DNA damage repair by the CDK4/6 inhibitor palbociclib delays irradiated intracranial atypical teratoid rhabdoid tumor and glioblastoma xenograft regrowth, Neuro Oncol. 18 (11) (2016) 1519–1528, <https://doi.org/10.1093/neuonc/now106>.
- [128] Children's Hospital Medical Center, Cincinnati. A Phase I/II Study of Ribociclib,a CDK4/6 Inhibitor, Following Radiation Therapy. ClinicalTrials.gov Identifier: NCT02607124. https://clinicaltrials.gov/ct2/show/NCT02607124.
- [129] Children's Hospital Medical Center, Cincinnati. A Study of Ribociclib and Everolimus Following Radiation Therapy in Children With Newly Diagnosed Nonbiopsied Diffuse Pontine Gliomas (DIPG) and RB+ Biopsied DIPG and High Grade

Gliomas (HGG). ClinicalTrials.govIdentifier:NCT03355794.https://www. clinicaltrials.gov/ct2/show/NCT03355794.

- [130] A. Sancar, L.A. Lindsey-Boltz, K. Unsal-Kacmaz, S. Linn, Molecular mechanisms of mammalian DNA repair and the DNA damage checkpoints, Annu Rev. Biochem 73 (2004) 39–85, [https://doi.org/10.1146/annurev.biochem.73.011303.073723.](https://doi.org/10.1146/annurev.biochem.73.011303.073723)
- [131] Z. Chen, Z. Xiao, W.Z. Gu, J. Xue, M.H. Bui, P. Kovar, G. Li, G. Wang, Z.F. Tao, Y. Tong, N.H. Lin, H.L. Sham, J.Y. Wang, T.J. Sowin, S.H. Rosenberg, H. Zhang, Selective Chk1 inhibitors differentially sensitize p53-deficient cancer cells to cancer therapeutics, Int J. Cancer 119 (12) (2006) 2784–2794, [https://doi.org/](https://doi.org/10.1002/ijc.22198) [10.1002/ijc.22198.](https://doi.org/10.1002/ijc.22198)
- [132] Eli Lilly and Company. A Study of Prexasertib (LY2606368) With Chemotherapy and Radiation in Participants With Head and Neck Cancer. ClinicalTrials.gov Identifier: NCT02555644. https://clinicaltrials.gov/ct2/show/study/ NCT02555644.
- [133] R.T. Bunch, A. Eastman, Enhancement of cisplatin-induced cytotoxicity by 7hydroxystaurosporine (UCN-01), a new G2-checkpoint inhibitor, Clin. Cancer Res 2 (5) (1996) 791-797. \https://www.ncbi.nlm.nih.gov/pubmed/981623
- [134] Pfizer. PF-00477736 Is Being Studied In Advanced Solid Tumors In Combination With Chemotherapy With Gemcitabine. ClinicalTrials.gov Identifier: NCT00437203. https://clinicaltrials.gov/ct2/show/NCT00437203.
- [135] J.D. Guster, S.V. Weissleder, C.J. Busch, M. Kriegs, C. Petersen, R. Knecht, E. Dikomey, T. Rieckmann, The inhibition of PARP but not EGFR results in the radiosensitization of HPV/p16-positive HNSCC cell lines, Radio. Oncol. 113 (3) (2014) 345–351, <https://doi.org/10.1016/j.radonc.2014.10.011>.
- [136] R. Patel, H.E. Barker, J. Kyula, M. McLaughlin, M.T. Dillon, U. Schick, H. Hafsi, A. Thompson, V. Khoo, K. Harrington, S. Zaidi, An orally bioavailable Chk1 inhibitor, CCT244747, sensitizes bladder and head and neck cancer cell lines to radiation, Radio. Oncol. 122 (3) (2017) 470–475, [https://doi.org/10.1016/j.](https://doi.org/10.1016/j.radonc.2016.12.026) [radonc.2016.12.026.](https://doi.org/10.1016/j.radonc.2016.12.026)
- [137] L. Booth, J. Roberts, A. Poklepovic, P. Dent, The CHK1 inhibitor SRA737 synergizes with PARP1 inhibitors to kill carcinoma cells, Cancer Biol. Ther. 19 (9) (2018) 786–796, <https://doi.org/10.1080/15384047.2018.1472189>.
- [138] L. Gorecki, M. Andrs, J. Korabecny, Clinical Candidates Targeting the ATR-CHK1-WEE1 Axis in Cancer, Cancers (Basel) 13 (4) (2021), [https://doi.org/10.3390/](https://doi.org/10.3390/cancers13040795) ancers130407
- [139] A.J. Davis, L. Chi, S. So, K.J. Lee, E. Mori, K. Fattah, J. Yang, D.J. Chen, BRCA1 modulates the autophosphorylation status of DNA-PKcs in S phase of the cell cycle, Nucleic Acids Res 42 (18) (2014) 11487–11501, [https://doi.org/10.1093/](https://doi.org/10.1093/nar/gku824) ar/øku824
- [140] Y. Zhou, T.T. Paull, DNA-dependent protein kinase regulates DNA end resection in concert with Mre11-Rad50-Nbs1 (MRN) and ataxia telangiectasia-mutated (ATM, J. Biol. Chem. 288 (52) (2013) 37112–37125, [https://doi.org/10.1074/](https://doi.org/10.1074/jbc.M113.514398) c.M113.514398.
- [141] J.A. Neal, V. Dang, P. Douglas, M.S. Wold, S.P. Lees-Miller, K. Meek, Inhibition of homologous recombination by DNA-dependent protein kinase requires kinase activity, is titratable, and is modulated by autophosphorylation, Mol. Cell Biol. 31 (8) (2011) 1719–1733, [https://doi.org/10.1128/MCB.01298-10.](https://doi.org/10.1128/MCB.01298-10)
- [142] B. Huang, Z.F. Shang, B. Li, Y. Wang, X.D. Liu, S.M. Zhang, H. Guan, W.Q. Rang, J. A. Hu, P.K. Zhou, DNA-PKcs associates with PLK1 and is involved in proper chromosome segregation and cytokinesis, J. Cell Biochem 115 (6) (2014) 1077–1088, <https://doi.org/10.1002/jcb.24703>.
- [143] W.Z. Tu, B. Li, B. Huang, Y. Wang, X.D. Liu, H. Guan, S.M. Zhang, Y. Tang, W. Q. Rang, P.K. Zhou, gammaH2AX foci formation in the absence of DNA damage: mitotic H2AX phosphorylation is mediated by the DNA-PKcs/CHK2 pathway, FEBS Lett. 587 (21) (2013) 3437–3443, [https://doi.org/10.1016/j.](https://doi.org/10.1016/j.febslet.2013.08.028) ebslet.2013.08.028
- [144] K.J. Lee, Y.F. Lin, H.Y. Chou, H. Yajima, K.R. Fattah, S.C. Lee, B.P. Chen, Involvement of DNA-dependent protein kinase in normal cell cycle progression through mitosis, J. Biol. Chem. 286 (14) (2011) 12796–12802, [https://doi.org/](https://doi.org/10.1074/jbc.M110.212969) [10.1074/jbc.M110.212969](https://doi.org/10.1074/jbc.M110.212969).
- [145] Z.F. Shang, B. Huang, Q.Z. Xu, S.M. Zhang, R. Fan, X.D. Liu, Y. Wang, P.K. Zhou, Inactivation of DNA-dependent protein kinase leads to spindle disruption and mitotic catastrophe with attenuated checkpoint protein 2 Phosphorylation in response to DNA damage, Cancer Res 70 (9) (2010) 3657–3666, [https://doi.org/](https://doi.org/10.1158/0008-5472.CAN-09-3362) [10.1158/0008-5472.CAN-09-3362](https://doi.org/10.1158/0008-5472.CAN-09-3362).
- [146] B.L. Ruis, K.R. Fattah, E.A. Hendrickson, The catalytic subunit of DNA-dependent protein kinase regulates proliferation, telomere length, and genomic stability in human somatic cells, Mol. Cell Biol. 28 (20) (2008) 6182-6195, https://doi. [10.1128/MCB.00355-08.](https://doi.org/10.1128/MCB.00355-08)
- [147] Y. Li, G. Zhou, I.G. Bruno, N. Zhang, S. Sho, E. Tedone, T.P. Lai, J.P. Cooke, J. W. Shay, Transient introduction of human telomerase mRNA improves hallmarks of progeria cells, Aging Cell 18 (4) (2019), e12979, [https://doi.org/10.1111/](https://doi.org/10.1111/acel.12979) [acel.12979.](https://doi.org/10.1111/acel.12979)
- [148] X. Cui, Y. Yu, S. Gupta, Y.M. Cho, S.P. Lees-Miller, K. Meek, Autophosphorylation of DNA-dependent protein kinase regulates DNA end processing and may also alter double-strand break repair pathway choice, Mol. Cell Biol. 25 (24) (2005) 10842–10852, [https://doi.org/10.1128/MCB.25.24.10842-10852.2005.](https://doi.org/10.1128/MCB.25.24.10842-10852.2005)
- [149] S. Zhang, S. Matsunaga, Y.F. Lin, B. Sishc, Z. Shang, J. Sui, H.Y. Shih, Y. Zhao, O. Foreman, M.D. Story, D.J. Chen, B.P. Chen, Spontaneous tumor development in bone marrow-rescued DNA-PKcs(3A/3A) mice due to dysfunction of telomere leading strand deprotection, Oncogene 35 (30) (2016) 3909–3918, [https://doi.](https://doi.org/10.1038/onc.2015.459) [org/10.1038/onc.2015.459](https://doi.org/10.1038/onc.2015.459).
- [150] B.S. Lee, E.J. Gapud, S. Zhang, Y. Dorsett, A. Bredemeyer, R. George, E. Callen, J. A. Daniel, O. Osipovich, E.M. Oltz, C.H. Bassing, A. Nussenzweig, S. Lees-Miller, M. Hammel, B.P. Chen, B.P. Sleckman, Functional intersection of ATM and DNAdependent protein kinase catalytic subunit in coding end joining during V(D)J

recombination, Mol. Cell Biol. 33 (18) (2013) 3568–3579, [https://doi.org/](https://doi.org/10.1128/MCB.00308-13) [10.1128/MCB.00308-13.](https://doi.org/10.1128/MCB.00308-13)

- [151] P. Douglas, S. Gupta, N. Morrice, K. Meek, S.P. Lees-Miller, DNA-PK-dependent phosphorylation of Ku70/80 is not required for non-homologous end joining, DNA Repair (Amst.) 4 (9) (2005) 1006–1018, [https://doi.org/10.1016/j.](https://doi.org/10.1016/j.dnarep.2005.05.003) [dnarep.2005.05.003](https://doi.org/10.1016/j.dnarep.2005.05.003).
- [152] I.S. Mohiuddin, M.H. Kang, DNA-PK as an Emerging Therapeutic Target in Cancer, Front Oncol. 9 (2019) 635, <https://doi.org/10.3389/fonc.2019.00635>.
- [153] D. Davidson, L. Amrein, L. Panasci, R. Aloyz, Small Molecules, Inhibitors of DNA-PK, Targeting DNA Repair, and Beyond, Front Pharm. 4 (2013) 5, [https://doi.org/](https://doi.org/10.3389/fphar.2013.00005) [10.3389/fphar.2013.00005](https://doi.org/10.3389/fphar.2013.00005).
- [154] S.J. Collis, T.L. DeWeese, P.A. Jeggo, A.R. Parker, The life and death of DNA-PK, Oncogene 24 (6) (2005) 949-961, https://doi.org/10.1038/sj.onc.1208
- [155] R.L. Gurung, H.K. Lim, S. Venkatesan, P.S. Lee, M.P. Hande, Targeting DNA-PKcs and telomerase in brain tumour cells, Mol. Cancer 13 (2014) 232, [https://doi.](https://doi.org/10.1186/1476-4598-13-232) [org/10.1186/1476-4598-13-232](https://doi.org/10.1186/1476-4598-13-232).
- [156] S. Durant, P. Karran, Vanillins–a novel family of DNA-PK inhibitors, Nucleic Acids Res 31 (19) (2003) 5501–5512, <https://doi.org/10.1093/nar/gkg753>.
- [157] E.T. Shinohara, L. Geng, J. Tan, H. Chen, Y. Shir, E. Edwards, J. Halbrook, E. A. Kesicki, A. Kashishian, D.E. Hallahan, DNA-dependent protein kinase is a molecular target for the development of noncytotoxic radiation-sensitizing drugs, Cancer Res 65 (12) (2005) 4987–4992, [https://doi.org/10.1158/0008-5472.](https://doi.org/10.1158/0008-5472.CAN-04-4250) N-04-4250
- [158] A.J. Khan, S.M. Misenko, A. Thandoni, D. Schiff, S.R. Jhawar, S.F. Bunting, B. G. Haffty, VX-984 is a selective inhibitor of non-homologous end joining, with possible preferential activity in transformed cells, Oncotarget 9 (40) (2018) 25833–25841, [https://doi.org/10.18632/oncotarget.25383.](https://doi.org/10.18632/oncotarget.25383)
- [159] J.H.L. Fok, A. Ramos-Montoya, M. Vazquez-Chantada, P.W.G. Wijnhoven, V. Follia, N. James, P.M. Farrington, A. Karmokar, S.E. Willis, J. Cairns, J. Nikkila, D. Beattie, G.M. Lamont, M.R.V. Finlay, J. Wilson, A. Smith, L.O. O'Connor, S. Ling, S.E. Fawell, E.B. Cadogan, AZD7648 is a potent and selective DNA-PK inhibitor that enhances radiation, chemotherapy and olaparib activity, Nat. Commun. 10 (1) (2019) 5065, <https://doi.org/10.1038/s41467-019-12836-9>.
- [160] R. Lamb, M. Fiorillo, A. Chadwick, B. Ozsvari, K.J. Reeves, D.L. Smith, R. B. Clarke, S.J. Howell, A.R. Cappello, U.E. Martinez-Outschoorn, M. Peiris-Pages, F. Sotgia, M.P. Lisanti, Doxycycline down-regulates DNA-PK and radiosensitizes tumor initiating cells: Implications for more effective radiation therapy, Oncotarget 6 (16) (2015) 14005–14025, [https://doi.org/10.18632/](https://doi.org/10.18632/oncotarget.4159) cotarget.4159.
- [161] F.T. Zenke, A. Zimmermann, C. Sirrenberg, H. Dahmen, V. Kirkin, U. Pehl, T. Grombacher, C. Wilm, T. Fuchss, C. Amendt, L.T. Vassilev, A. Blaukat, Pharmacologic Inhibitor of DNA-PK, M3814, Potentiates Radiotherapy and Regresses Human Tumors in Mouse Models, Mol. Cancer Ther. 19 (5) (2020) 1091–1101, [https://doi.org/10.1158/1535-7163.MCT-19-0734.](https://doi.org/10.1158/1535-7163.MCT-19-0734)
- [162] B. Vormoor, Y.T. Schlosser, H. Blair, A. Sharma, S. Wilkinson, D.R. Newell, N. Curtin, Sensitizing Ewing sarcoma to chemo- and radiotherapy by inhibition of the DNA-repair enzymes DNA protein kinase (DNA-PK) and poly-ADP-ribose polymerase (PARP) 1/2, Oncotarget 8 (69) (2017) 113418–113430, [https://doi.](https://doi.org/10.18632/oncotarget.21300) [org/10.18632/oncotarget.21300](https://doi.org/10.18632/oncotarget.21300).
- [163] T. Hirai, S. Saito, H. Fujimori, K. Matsushita, T. Nishio, R. Okayasu, M. Masutani, Radiosensitization by PARP inhibition to proton beam irradiation in cancer cells, Biochem Biophys. Res Commun. 478 (1) (2016) 234–240, [https://doi.org/](https://doi.org/10.1016/j.bbrc.2016.07.062) [10.1016/j.bbrc.2016.07.062](https://doi.org/10.1016/j.bbrc.2016.07.062).
- [164] S.A. Jannetti, G. Carlucci, B. Carney, S. Kossatz, L. Shenker, L.M. Carter, B. Salinas, C. Brand, A. Sadique, P.L. Donabedian, K.M. Cunanan, M. Gonen, V. Ponomarev, B.M. Zeglis, M.M. Souweidane, J.S. Lewis, W.A. Weber, J. L. Humm, T. Reiner, PARP-1-Targeted Radiotherapy in Mouse Models of Glioblastoma, J. Nucl. Med 59 (8) (2018) 1225–1233, [https://doi.org/10.2967/](https://doi.org/10.2967/jnumed.117.205054) umed.117.205054.
- [165] H. Ryu, H.J. Kim, J.Y. Song, S.G. Hwang, J.S. Kim, J. Kim, T.H.N. Bui, H.K. Choi, J. Ahn, A Small Compound KJ-28d Enhances the Sensitivity of Non-Small Cell Lung Cancer to Radio- and Chemotherapy, Int J. Mol. Sci. 20 (23) (2019), [https://](https://doi.org/10.3390/ijms20236026) [doi.org/10.3390/ijms20236026.](https://doi.org/10.3390/ijms20236026)
- [166] C. Guillot, V. Favaudon, Z. Herceg, C. Sagne, S. Sauvaigo, P. Merle, J. Hall, I. Chemin, PARP inhibition and the radiosensitizing effects of the PARP inhibitor ABT-888 in in vitro hepatocellular carcinoma models, BMC Cancer 14 (2014) 603,<https://doi.org/10.1186/1471-2407-14-603>.
- [167] L. Wang, K.A. Mason, K.K. Ang, T. Buchholz, D. Valdecanas, A. Mathur, C. Buser-Doepner, C. Toniatti, L. Milas, MK-4827, a PARP-1/-2 inhibitor, strongly enhances response of human lung and breast cancer xenografts to radiation, Invest N. Drugs 30 (6) (2012) 2113–2120, [https://doi.org/10.1007/s10637-011-](https://doi.org/10.1007/s10637-011-9770-x) [9770-x](https://doi.org/10.1007/s10637-011-9770-x).
- [168] E.C. Bourton, P.A. Ahorner, P.N. Plowman, S.A. Zahir, H. Al-Ali, C.N. Parris, The PARP-1 inhibitor Olaparib suppresses BRCA1 protein levels, increases apoptosis and causes radiation hypersensitivity in $\text{BRCA1}(+/-)$ lymphoblastoid cells, J. Cancer 8 (19) (2017) 4048-4056, https://doi.org/10.7150/jca.2133
- [169] M. Alotaibi, K. Sharma, T. Saleh, L.F. Povirk, E.A. Hendrickson, D.A. Gewirtz, Radiosensitization by PARP Inhibition in DNA Repair Proficient and Deficient Tumor Cells: Proliferative Recovery in Senescent Cells, Radiat. Res 185 (3) (2016) 229–245, [https://doi.org/10.1667/RR14202.1.](https://doi.org/10.1667/RR14202.1)
- [170] M.R. Mirza, B.J. Monk, J. Herrstedt, A.M. Oza, S. Mahner, A. Redondo, M. Fabbro, J.A. Ledermann, D. Lorusso, I. Vergote, N.E. Ben-Baruch, C. Marth, R. Madry, R. D. Christensen, J.S. Berek, A. Dorum, A.V. Tinker, A. du Bois, A. Gonzalez-Martin, E.-O.N. Investigators, Niraparib Maintenance Therapy in Platinum-Sensitive, Recurrent Ovarian Cancer, N. Engl. J. Med 375 (22) (2016) 2154–2164, [https://](https://doi.org/10.1056/NEJMoa1611310) doi.org/10.1056/NEJMoa1611310.
- [171] Centre Francois Baclesse. Phase I/IIa Study of Concomitant Radiotherapy With Olaparib and Temozolomide in Unresectable High Grade Gliomas Patients (OLA-TMZ-RTE-01). ClinicalTrials.govIdentifier:NCT03212742.https://clinicaltrials. gov/ct2/show/NCT03212742.
- [172] Massachusetts General Hospital. Radiation, Immunotherapy and PARP Inhibitor in Triple Negative Breast Cancer (NADiR). ClinicalTrials.gov Identifier: NCT04837209. https://clinicaltrials.gov/ct2/show/NCT04837209.
- [173] National Cancer Institute (NCI). Veliparib, Radiation Therapy, and Temozolomide in Treating Patients With Newly Diagnosed Malignant Glioma Without H3 K27M or BRAFV600 Mutations. ClinicalTrials.gov Identifier: NCT03581292. https:// clinicaltrials.gov/ct2/show/NCT03581292.
- [174] L. Sun, E. Moore, R. Berman, P.E. Clavijo, A. Saleh, Z. Chen, C. Van Waes J. Davies, J. Friedman, C.T. Allen, WEE1 kinase inhibition reverses G2/M cell cycle checkpoint activation to sensitize cancer cells to immunotherapy, Oncoimmunology 7 (10) (2018), e1488359, [https://doi.org/10.1080/](https://doi.org/10.1080/2162402X.2018.1488359) 2162402X.2018.14883
- [175] Y.Y. Lee, Y.J. Cho, S.W. Shin, C. Choi, J.Y. Ryu, H.K. Jeon, J.J. Choi, J.R. Hwang, C.H. Choi, T.J. Kim, B.G. Kim, D.S. Bae, W. Park, J.W. Lee, Anti-Tumor Effects of Wee1 Kinase Inhibitor with Radiotherapy in Human Cervical Cancer, Sci. Rep. 9 (1) (2019) 15394,<https://doi.org/10.1038/s41598-019-51959-3>.
- [176] R. Havelek, J. Cmielova, K. Kralovec, L. Bruckova, Z. Bilkova, I. Fousova, Z. Sinkorova, J. Vavrova, M. Rezacova, Specific inhibition of Wee1 kinase and Rad51 recombinase: a strategy to enhance the sensitivity of leukemic T-cells to ionizing radiation-induced DNA double-strand breaks, Biochem Biophys. Res Commun. 453 (3) (2014) 569–575, [https://doi.org/10.1016/j.bbrc.2014.09.123.](https://doi.org/10.1016/j.bbrc.2014.09.123)
- [177] J. PosthumaDeBoer, T. Wurdinger, H.C. Graat, V.W. van Beusechem, M.N. Helder, B.J. van Royen, G.J. Kaspers, WEE1 inhibition sensitizes osteosarcoma to radiotherapy, BMC Cancer 11 (2011) 156, [https://doi.org/10.1186/1471-2407-](https://doi.org/10.1186/1471-2407-11-156)
- [11-156.](https://doi.org/10.1186/1471-2407-11-156) [178] C.J. Matheson, S. Venkataraman, V. Amani, P.S. Harris, D.S. Backos, A. M. Donson, M.F. Wempe, N.K. Foreman, R. Vibhakar, P. Reigan, A WEE1 Inhibitor Analog of AZD1775 Maintains Synergy with Cisplatin and Demonstrates Reduced Single-Agent Cytotoxicity in Medulloblastoma Cells, ACS Chem. Biol. 11 (4) (2016) 921–930, <https://doi.org/10.1021/acschembio.5b00725>.
- [179] G. Chen, B. Zhang, H. Xu, Y. Sun, Y. Shi, Y. Luo, H. Jia, F. Wang, Suppression of Sirt1 sensitizes lung cancer cells to WEE1 inhibitor MK-1775-induced DNA damage and apoptosis, Oncogene 36 (50) (2017) 6863-6872, https://doi.org/ [10.1038/onc.2017.297.](https://doi.org/10.1038/onc.2017.297)
- [180] B.S. Chera, S.H. Sheth, S.A. Patel, D. Goldin, K.E. Douglas, R.L. Green, C.J. Shen, G.P. Gupta, D.T. Moore, J.E. Grilley Olson, J.M. Weiss, Phase 1 trial of adavosertib (AZD1775) in combination with concurrent radiation and cisplatin for intermediate-risk and high-risk head and neck squamous cell carcinoma, Cancer 127 (23) (2021) 4447–4454, [https://doi.org/10.1002/cncr.33789.](https://doi.org/10.1002/cncr.33789)
- [181] University of Michigan Rogel Cancer Center. Dose Escalation Trial of AZD1775 and Gemcitabine (+Radiation) for Unresectable Adenocarcinoma of the Pancreas. ClinicalTrials.govIdentifier:NCT02037230.https://clinicaltrials.gov/ct2/show/ NCT02037230.
- [182] G.B. Mills, R. Schmandt, M. McGill, A. Amendola, M. Hill, K. Jacobs, C. May, A. M. Rodricks, S. Campbell, D. Hogg, Expression of TTK, a novel human protein kinase, is associated with cell proliferation, J. Biol. Chem. 267 (22) (1992) 16000–16006. 〈<https://www.ncbi.nlm.nih.gov/pubmed/1639825>〉.
- [183] J. Daniel, J. Coulter, J.H. Woo, K. Wilsbach, E. Gabrielson, High levels of the Mps1 checkpoint protein are protective of aneuploidy in breast cancer cells, Proc. Natl. Acad. Sci. USA 108 (13) (2011) 5384–5389, [https://doi.org/10.1073/](https://doi.org/10.1073/pnas.1007645108) [pnas.1007645108](https://doi.org/10.1073/pnas.1007645108).
- [184] S. Simon Serrano, W. Sime, Y. Abassi, R. Daams, R. Massoumi, M. Jemaa, Inhibition of mitotic kinase Mps1 promotes cell death in neuroblastoma, Sci. Rep. 10 (1) (2020) 11997, <https://doi.org/10.1038/s41598-020-68829-y>.
- [185] B.C. Chandler, L. Moubadder, C.L. Ritter, M. Liu, M. Cameron, K. Wilder-Romans, A. Zhang, A.M. Pesch, A.R. Michmerhuizen, N. Hirsh, M. Androsiglio, T. Ward, E. Olsen, Y.S. Niknafs, S. Merajver, D.G. Thomas, P.H. Brown, T.S. Lawrence, S. Nyati, C. Speers, TTK inhibition radiosensitizes basal-like breast cancer through impaired homologous recombination, J. Clin. Invest 130 (2) (2020) 958–973, s://doi.org/10.1172/JCI130435.
- [186] U.B. Maachani, T. Kramp, R. Hanson, S. Zhao, O. Celiku, U. Shankavaram, R. Colombo, N.J. Caplen, K. Camphausen, A. Tandle, Targeting MPS1 Enhances Radiosensitization of Human Glioblastoma by Modulating DNA Repair Proteins, Mol. Cancer Res 13 (5) (2015) 852-862, https://doi.org/10.1158/1541-7 [MCR-14-0462-T.](https://doi.org/10.1158/1541-7786.MCR-14-0462-T)
- [187] B.A. Tannous, M. Kerami, P.M. Van der Stoop, N. Kwiatkowski, J. Wang, W. Zhou, A.F. Kessler, G. Lewandrowski, L. Hiddingh, N. Sol, T. Lagerweij, L. Wedekind, J. M. Niers, M. Barazas, R.J. Nilsson, D. Geerts, P.C. De Witt Hamer, C. Hagemann, W.P. Vandertop, T. Wurdinger, Effects of the selective MPS1 inhibitor MPS1-IN-3 on glioblastoma sensitivity to antimitotic drugs, J. Natl. Cancer Inst. 105 (17) (2013) 1322–1331, [https://doi.org/10.1093/jnci/djt168.](https://doi.org/10.1093/jnci/djt168)
- [188] Bayer. Phase I Study of Oral BAY 1217389 in Combination With Intravenous Paclitaxel. ClinicalTrials.govIdentifier:NCT02366949.https://www.clinicaltrials. gov/ct2/show/NCT02366949.
- [189] Bayer. Phase I Dose Escalation of Oral BAY1161909 in Combination With Intravenous Paclitaxel.ClinicalTrials.govIdentifier:NCT02138812.〈https://www. clinicaltrials.gov/ct2/show/NCT02138812〉.
- [190] Boston Pharmaceuticals. Study of Paclitaxel in Combination With BOS172722 in Patients With Advanced Nonhaematologic Malignancies. ClinicalTrials.gov Identifier: NCT03328494. https://clinicaltrials.gov/ct2/show/NCT03328494.
- [191] L.J. Mo, M. Song, Q.H. Huang, H. Guan, X.D. Liu, D.F. Xie, B. Huang, R.X. Huang, P.K. Zhou, Exosome-packaged miR-1246 contributes to bystander DNA damage

S. Sriramulu et al.

by targeting LIG4, Br. J. Cancer 119 (4) (2018) 492–502, [https://doi.org/](https://doi.org/10.1038/s41416-018-0192-9) [10.1038/s41416-018-0192-9.](https://doi.org/10.1038/s41416-018-0192-9)

- [192] G.J. Williams, M. Hammel, S.K. Radhakrishnan, D. Ramsden, S.P. Lees-Miller, J. A. Tainer, Structural insights into NHEJ: building up an integrated picture of the dynamic DSB repair super complex, one component and interaction at a time, DNA Repair (Amst.) 17 (2014) 110–120, [https://doi.org/10.1016/j.](https://doi.org/10.1016/j.dnarep.2014.02.009) $2014.02.00$
- [193] M. O'Driscoll, K.M. Cerosaletti, P.M. Girard, Y. Dai, M. Stumm, B. Kysela, B. Hirsch, A. Gennery, S.E. Palmer, J. Seidel, R.A. Gatti, R. Varon, M.A. Oettinger, H. Neitzel, P.A. Jeggo, P. Concannon, DNA ligase IV mutations identified in patients exhibiting developmental delay and immunodeficiency, Mol. Cell 8 (6) (2001) 1175–1185, [https://doi.org/10.1016/s1097-2765\(01\)00408-7](https://doi.org/10.1016/s1097-2765(01)00408-7).
- [194] E. Riballo, A.J. Doherty, Y. Dai, T. Stiff, M.A. Oettinger, P.A. Jeggo, B. Kysela, Cellular and biochemical impact of a mutation in DNA ligase IV conferring clinical radiosensitivity, J. Biol. Chem. 276 (33) (2001) 31124–31132, [https://](https://doi.org/10.1074/jbc.M103866200) doi.org/10.1074/jbc.M103866200.
- [195] K.D. Mumbrekar, H.V. Goutham, B.M. Vadhiraja, S.R. Bola Sadashiva, Polymorphisms in double strand break repair related genes influence radiosensitivity phenotype in lymphocytes from healthy individuals, DNA Repair (Amst.) 40 (2016) 27–34, [https://doi.org/10.1016/j.dnarep.2016.02.006.](https://doi.org/10.1016/j.dnarep.2016.02.006)
- [196] H.M. Tseng, D. Shum, B. Bhinder, S. Escobar, N.J. Veomett, A.E. Tomkinson, D. Y. Gin, H. Djaballah, D.A. Scheinberg, A high-throughput scintillation proximitybased assay for human DNA ligase IV, Assay. Drug Dev. Technol. 10 (3) (2012) 235–249, [https://doi.org/10.1089/adt.2011.0404.](https://doi.org/10.1089/adt.2011.0404)
- [197] H. Ma, A. Takahashi, Y. Yoshida, A. Adachi, T. Kanai, T. Ohno, T. Nakano, Combining carbon ion irradiation and non-homologous end-joining repair inhibitor NU7026 efficiently kills cancer cells, Radiat. Oncol. 10 (2015) 225, <https://doi.org/10.1186/s13014-015-0536-z>.
- [198] V.T. Chu, T. Weber, B. Wefers, W. Wurst, S. Sander, K. Rajewsky, R. Kuhn, Increasing the efficiency of homology-directed repair for CRISPR-Cas9-induced precise gene editing in mammalian cells, Nat. Biotechnol. 33 (5) (2015) 543–548, <https://doi.org/10.1038/nbt.3198>.
- [199] G. Li, X. Zhang, C. Zhong, J. Mo, R. Quan, J. Yang, D. Liu, Z. Li, H. Yang, Z. Wu, Small molecules enhance CRISPR/Cas9-mediated homology-directed genome editing in primary cells, Sci. Rep. 7 (1) (2017) 8943, [https://doi.org/10.1038/](https://doi.org/10.1038/s41598-017-09306-x) [s41598-017-09306-x](https://doi.org/10.1038/s41598-017-09306-x).
- [200] M. Srivastava, M. Nambiar, S. Sharma, S.S. Karki, G. Goldsmith, M. Hegde, S. Kumar, M. Pandey, R.K. Singh, P. Ray, R. Natarajan, M. Kelkar, A. De, B. Choudhary, S.C. Raghavan, An inhibitor of nonhomologous end-joining abrogates double-strand break repair and impedes cancer progression, Cell 151 (7) (2012) 1474–1487,<https://doi.org/10.1016/j.cell.2012.11.054>.
- [201] L. Nezi, A. Musacchio, Sister chromatid tension and the spindle assembly checkpoint, Curr. Opin. Cell Biol. 21 (6) (2009) 785–795, [https://doi.org/](https://doi.org/10.1016/j.ceb.2009.09.007) [10.1016/j.ceb.2009.09.007](https://doi.org/10.1016/j.ceb.2009.09.007).
- [202] B.T. Roberts, K.A. Farr, M.A. Hoyt, The Saccharomyces cerevisiae checkpoint gene BUB1 encodes a novel protein kinase, Mol. Cell Biol. 14 (12) (1994) 8282–8291, [https://doi.org/10.1128/mcb.14.12.8282-8291.1994.](https://doi.org/10.1128/mcb.14.12.8282-8291.1994)
- [203] A.A. Alizadeh, M.B. Eisen, R.E. Davis, C. Ma, I.S. Lossos, A. Rosenwald, J. C. Boldrick, H. Sabet, T. Tran, X. Yu, J.I. Powell, L. Yang, G.E. Marti, T. Moore, J. Hudson Jr., L. Lu, D.B. Lewis, R. Tibshirani, G. Sherlock, L.M. Staudt, Distinct types of diffuse large B-cell lymphoma identified by gene expression profiling, Nature 403 (6769) (2000) 503–511, [https://doi.org/10.1038/35000501.](https://doi.org/10.1038/35000501)
- [204] C. Sotiriou, S.Y. Neo, L.M. McShane, E.L. Korn, P.M. Long, A. Jazaeri, P. Martiat, S.B. Fox, A.L. Harris, E.T. Liu, Breast cancer classification and prognosis based on gene expression profiles from a population-based study, Proc. Natl. Acad. Sci. USA 100 (18) (2003) 10393–10398, <https://doi.org/10.1073/pnas.1732912100>.
- [205] M. Shichiri, K. Yoshinaga, H. Hisatomi, K. Sugihara, Y. Hirata, Genetic and epigenetic inactivation of mitotic checkpoint genes hBUB1 and hBUBR1 and their relationship to survival, Cancer Res 62 (1) (2002) 13–17. 〈[https://www.ncbi.nlm.](https://www.ncbi.nlm.nih.gov/pubmed/11782350) [nih.gov/pubmed/11782350](https://www.ncbi.nlm.nih.gov/pubmed/11782350)〉.
- [206] A.G. Morales, J.A. Pezuk, M.S. Brassesco, J.C. de Oliveira, R.G. de Paula Queiroz, H.R. Machado, C.G. Carlotti Jr., L. Neder, H.F. de Oliveira, C.A. Scrideli, L. G. Tone, BUB1 and BUBR1 inhibition decreases proliferation and colony formation, and enhances radiation sensitivity in pediatric glioblastoma cells, Childs Nerv. Syst. 29 (12) (2013) 2241–2248, [https://doi.org/10.1007/s00381-](https://doi.org/10.1007/s00381-013-2175-8) [013-2175-8.](https://doi.org/10.1007/s00381-013-2175-8)
- [207] S.J. Suijkerbuijk, G.J. Kops, Preventing aneuploidy: the contribution of mitotic checkpoint proteins, Biochim Biophys. Acta 1786 (1) (2008) 24–31, [https://doi.](https://doi.org/10.1016/j.bbcan.2008.04.001) [org/10.1016/j.bbcan.2008.04.001.](https://doi.org/10.1016/j.bbcan.2008.04.001)
- [208] C. Speers, S. Zhao, M. Liu, H. Bartelink, L.J. Pierce, F.Y. Feng, Development and Validation of a Novel Radiosensitivity Signature in Human Breast Cancer, Clin. Cancer Res 21 (16) (2015) 3667–3677, [https://doi.org/10.1158/1078-0432.CCR-](https://doi.org/10.1158/1078-0432.CCR-14-2898)
- [14-2898](https://doi.org/10.1158/1078-0432.CCR-14-2898). [209] M. Jessulat, R.H. Malty, D.H. Nguyen-Tran, V. Deineko, H. Aoki, J. Vlasblom, K. Omidi, K. Jin, Z. Minic, M. Hooshyar, D. Burnside, B. Samanfar, S. Phanse, T. Freywald, B. Prasad, Z. Zhang, F. Vizeacoumar, N.J. Krogan, A. Freywald, M. Babu, Spindle Checkpoint Factors Bub1 and Bub2 Promote DNA Double-Strand Break Repair by Nonhomologous End Joining, Mol. Cell Biol. 35 (14) (2015) 2448–2463, <https://doi.org/10.1128/MCB.00007-15>.
- [210] C. Yang, H. Wang, Y. Xu, K.L. Brinkman, H. Ishiyama, S.T. Wong, B. Xu, The kinetochore protein Bub1 participates in the DNA damage response, DNA Repair (Amst.) 11 (2) (2012) 185–191, [https://doi.org/10.1016/j.dnarep.2011.10.018.](https://doi.org/10.1016/j.dnarep.2011.10.018)
- [211] L. Santarpia, T. Iwamoto, A. Di Leo, N. Hayashi, G. Bottai, M. Stampfer, F. Andre, N.C. Turner, W.F. Symmans, G.N. Hortobagyi, L. Pusztai, G. Bianchini, DNA repair gene patterns as prognostic and predictive factors in molecular breast cancer subtypes, Oncologist 18 (10) (2013) 1063–1073, [https://doi.org/](https://doi.org/10.1634/theoncologist.2013-0163) [10.1634/theoncologist.2013-0163.](https://doi.org/10.1634/theoncologist.2013-0163)
- [212] P.G. Pilie, C. Tang, G.B. Mills, T.A. Yap, State-of-the-art strategies for targeting the DNA damage response in cancer, Nat. Rev. Clin. Oncol. 16 (2) (2019) 81–104, <https://doi.org/10.1038/s41571-018-0114-z>.