

# An update on PARP inhibitors for the treatment of cancer

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**Abstract:** The development of poly (adenosine diphosphate [ADP]) ribose polymerase (PARP) inhibitors (PARPi) has progressed greatly over the last few years and has shown encouraging results in the *BRCA1/2* mutation-related cancers. This article attempts to summarize the rationale and theory behind PARPi, the clinical trials already reported, as well as ongoing studies designed to determine the role of PARPi in patients with and without germline mutations of *BRCA* genes. Future plans for PARPi both as monotherapy and in combination with standard cytotoxics, other biological agents, and as radiosensitizers are also covered. The widening scope of PARPi adds another important targeted agent to the growing list of molecular inhibitors; future and ongoing trials will identify the most effective role for PARPi, including for patients other than *BRCA* germline mutation carriers.

**Keywords:** PARPi, *BRCA* genes, germline mutations, cytotoxics, radiosensitizers, *BRCA* germline mutation carriers

## Introduction

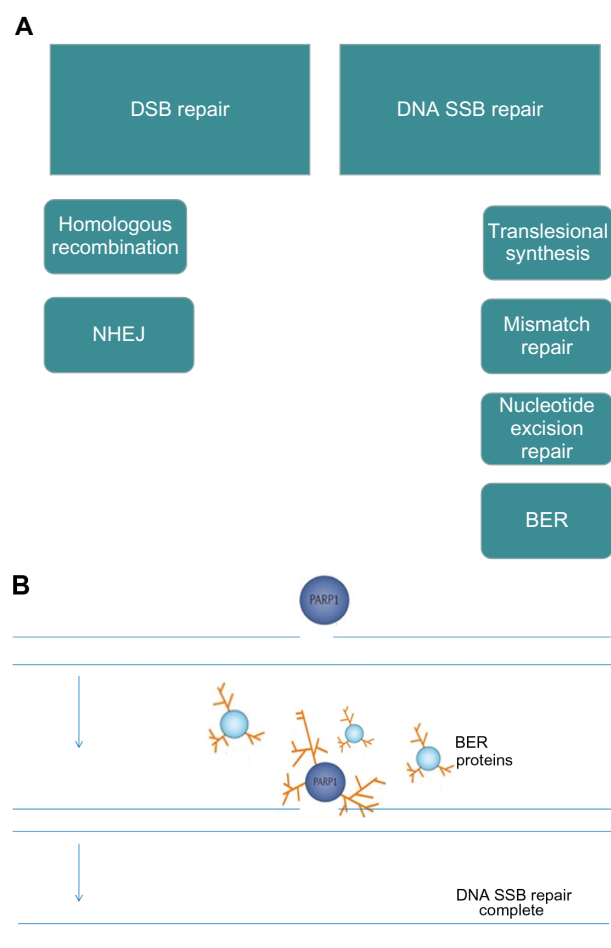
The increasing understanding of the functions of the poly (adenosine diphosphate [ADP]) ribose polymerase (PARP) enzymes in DNA repair among other things has led to the investigation of specific inhibitors of PARP in the cancer therapeutics setting.

The PARP proteins are a family of 17 enzymes involved in a wide range of cellular functions including DNA transcription, DNA damage response, genomic stability maintenance, cell cycle regulation, and cell death.<sup>1</sup> PARP enzymes are a subgroup of the ADP-ribosyltransferases (ARTs), which transfer ADP-ribose units from nicotinamide adenine dinucleotide (NAD<sup>+</sup>) molecules to form polymers of ADP ribose (PAR) units. These covalently bind to side chains of specific amino acid residues of acceptor proteins including PARP-1 itself.<sup>2</sup> PARP-1 is the most abundant and best characterized protein of this group. In oncology, its integral role in the repair of single-strand DNA breaks (SSBs) via the base excision repair (BER) pathway has been a focus of interest.

PARP-1 recognizes and binds to sites of SSBs (Figure 1B). Not all the PARP enzymes have been fully investigated, but some of them have been shown to catalyze mono-ADP-ribosylation, and thus do not fit the definition of a polymerase.<sup>3</sup> Therefore, there have been calls to revise the nomenclature of PARP enzymes to more accurately reflect these differences.<sup>2</sup> For the purposes of this review, we will use the term PARP, referring to the PARP-1 enzyme.

In cancer therapeutics, accumulation of SSBs with PARP inhibition leads to the development of double-strand DNA breaks (DSBs), which require competent

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**Figure 1** (A) DNA repair pathways; (B) PARP senses DNA SSBs and utilizes NAD<sup>+</sup> as a substrate to form PAR, which attach to a range of target proteins including PARP-1 itself and BER proteins. This posttranslational modification is termed PARylation.

**Notes:** (A) There are six DNA repair pathways, two for repair of DNA DSBs and four for repair of SSBs. PARP is known to be involved in BER and has been shown to be a negative regulator of NHEJ (low-fidelity, error-prone pathway). PARP inhibition therefore leads to disruption of BER and in HR-deficient cells, increased reliance on error-prone NHEJ.<sup>53</sup>

**Abbreviations:** BER, base excision repair; DSBs, double-strand DNA breaks; HR, homologous recombination; NAD<sup>+</sup>, nicotinamide adenine dinucleotide; NHEJ, nonhomologous end joining; PAR, polymers of ADP-ribose; PARP, poly (adenosine diphosphate [ADP]) ribose polymerase; SSBs, single-strand DNA breaks.

homologous recombination (HR) repair to allow cell survival.<sup>1</sup> PARP has also been shown to be involved in DSB repair pathways.

Much of the development of PARP inhibitors (PARPi) has been focused on targeting cancers associated with a mutation of the breast cancer-related genes, *BRCA1* and *BRCA2*, which are proteins that are integral to the HR repair pathway.<sup>4</sup> *BRCA* mutation carriers have a single functioning *BRCA* gene (wild type individuals have two copies of the *BRCA* gene), and when this is lost by mutation, cells are unable to undertake HR;<sup>5</sup> this 'second hit' makes a cell susceptible to tumor formation. The selective targeting of *BRCA*-deficient cancer cells by PARPi demonstrates the concept of synthetic

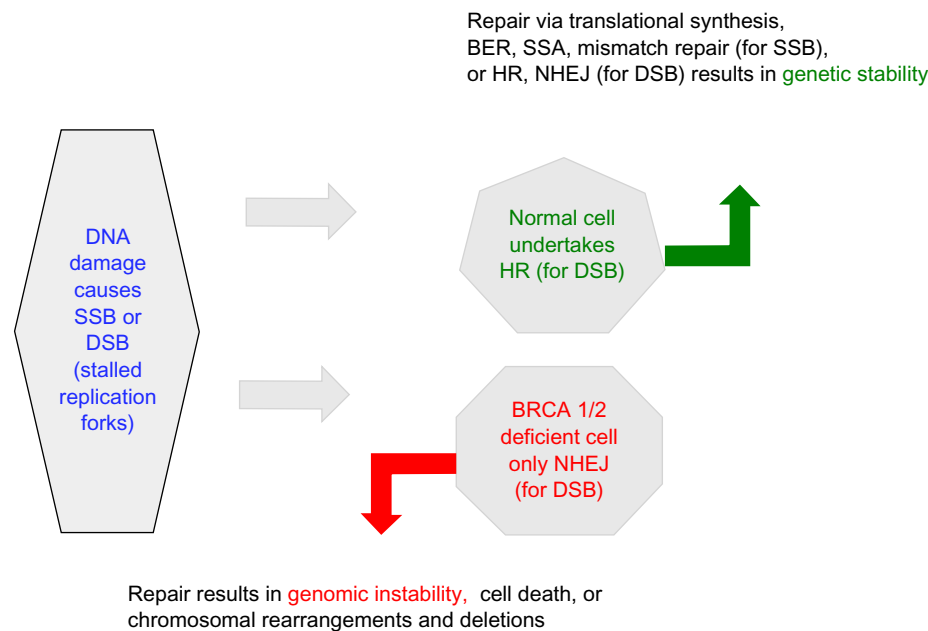
lethality, which was first described in 1922 by Bridges,<sup>6</sup> whereby a cell harboring one of two gene or protein defects is viable while a cell containing both defects is nonviable (Figure 2).<sup>7</sup>

Epidemiological studies have demonstrated a tight relationship between *BRCA1/2* mutations and the development of breast and ovarian cancers,<sup>8</sup> as well as other cancers.<sup>9</sup> PARPi selectively target *BRCA*-deficient cancer cells while sparing normal tissues, which retain a normal copy of the *BRCA* gene and can undergo normal HR repair of persistent SSBs not repaired by PARP.<sup>5</sup> Apart from blocking SSB repair, PARPi are also thought to act by 'trapping' PARP at sites of DNA damage, resulting in a cytotoxic PARP–DNA complex (Figure 3).<sup>10</sup> PARP localization to a point of DNA damage normally stimulates attraction of DNA repair proteins, which in turn prompt PARP dissociation from DNA, allowing repair to take place. The dissociation requires PARP to 'auto-PARylate' and is prevented by PARPi. In vitro studies by Murai et al demonstrated that PARP inhibition (by olaparib) and the PARP-1–DNA complexes generated are more cytotoxic than genetic depletion of PARP-1,<sup>10</sup> suggesting that there is more than one mechanism of action for PARPi. Interestingly, it has also been shown that the degree of PARP trapping varies between PARPi, niraparib being more potent than olaparib, which is more potent than veliparib. PARP trapping did not correlate with the catalytic inhibitory properties of each drug.<sup>10</sup>

## PARPi use in ovarian cancer

The Phase I trial of olaparib (AZD2281)<sup>11</sup> was the first clinical demonstration of PARPi activity in the *BRCA* mutation-associated cancers (Figure 4). In this study, 60 patients with refractory cancers were recruited to evaluate the pharmacokinetics and pharmacodynamics of olaparib. Twenty-two patients (37%) carried a *BRCA 1/2* mutation. A clinical benefit rate (CBR) of 63% was observed in this group of patients (12/19). This led to the enrollment of an expanded cohort of 50 patients with *BRCA1/2* mutation-associated gynecologic malignancies (ovarian, primary peritoneal, and fallopian tube cancers) where a CBR of 46% was observed. A Response Evaluation Criteria in Solid Tumors (RECIST) radiological response or CA125 response was seen in 40% of patients.<sup>12</sup> This early phase study confirmed the *BRCA1/2* mutation to be a predictive biomarker of response to PARPi.

In addition, it confirmed a correlation between PARPi response and platinum sensitivity, as platinum-sensitive patients displayed a 69% CBR compared with 46% and 23% in platinum-resistant and refractory patients, respectively.



**Figure 2** BRCA-deficient cells are highly reliant on BER as well as SSA and the error prone NHEJ pathway for DNA repair, both of which are influenced by PARPi.

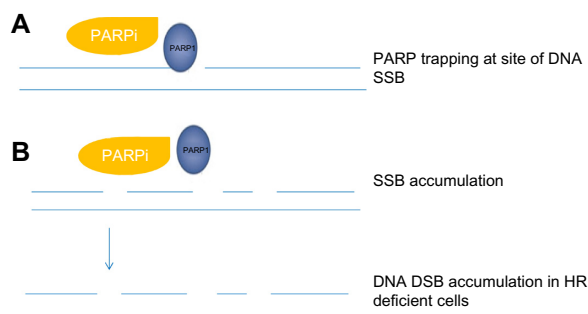
**Abbreviations:** BER, base excision repair; DSB, double-strand DNA breaks; HR, homologous recombination; NHEJ, nonhomologous end joining; SSA, single-strand annealing; SSBs, single-strand DNA breaks; PARPi - poly (adenosine diphosphate [ADP]) ribose polymerase inhibitors.

Platinum agents cause intra- and inter-strand DNA crosslinks, which lead to DNA DSBs. In cancer cells with HR deficiency (HRD), such as patients with mutant *BRCA*, cells become prone to further DNA damage and cell death. There is growing evidence that defects other than that of *BRCA1/2* can lead to HRD, which might explain the broader success of platinum agents in cancer,<sup>13</sup> the corollary being that cancers sensitive to platinum agents may also be sensitive to PARPi, widening their use beyond the *BRCA1/2* mutation carriers (Figure 4). McCabe et al demonstrated via in vitro studies

that deficiencies of several proteins in the HR DNA repair pathway, such as the DNA damage sensors ATM (ataxia telangiectasia mutated) and ATR (ataxia telangiectasia and RAD3-related protein), leads to HRD and subsequent PARPi sensitivity.<sup>13</sup> This concept, known as 'BRCAness' has been used to describe the phenotype arising in sporadic cancers that have intact *BRCA1/2* genes but share features with the *BRCA1/2* mutation-related tumors, such as profound platinum sensitivity.<sup>14</sup>

## Tumors with BRCA-like features

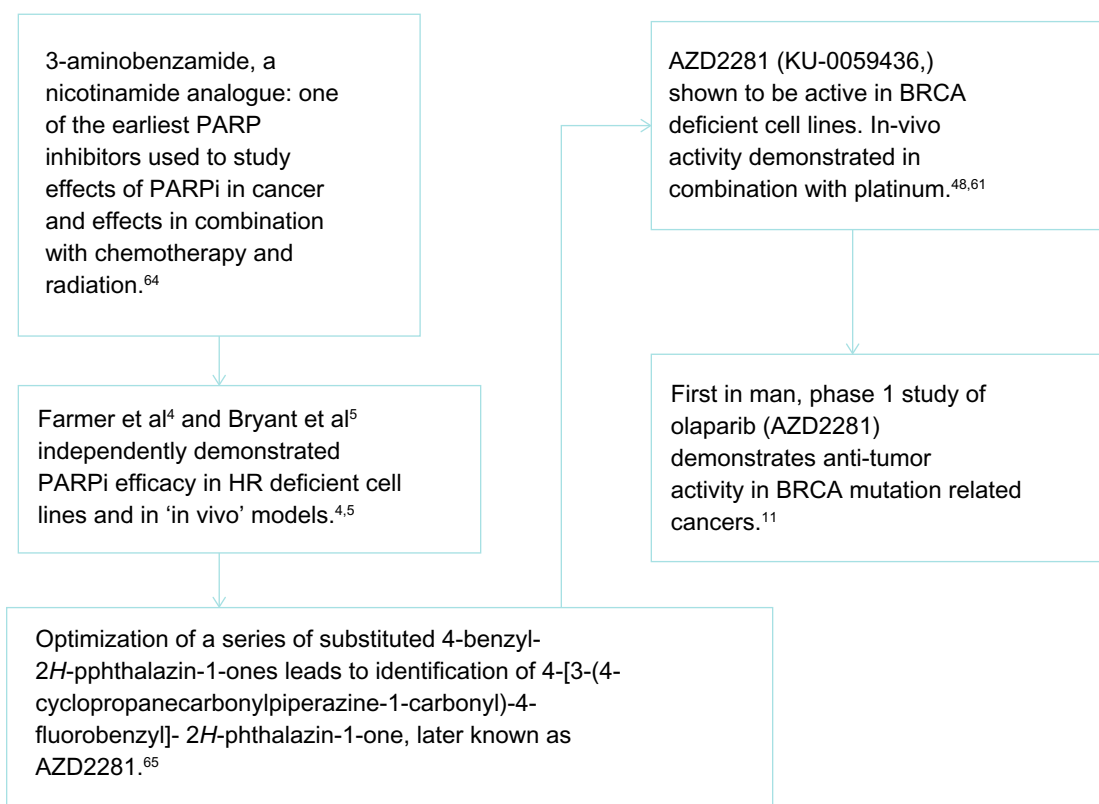
'BRCAness' can be illustrated by the responses in high-grade serous ovarian cancers (HGSOC), where nearly 20% of cases occur in women with germline *BRCA1/2* mutations, but a further 35% are thought to have acquired defects of HR<sup>15</sup> and display the same sensitivity to platinum chemotherapy, as well as to PARPi. This was highlighted in a Phase II trial of olaparib, which recruited both *BRCA1/2* mutated as well as wild type patients with breast cancer and HGSOC. In the cohort of patients who had platinum-sensitive HGSOC without *BRCA1/2* mutation, the objective response rate (ORR) was 50% (10/20).<sup>16</sup> In addition, among 38 *BRCA* wild type patients with ovarian cancer, a 26% response rate in CA125 was observed and a combined RECIST or CA125 response rate of 30% was recorded. All these responders had HGSOC. Interestingly, responses were also seen in the platinum-resistant cohort without *BRCA 1/2* mutation



**Figure 3** PARP inhibitor – mechanism of action.

**Notes:** (A) PARP attaches to site(s) of DNA damage and is unable to dissociate due to inhibition of PARP autoPARylation causing PARP trapping and cytotoxic PARP–DNA complexes. The degree of 'PARP trapping' appears to vary between the different inhibitors currently under investigation and may explain the differences in toxicity profile observed, with olaparib and niraparib having greater potency compared to veliparib.<sup>34</sup> (B) SSBs accumulate, leading to DSBs, which are unrepaired in cells deficient in HR proteins, ultimately leading to cell death.

**Abbreviations:** DSBs, double-strand DNA breaks; HR, homologous recombination; PARP, poly (adenosine diphosphate [ADP]) ribose polymerase; SSBs, single-strand DNA breaks.



**Figure 4** Schema of development of first PARPi, olaparib (AZD2281).

**Abbreviations:** HR, homologous recombination; PARP, poly (adenosine diphosphate [ADP]) ribose polymerase; PARPi, PARP inhibitor.

(17% response rate), suggesting incomplete crossover of platinum sensitivity and PARPi response.<sup>16</sup> Similar findings were found by Ledermann et al in a randomized double-blind placebo-controlled Phase II trial of maintenance olaparib in HGSOC, in which patients had responded to two or more lines of platinum chemotherapy.<sup>17</sup>

Findings such as this have led to Phase III trials of maintenance PARPi treatment after chemotherapy for all ovarian cancer patients; many of the PARPi combination trials are recruiting patients with HGSOC regardless of *BRCA* status. Currently recruiting is a randomized double-blind placebo-controlled trial of niraparib (MK-4827) as maintenance therapy in platinum-sensitive ovarian cancer patients. This study is recruiting patients who have either a *BRCA1/2* mutation or tumors with high-grade serous histology and have had a response to their most recent platinum-containing chemotherapy (NCT01847274). In contrast, the SOLO1 and SOLO2 trials are ongoing Phase III randomized, placebo-controlled trials investigating the use of olaparib maintenance in the first-line (SOLO 1 NCT01844986) and relapsed setting (SOLO 2 NCT01874353) and are only recruiting *BRCA* mutation carriers with ovarian cancer.

The ARIEL2 and NOVA trials are Phase II trials investigating rucaparib and niraparib, respectively, in platinum-sensitive relapsed ovarian cancer (NCT01891344). One of the aims of ARIEL2 is to identify a molecular signature associated with HRD in HGSOC and fallopian tube cancers, which can then be used to identify patients most likely to respond to rucaparib. A Phase III trial (ARIEL3) is planned once this signature is identified to prospectively assess its use in this patient population.

## PARPi use in breast cancer

In breast cancer, a Phase II trial of single-agent olaparib in patients with *BRCA1/2* mutation and advanced disease investigated two dose levels (cohort 1: 400 mg twice daily [bd] and cohort 2: 100 mg bd) and showed an ORR of 42% (11/26) and 25% (6/24), respectively. Progression-free survival (PFS) was 5.7 months in cohort 1 and 3.8 months in cohort 2.<sup>18</sup>

Breast cancer patients with *BRCA1* mutations also display marked platinum sensitivity; treatment with neoadjuvant cisplatin in such individuals results in high pathological complete response rates (72%–90%).<sup>19,20</sup> The majority of breast cancers in *BRCA* mutation carriers are found to be negative

for the estrogen receptor (ER), progesterone receptor, and human epidermal growth factor receptor (HER2), and are termed triple negative breast cancer (TNBC). Non-*BRCA*-mutated TNBC appears to harbor a 'BRCAness' phenotype due to HRD;<sup>21</sup> responses to PARPi in this population are being investigated in a number of Phase I and II trials. However, unlike HGSOc patients with *BRCA* mutations, TNBC patients are thought to have a poorer prognosis compared to hormone receptor-positive breast cancers. Sporadic TNBC has been shown to have a two-fold lower expression of *BRCA1* compared to ER-positive cancers.<sup>22</sup> Lower *BRCA1* expression is possibly related to a number of factors, such as upregulation of *BRCA1* transcriptional repressors such as ID4 and HMG1, and of microRNAs (miR) that negatively regulate *BRCA1* expression such as miR-182.<sup>23</sup> Many TNBCs share features with the basal-like phenotype classically associated with *BRCA1*-related cancers, yet only 20% of cases have a germline *BRCA1/2* mutation. Although the majority of TNBCs are *BRCA*-mutation-negative, the shared phenotypic characteristics and molecular aberrations such as those mentioned above lead to HRD, which could sensitize cells to PARP inhibition. The basis of current investigations with respect to TNBC and its response to PARPi revolve around HRD. Potential mechanisms include *BRCA1* promoter methylation leading to gene silencing, aberrations of MRE11 protein (involved in DSB detection),<sup>24</sup> and ATM, all of which play important roles in HR.

A three-arm, double-blinded, placebo-controlled Phase III trial of veliparib (ABT-888) in TNBC with carboplatin and paclitaxel after neoadjuvant chemotherapy (doxorubicin and cyclophosphamide) in early-stage TNBC is ongoing (NCT02032277).

## PARPi in other cancers

### Endometrial cancer

The most common genetic aberration seen in up to 80% of endometrioid endometrial cancer (EEC) is loss of the *PTEN* tumor suppressor gene.<sup>25</sup> However, apart from *PTEN*'s function in the regulation of the PI3 kinase/AKT/mTOR signaling pathway, it is apparent that *PTEN* competence is also important in maintaining genetic stability.<sup>26</sup> Dedes et al showed *PTEN*-deficient EEC cells lack HR DNA repair,<sup>27</sup> which sensitizes the EEC cells to PARPi in a similar fashion to that seen with *BRCA*-mutated cancers. These *in vitro* results have also been reflected by *in vivo* models of *PTEN*-deficient tumors treated with PARPi where tumor growth was significantly suppressed by PARPi treatment.<sup>28</sup> Although this relationship between *PTEN* loss and sensitivity to PARPi has not been

investigated in the context of a clinical trial, a case report of a patient with metastatic endometrial cancer who was treated with olaparib in a Phase I trial showed significant clinical response.<sup>29</sup> Subsequent tumor biopsies confirmed no *BRCA* mutation but did demonstrate *PTEN* loss.<sup>29</sup> Together with preclinical studies, this suggests a potential role for PARPi treatment for *PTEN*-deficient tumors, including endometrial cancer among others. *PTEN* loss may perhaps also serve as a predictive biomarker for PARPi sensitivity.

### Prostate cancer

A minority of prostate cancers are attributed to *BRCA1/2* mutation, and in these cases, PARPi sensitivity may be observed with striking clinical responses and lengthened overall survival (OS).<sup>11</sup> Akin to endometrial cancer, 20% of prostate cancers show loss of *PTEN* expression and similar preclinical studies of *PTEN*-mutant prostate cancer cell lines have suggested that *PTEN* loss predicts sensitivity to PARPi with olaparib and veliparib.<sup>13</sup>

The Phase I trial of niraparib reported in 2013 set out to assess the relationship between *PTEN* loss in prostate cancer and response.<sup>30</sup> Overall, niraparib showed good tolerability, and as seen in ovarian cancer patients, clinical responses were achieved in both *BRCA* mutation carriers (~50% CBR) as well as sporadic (*BRCA* wild type) cancers. Surprisingly, molecular defects such as *PTEN* loss or *ETS* gene rearrangements, often found in castration-resistant prostate cancer (CRPC), did not correlate with measures of antitumor activity in this Phase I study.<sup>30</sup> It is possible that CRPC tumor cells harbor other defects of HR such as ATM, CHEK2, and FANCD1.<sup>31,32</sup> The use of PARPi in prostate cancer is thus being further studied in combination trials such as that due to start this year comparing olaparib alone to olaparib and abiraterone in metastatic CRPC regardless of *BRCA* status (NCT01972217).

### Pancreatic cancer

Although hereditary pancreatic cancer is rare, 10% of cases are due to inherited genetic factors,<sup>33,34</sup> the most common group being related to *BRCA* mutations. These account for up to 17% of inherited pancreatic cancer.<sup>35</sup> *BRCA2* mutation increases the risk of pancreatic cancer development by 3.5-fold.<sup>36</sup> As *BRCA*-associated pancreatic cancer is rare, the data are sparse regarding the use of PARPi, though the responses of patients treated within mixed cohorts in early phase trials of PARPi combinations have been encouraging, with *BRCA*-mutant patients achieving meaningful partial responses of 1–2 years.<sup>37,38</sup>



In a Phase II study of olaparib monotherapy for *BRCA* mutation-associated solid tumors, platinum-based chemotherapy was used first-line in six metastatic pancreatic cancer (MPC) patients; five (83%) achieved a partial response or complete response by RECIST criteria<sup>38</sup> suggesting that *BRCA*-associated MPC may share the platinum sensitivity observed in other *BRCA*-associated malignancies.

Another trial investigating the use of PARPi in *BRCA*-associated MPC, with rucaparib monotherapy (NCT02042378), continues to recruit, and a third opening imminently involves maintenance olaparib in patients whose disease has not progressed on platinum-based first-line chemotherapy (NCT02184195).

## PARPi in combination therapy

### PARPi and ionizing radiation

Like platinum agents, PARPi have been shown to act as radiosensitizers in preclinical studies using both in vitro and in vivo models.<sup>39</sup> By combining ionizing radiation with PARPi, the SSB induced by irradiation go unrepaired by PARP-associated BER, leading to cell death and delay in tumor growth. In vitro studies in lung<sup>40</sup> and prostate cancers have confirmed this effect, although as described above, in prostate cancer the mechanisms for PARPi sensitization may be as a consequence of PTEN deficiency or upregulation of the fusion gene *TPR52-ERG*.<sup>40–42</sup>

Two Phase I trials are underway comparing concomitant PARPi and radiotherapy with radiotherapy alone in esophageal and locally advanced pancreatic cancers (NCT01460888, NCT01908478).

### PARPi and chemotherapy

Extensive preclinical studies have shown that PARPi enhance the cytotoxic effects of chemotherapy. The knockout of PARP-1 drastically impairs DNA repair following damage caused by ionizing radiation or cytotoxic treatment.<sup>43</sup> Although PARP is expressed in all cells, its expression and activity is increased in actively proliferating cells. Several hematological malignancies as well as solid tumors have been found to have higher levels of PARP expression compared with normal cells.<sup>44,45</sup> Therefore, PARPi could selectively target tumor cells. Preclinical studies have shown that PARP inhibition potentiates the cytotoxic effects of agents that cause SSBs in DNA, such as alkylating agents (eg, temozolomide<sup>46</sup> or cyclophosphamide) and topoisomerase inhibitors (eg, irinotecan<sup>47</sup>). However, platinum agents cause DNA crosslinks which are usually repaired either by nucleotide/

BER or HR. Therefore, theoretically and in vitro, combined therapy cytotoxics such as these, and PARPi even in normal cells should result in synergism.<sup>48</sup> Early combination trials of PARPi and cytotoxics in vivo have been reported<sup>49–51</sup> with encouraging results although, as might be expected, responses were more commonly seen in *BRCA* mutation carriers, suggesting that they have the greatest effect in patients with inherent DNA repair defects.<sup>50</sup>

Table 1 details trials of PARPi currently underway in combination with chemotherapy. Phase II studies of veliparib with topotecan (in refractory ovarian or primary peritoneal cancer [NCT01012817]) and veliparib with metronomic cyclophosphamide (in refractory gynecological cancers, TNBC, and low-grade non-Hodgkin lymphoma [NCT01306032]) are currently ongoing. A detailed genetic study of DNA repair defects as part of the latter study may identify HRD other than *BRCA* mutations to predict PARPi sensitivity. Myelosuppression is the main toxicity encountered in these combination studies, and as expected, is more significant when PARPi are combined with topotecan.<sup>49,51,52</sup> Similarly, two completed trials, olaparib with paclitaxel in TNBC<sup>45</sup> and rucaparib with temozolomide in melanoma,<sup>53</sup> have shown trends toward higher response rates and longer PFS than historical controls but both required dose reductions of the cytotoxic agents for myelosuppression.

Current PARPi and chemotherapy combination trials are summarized in Table 1.

## PARPi and targeted agents

### Combination with antiangiogenics

PARPi in combination with targeted antiangiogenic agents have been studied in the breast and ovarian cancer population. Like PARPi, antiangiogenics such as bevacizumab (anti-vascular endothelial growth factor [VEGF] antibody) and cediranib (a small-molecule inhibitor of VEGF receptor [VEGFR]-1/2/3) have been shown to have single-agent activity in recurrent ovarian cancer, with a 18% response rate to bevacizumab<sup>54</sup> and 19% response rate to cediranib.<sup>55</sup> Preclinical studies have suggested that PARPi also have an antiangiogenic effect,<sup>56</sup> which prompts speculation that synergism may be achieved by combination therapy.

One of the first trials to assess this combination was the Phase I trial of olaparib and cediranib in recurrent ovarian cancer or TNBC.<sup>57</sup> In 18 RECIST evaluable ovarian cancer patients, the ORR was 44% and CBR was 61%. These responses were not limited to the *BRCA* mutation carriers. No responses were seen in seven evaluable breast

**Table I** Currently recruiting trials of PARPi and chemotherapy combinations

PARPi/chemo combination	Phase of study	Population
BMN673 + TMZ NCT02116777	Phase I dose escalation followed by Phase II	Phase II: patients aged > 12 months and ≤21 months, relapsed/refractory solid tumors Phase I: > 12 months and ≤30 months, Ewing's sarcoma, PNET, ALL
Olaparib + C AUC4 NCT01237067	Phase I	Recurrent/refractory gynecologic tumors and recurrent/refractory metastatic BC BRCA mutation BC that is locally advanced or metastatic, even if not treated previously
AZD2281 + C NCT01445418	Phase I	BRCA +ve BC and OC Sporadic OC Sporadic TNBC
Niraparib + TMZ NCT02044120	Phase I	Previously treated, incurable Ewing's sarcoma
Veliparib + pegylated liposomal doxorubicin NCT01145430	Phase I	Recurrent OC, fallopian tube cancer, PPC, metastatic TNBC (BRCA +ve and -ve)
Veliparib + floxuridine NCT01749397	Phase I	Metastatic epithelial ovarian, PPC, or fallopian tube cancer
Olaparib + TMZ NCT01390571	Phase I	Relapsed glioblastoma
TMZ +/- veliparib NCT01638546	Phase II, randomized placebo-controlled	Relapsed or refractory small cell lung cancer
E7449 single agent or with TMZ or with C and P NCT01618136	Phase I dose escalation followed by Phase II	Single agent: advanced solid tumors Combination arms: B cell malignancies (TMZ) or advanced solid tumors (C + P)
Veliparib + topotecan NCT01012817	Phase I dose escalation followed by Phase II	Phase I: advanced solid tumors Phase II: recurrent ovarian or PPC
Veliparib + C, P, and bevacizumab NCT00989651	Phase I	Newly diagnosed stage II, III, or IV OC, fallopian tube cancer, or PPC

**Abbreviations:** ALL, acute lymphoblastic leukemia; AUC, area under the curve; BC, breast cancer; C, carboplatin; OC, ovarian cancer; P, paclitaxel; PNET, peripheral primitive neuroectodermal tumor; PPC, primary peritoneal cancer; TMZ, temozolomide; TNBC, triple negative breast cancer.

cancer patients, though two patients had stable disease (both *BRCA* mutation–positive) for more than 24 weeks. Grade 3 or 4 toxicities were observed in 75% of patients, the most common being hypertension (25%), fatigue (18%), and neutropenia (11%).<sup>57</sup>

Results from a Phase II trial comparing olaparib alone to olaparib with cediranib in ovarian cancer (NCT01116648) were presented earlier this year (American Society of Clinical Oncology, 2014) and showed a remarkable improvement in ORR (79.6% vs 47.8% and PFS 17.7 months vs 9.0 months, HR: 0.42) for the combination arm versus single-agent olaparib.<sup>58</sup> There was a 39%, 23%, and 27% incidence of grade 3 hypertension, diarrhea, and fatigue, respectively, in the combination arm; 77% of patients in the combination arm were dose reduced compared to 24% in the olaparib-alone arm. However, the doses of both agents were relatively high at 30 mg daily cediranib and 200 mg bd olaparib capsules in the combination arm and 400 mg bd olaparib capsules in the single agent arm. OS data are not yet mature, but analysis of the population in relation to *BRCA* mutation status revealed

that the improvement in PFS with combination olaparib/cediranib was much more marked in the *BRCA* wild type or unknown patients than those with *BRCA* mutations.<sup>58</sup>

## Combination with PI3K inhibition

Study into the molecular defects of TNBC has identified aberrations involving the PI3 kinase/AKT/mTOR pathway such as *PTEN* loss or activating mutations of *PIK3CA* gene encoding PI3K.<sup>59</sup> Inhibition of PI3K in TNBC cells causes a downregulation of *BRCA1/2* and increased PARP activity, suggesting an increased dependency on *BRCA*-mediated DNA repair and a potential sensitizing effect to treatment with PARPi when PI3K is inhibited. Additionally, in TNBC xenograft models, dual inhibition of PI3K and PARP significantly suppressed tumor growth by downregulating *BRCA1/2*.<sup>59</sup>

On the basis of this work, a Phase I trial of a PI3K inhibitor (buparlisib) combined with olaparib is underway in TNBC and HGSOc patients (NCT01623349). Translational endpoints in this study will be vital to determine predictive biomarkers of response.

## Future directions

Clinical trials of PARPi in cancer patients have already shown that these agents are active in patients irrespective of *BRCA* mutation status. In ovarian cancer, the growing understanding of multiple DNA repair defects other than *BRCA* mutations makes it likely that these targeted inhibitors can be applied to many *BRCA* wild type as well as mutation carrier patients. Predictive tools such as molecular profiles to identify responsive patients remain an important translational focus. Preclinical studies have already identified potential markers of PARPi sensitivity other than HR defects but these have yet to be proven clinically useful. Mechanisms of resistance to PARPi are also being unraveled. To date, three have been described: secondary mutations in *BRCA1/2* that partially restore the gene reading frame and protein function,<sup>60</sup> overexpression of the multidrug efflux transporter protein Pgp,<sup>61</sup> and loss of 53BP1 with partial restoration of HR.<sup>62</sup> These relationships are not clear, as 36%–45% of *BRCA* mutation carriers with ovarian cancer who progressed on PARPi therapy went on to respond to further chemotherapy, independent of previous PARPi platinum sensitivity or response to PARPi.<sup>63</sup> Understanding these interactions and identifying predictive markers of sensitivity and resistance will inform the treatment strategy for PARPi agents and cytotoxics in the future.

In this era of personalized medicine, combinations of chemotherapy and molecularly targeted agents, such as PARPi, have already become standard treatment in some settings, eg, FOLFOX (fluorouracil and oxaliplatin) and cetuximab in *KRAS* wild type metastatic colorectal cancer and docetaxel with trastuzumab in HER2-positive breast cancer. Current combination treatment trials will hopefully identify the most effective role for PARPi combinations.

## Disclosure

Dr Benaffif reports no conflicts of interest in this work. Dr Hall has received honorarium for advisory boards from Roche, Astra Zeneca, Glaxo Smith Kline, Boehringer Ingelheim, and Merck.

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