New Targeted Agents in Gynecologic Cancers: Synthetic Lethality, Homologous Recombination Deficiency, and PARP Inhibitors

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Opinion statement
Inhibitors of poly (ADP-ribose) polymerase (PARP) have emerged as a new class of anti-cancer drugs, specifically for malignancies bearing aberrations of the homologous recombination pathway, like those with mutations in the BRCA 1 and BRCA 2 genes. Olaparib, a potent PARP1 and PARP2 inhibitor, has been shown to significantly increase progression-free survival (PFS) in women with recurrent ovarian cancer related to a germline BRCA mutation and is currently approved fourth-line treatment in these patients. PARP inhibitors (PARPi) target the genetic phenomenon known as synthetic lethality to exploit faulty DNA repair mechanisms. While ovarian cancer is enriched with a population of tumors with known homologous recombination defects, investigations are underway to help identify pathways in other gynecologic cancers that may demonstrate susceptibility to PARPi through synthetically lethal mechanisms. The ARIEL2 trial prospectively determined a predictive assay to identify patients with HRD. The future of cancer therapeutics will likely incorporate these HRD assays to determine the best treatment plan for patients. While the
role of PARPi is less clear in non-ovarian gynecologic cancers, the discovery of a predictive assay for HRD may open the door for clinical trials in these other gynecologic cancers enriched with patients with HRD. Identification of patients with tumors deficient in homologous repair or have HRD-like behavior moves cancer treatment towards individualized therapies in order to maximize treatment effect and quality of life for women living with gynecologic cancers.

Introduction

Each year, over 1 million women worldwide are newly diagnosed with gynecologic malignancy, and almost 500,000 women will die from gynecologic cancer [1]. The last decade of research in the treatment of gynecologic cancers has seen the development of multimodal options, including anti-angiogenic biologics, like bevacizumab, and targeted therapies, like olaparib, a PARPi recently approved for the fourth-line treatment of BRCA-deficient ovarian cancers. Olaparib is the first PARPi approved by the Food and Drug Administration (FDA) for clinical use in the treatment of cancer in the USA, based only on phase II efficacy and safety data. PARP inhibitors have the benefit of being an oral medication which minimizes the impact on quality of life for patients with recurrent cancer. The significant clinical impact of PARP inhibition, a manifestation of sound translational rationale behind therapeutic development, can be attributed to the exploitation of synthetic lethality as a mechanism that selectively targets a specific population of cancer cells, those deficient in tumor suppressor genes BRCA 1 and BRCA 2. The purpose of this article is to provide a background on synthetic lethality, an overview of clinical trials investigating the use of PARPi in gynecologic cancers, highlighting the approval of olaparib in the treatment of recurrent BRCA-deficient ovarian cancer, and identification of patients with a homologous recombination-deficient (HRD) profile to better tailor treatment of women with gynecologic cancers in the future (Table 1).

Synthetic Lethality

During the early twentieth century, the American geneticist Calvin Bridges (1889–1938) (Fig. 1) noted that when crossing the fruit fly, Drosophila melanogaster, certain non-allelic genes were lethal only in combination even when homozygous parents were viable [2]. Twenty years later, the term “synthetic lethality” was coined by Bridges’ colleague, Theodosius Dobzhansky, who reported the same observations in Drosophila pseudoobscura [3]. The ancient Greek meaning of synthetic is the combination of two entities to form something new [4••]. Thus, synthetic lethality occurs when a genetic defect or defective protein is compatible with cell viability but is lethal when combined (i.e., synthesized) with another genetic/protein defect. By way of contrast, genetic combinations resulting in non-lethal growth impairment are synthetic sick.

In their description of induced essentiality, Tischler et al. provided a hypothesis to account for synthetic lethal and oncogene addiction effects in nature [5]. Lord et al. note that as tumor cells acquire more mutations, significant deleterious effects may be minimized through molecular networks within a cell to facilitate compensatory alterations that permit cell survival via escape or functional buffering [3]. As an example, oncogenes leading to increased cell proliferation induce a state of replicative stress which results in slowing or stalling at the replication forks, ultimately leading to DNA damage deleterious to cancer cells. To minimize this effect, oncogene activation is often associated with compensatory molecular changes mediated by the ataxia telangiectasia mutated (ATM) and ataxia telangiectasia and Rad3-related (ATR) protein kinases.

Exploitation of synthetic lethality to eradicate cancer was initially suggested by Hartwell et al. who outlined strategies through which drugs could be profiled for their ability to selectively kill cells in a molecular context that matches those found in malignant neoplasms [6]. Kaelin advanced this idea in noting that because targeting a gene that is synthetic lethal to a cancer-relevant mutation should kill only cancer cells and spare normal cells, synthetic lethality provides a conceptual framework for the development of cancer-specific agents [7]. Theoretically, the development of synthetic lethal resistance occurs not through modulation of the drug target but rather through modulation of the synthetic lethal partner. The most robust demonstration of the principle of harnessing synthetic lethality comes from the treatment of cancers resulting from loss of BRCA gene function.

The development of PARPi as therapeutic options for cancer treatment capitalizes on the role of PARP in DNA
Table 1. Important phase II trials of parp inhibitors and ovarian cancer

| Trial (year) | Study Design | N  | Eligibility                                                                 | Drug dose                                      | ORR          | PFS           | Grade 3/4 AEs                  |
|-------------|--------------|----|-----------------------------------------------------------------------------|-----------------------------------------------|--------------|---------------|------------------------------|
| Lederman et al. (2013) Study 19 [36] | Randomized phase II | 265 | Recurrent HGSOC · Platinum sensitive · >2 prior regimens · ± gBRCA mutation | Olaparib 400 mg BID · Placebo                  | 12 % versus 4 % | 8.8 versus 4.8 months | Nausea, fatigue, emesis, anemia |
| Coleman et al. (2014) [56] | Phase II | 52  | Recurrent HGSOC · ± platinum sensitive · ≤3 prior regimens · ± gBRCA mutation | Veliparib 400 mg BID × 28d for 6 cycles         | 26 %         | 8.1 months OS: 19 months | Nausea, emesis, neutropenia, TCP |
| Liu et al. (2014) [54] | Randomized phase II | 90  | Recurrent disease · HGSOC or endometrioid (any if BRCA mutation) · Platinum sensitive · <3 prior regimens · ± gBRCA mutation | Olaparib 400 mg BID · Olaparib 200 mg BID + Cediranib 30 mg daily | 48 % versus 80 % | 9.0 versus 17.7 months | Fatigue, diarrhea, hypertension |
| Oza et al. (2015) Study 41 [48] | Randomized phase II | 162 | Recurrent HGSOC · Platinum sensitive · <3 prior regimens · ± gBRCA mutation | Carboplatin/paclitaxel + Olaparib 200 mg BID (d1-10) → Olaparib 400 mg bid | 64 % versus 58 % | 12.2 versus 9.6 months | Neutropenia, anemia |
| McNeish et al. (2015) ARIEL2 [20] | Phase II | 204 | Recurrent HGSOC · Platinum sensitive · ≥1 prior Platinum regimens · ± gBRCA mutation | Rucaparib 600 mg BID +BRCA: 82 % BRCA-like: 45 % Biomarker: 21 % | +BRCA: 82 % | 9.4 months 7.1 months 3.7 months | Anemia, transaminitis, fatigue |
| Kaufman et al. (2015) Study 42 [37] | Phase II | 298 | Recurrent disease, ovarian cancer (N = 193) · ± gBRCA mutation | Olaparib 400 mg BID | Ovarian cancer: 31 % | 7 months OS: 16.6 months | Anemia, abdominal pain, fatigue |

AE adverse events. BID twice daily. gBRCA germline breast cancer gene. HGSOC high-grade serous ovarian cancer. ORR response rate. OS overall survival. PFS progression-free survival. TCP thrombocytopenia
repair and the cancers already deficient in homologous recombination, like BRCA-related breast and ovarian cancers (Fig. 2, [8]). DNA undergoes constant damaging sequence alterations due to toxic byproducts of the cell cycle, environmental insult, and errors in replication. Several mechanisms have evolved to repair these errors, including (1) nucleotide excision repair, (2) base excision repair (BER), (3) homologous recombination (HR), and (4) non-homologous end-joining (NHEJ).

PARP enzymes are found in the cellular nucleus where they are activated by DNA damage to identify DNA single-strand breaks (SSB) [9]. PARP is thought to have four domains: an N-terminal DNA-binding domain comprised of two zinc finger motifs, a C-terminal catalytic domain, a central auto-modification domain, and a caspase-cleaved domain [10]. After binding to a site of SSB, PARP undergoes a conformational change allowing for the C-terminal catalytic domain to transfer ADP-ribose moieties from nicotinamide-adenine-dinucleotide (NAD⁺) to form PAR chains that recruit other DNA repair proteins (e.g., DNA ligase III, DNA polymerase beta) to form the BER multiplex, ultimately repairing SSB (Fig. 2). The formation of PAR chains also appears to play a role in multiple other cellular tasks, like gene expression and signal response [11–13], as well as double-stranded DNA repair [14–16]. PARPi interrupts the DNA repair process by impairing two specific mechanisms of PARP: (1) binding sites of SSB with its own zinc finger domains thereby directly blocking DNA access by PARP and (2) preventing the transfer of ADP-ribose to form PAR chains thereby blocking the formation of the BER multiplex. The summation of these

Fig. 1. American geneticist, Calvin Blackman Bridges in 1927. Photograph courtesy of Cold Spring Harbor Laboratory Archives, used with permission.

Fig. 2. PARP inhibition mechanism of action—blockade of the base excision pathway. Poly(ADP-ribose) polymerase (PARP) recognizes and binds to sites of DNA damage through its zinc-finger domains and recruits proteins involved in DNA repair through polyADP-ribose catalyzation. PARP inhibitors function by trapping PARP to sites of DNA damage and blocking the enzymatic transformation required for polyADP-ribosylation. Adapted from Tewari KS, Monk BJ, “Translational Science,” In: The 21st Century Handbook of Clinical Ovarian Cancer, page 80. © Springer International Publishing Switzerland 2015, with permission of Springer.
actions leads to the progression of SSB to DSBs at replication forks, and in cells without intact homologous repair, chromosomal integrity is destroyed, the cell cycle arrests, and apoptosis results.

The loss of high-fidelity homologous recombination is partly due to an inability to localize DNA polymerase RAD51 [3]. Preclinical studies showed that treatment of BRCA-deficient cells with PARP inhibition induced the presence of nuclear RAD51 foci, an indication of double-strand DNA repair [17]. Indeed, subsequent in vitro studies demonstrated that cells with BRCA mutations are 1000 times more sensitive to PARPi compared to wild-type cells [18, 19]. These observations provided the translational impetus to begin phase I and II clinical trials with PARPi in breast, ovarian, and prostate cancers. In the most recent gynecologic cancer clinical trials of PARPi, specifically in the ARIEL2 trial, tumors with deficiencies in RAD51 demonstrated a BRCA-like HRD phenotype with high genomic loss of heterozygosity (LOH) and increased response to rucaparib [20••].

While the focus of PARPi has been in the treatment of BRCA-related ovarian cancer, their therapeutic use in other gynecologic cancers is under investigation. Up to 80% of sporadic endometrial cancers have been associated with activation of the phosphatidylinositide 3-kinase (PI3-kinase) pathway via mutations in phosphatase and tensin homologue (PTEN) [21, 22], and early studies in mouse embryonic fibroblasts showed that PTEN inactivation induced genomic instability due to defective RAD51-mediated HR DNA repair [23]. Two in vitro studies followed demonstrating sensitivity of PTEN-deficient cells to PARP inhibition [24, 25]. Compared to the work done in ovarian cancer, the basic science support is less robust; therefore, only a handful of phase I and phase II clinical trials are active in uterine cancer. A phase 0 trial, the Preoperative Olaparib Endometrial Carcinoma Study (POLEN, NCT 02506816) will be recruiting patients to assess the biological impact of PARP inhibition during the period of time between diagnosis and surgery.

The role and application of PARP inhibition in malignancies of the cervix, vagina, and vulva has yet to be clearly determined. To date, no clinical trials have been conducted in the treatment of vaginal and vulvar cancers using PARP inhibition. There is some preclinical evidence of the mediation of PARP activity by HPV infection [26–28]. Specifically, in a series of head and neck squamous cell carcinomas, repair of DNA DSB was significantly delayed in HPV+ tumors, which correlated with increased in vitro sensitivity to veliparib [28]. Veliparib is currently under study in a phase I and phase II trial in advanced cervical cancer (see below). Olaparib is being investigated in a phase I trial in recurrent/refractory cervical cancer (NCT01237067), which seeks to determine the safety and efficacy of combined carboplatin and olaparib on different doses and schedules in women with recurrent/refractory cervical cancer, as well as uterine, ovarian, and breast cancer and in men with metastatic breast cancer and BRCA mutation.

Olaparib
Olaparib (AZD2281) is an oral PARP-1 and PARP-2 inhibitor manufactured by AstraZeneca that was approved for fourth-line treatment of recurrent BRCA-related ovarian cancer in December 2014. In the original dose-escalation phase I clinical trial published in 2009, Fong et al. reported a 47% overall response rate (ORR) and a 64% clinical benefit rate (CBR—tumor marker or radiologic response or stable disease) of ≥4 months in a total of 60 heavily pretreated patients with refractory breast, ovarian, or prostate cancer, 22 of whom had BRCA1 and BRCA 2 germline mutations [29•]. Additionally, a supplemental expansion cohort of 50 patients with BRCA1/2 mutations was later recruited into the study. The maximum-tolerated dose (MTD) of olaparib was 400-mg BID after reversible dose-limiting toxicity was seen in one of eight patients receiving 400-mg BID dosing (grade 3 mood alterations and fatigue) and two of five patients receiving 600-mg BID dosing (grade 4 thrombocytopenia, grade 3 somnolence). The study also demonstrated that CBR was associated with platinum sensitivity (23% in platinum refractory, 45% in platinum resistant versus 69% in platinum sensitive patients) [30].

After these initial findings, two phase II proof-of-concept trials were initiated [31, 32], followed by phase II trials of olaparib for maintenance therapy and in combination with cytotoxic chemotherapy agents. The phase II proof of concept trial of olaparib in gynecologic cancer patients found a dose-response to olaparib in 57 women with BRCA-related recurrent ovarian cancer (ORR 33% with 400 mg BID vs 13% with 100 mg BID) [32]. Grade 3/4 toxicities in the higher dose group were two cases of nausea and one case each of fatigue and anemia. At 16 weeks of treatment, prevalence of progression was higher in the lower dose group (65% vs 33%). In the first phase II trial confirming olaparib efficacy in sporadic ovarian cancers, Gelmon et al. reported an ORR in seven of 17 cases (41%) of BRCA-mutated ovarian cancer, compared to 11 of 46 cases of sporadic cases (24%) [33]. Platinum sensitivity also correlated with response in this study: among sensitive
patients, 60% of patients with mutations and 50% of patients without mutations responded while only 4% of patients with platinum resistant and non-BRCA-associated cancer had a response.

Olaparib monotherapy has been tested in two randomized phase II studies, directly against pegylated liposomal doxorubicin (PLD) in the recurrent germline BRCA mutation population and against placebo in the maintenance setting in the platinum-sensitive population [34, 35]. The first study showed comparable PFS of PLD (50 mg/m² IV q28 days) compared to two doses of olaparib (200-mg BID or 400-mg BID continuously) in the recurrent BRCA-deficient ovarian cancer. Though the differences were not statistically significant, ORR also differed between the three groups (18, 25, and 31%, respectively). In the maintenance study (also known as Study 19), patients received placebo or olaparib 400-mg BID after platinum-based chemotherapy. Compared to placebo, olaparib conferred a 3.6-month advantage in PFS (8.4 versus 4.8 months), and among BRCA mutation carriers, this benefit in PFS was the greatest (11.2 months compared to 4.3 months) [36••]. Patients with sporadic cancers had a more modest improvement in PFS of 7.4 months versus 5.5 months. There was no benefit in overall survival (OS).

The multicenter phase II study that led to the FDA approval of olaparib as fourth-line therapy for women with BRCA-related ovarian cancer is known as Study 42 by Kaufman et al. [37••]. The trial enrolled patients with a germline BRCA1/2 mutation and recurrent cancer with three or more prior lines of chemotherapy and administered olaparib 400-mg BID; their primary endpoint was tumor response rate. A total of 298 patients received treatment, including 193 women with ovarian cancer who achieved an ORR of 34% and a median response duration of 8 months. The study further strengthened the utility of olaparib among this heavily pretreated population. On December 19, 2014, the FDA granted accelerated approval to olaparib as fourth-line therapy for women with recurrent ovarian cancer and germline BRCA mutations as determined by the FDA-approved companion diagnostic BRCA analysis test by Myriad, the first approval of its kind for a laboratory-developed test [38].

Given the efficacy and safety of olaparib in phase II trials, several phase III clinical trials of olaparib in ovarian cancer are underway (Fig. 3). The study of Olaparib in ovarian cancer (SOLO) studies are all in patients with BRCA-deficient cancers. SOLO1 (NCT01844986), the follow-up to Study 19, is the first double-blind, placebo-controlled phase III trial in newly diagnosed ovarian cancer of olaparib maintenance post-platinum-based chemotherapy. With an estimated enrollment of 344, it randomizes patients 2:1 to olaparib to placebo, and its primary endpoint is PFS. SOLO2 (NCT01874353) is the follow-up study to Study 41 investigating maintenance olaparib versus placebo in recurrent platinum-sensitive disease. In a similar fashion, patients will be randomized 2:1 to olaparib versus placebo with an estimated enrollment of 264. Primary endpoint is PFS. SOLO3 (NCT02282020) will assess olaparib monotherapy versus physician’s choice single agent cytotoxic chemotherapy in recurrent platinum-sensitive disease. Its estimated enrollment is 411, and primary endpoint is PFS. Notably, the FDA approval of olaparib for use in this population is contingent upon the results of SOLO3.

Olaparib has been studied in combination with chemotherapeutic agents, and a number of preclinical studies have demonstrated synergy between PARPi and platinum-containing compounds [39–41]. However, the overlapping myelosuppressive toxicities of olaparib and chemotherapeutics have limited the achievement of full-dose chemotherapy in these phase I trials [42–47]. Phase II studies reinforced the activity of PARPi in patients with BRCA mutations. Lee et al. studied olaparib in combination with carboplatin for up to 8 cycles of therapy, followed by single-agent maintenance olaparib until progression, in 45 cases of either breast or ovarian cancer and germline BRCA mutation [47]. For women with ovarian cancer, the ORR among the platinum-sensitive cases was 71%, platinum-resistant cases was 25%, with an overall ORR of 44%; an additional 41% had stable disease from 3 to 25+ months. Olaparib was studied in a randomized phase II trial, known as Study 41, in combination with carboplatin/paclitaxel compared to chemotherapy alone in platinum-sensitive recurrent ovarian cancer [48•]. PFS was significant for the combination arm (12.2 versus 9.6 months) with the greatest benefit for those with known BRCA mutations. When studied in combination with carboplatin and metronomic paclitaxel in 12 patients with at least three prior therapies, four had complete response, four had partial response or stable disease, and two had progression [49]. A planned expansion of this phase Ib study will recruit an additional 40 patients.

Olaparib has also been combined with antiangiogenic agents. Homologous recombination can be suppressed by hypoxia through the regulation of hypoxia-inducible factor 1α and nuclear factor κB (NF-κB) [50]. Bevacizumab (10 mg/kg IV q14 days) was combined with dose escalations of continuous olaparib
A. **SOLO1 (phase III)**

Upfront maintenance treatment, randomized within 8 weeks of last dose of chemotherapy

- Newly diagnosed stage III/IV high grade ovarian cancer – completed upfront platinum-based chemotherapy, +BRCA 1/2 mutation
  - Olaparib 300mg PO BID
  - Placebo PO BID
  - Until disease progression

B. **SOLO2 (phase III)**

For chemotherapy immediately prior to randomization, patients with ≥4 cycles

- Recurrent platinum sensitive high grade serous ovarian cancer – completed ≥2 prior lines of platinum-based therapy +BRCA 1/2 mutation
  - Olaparib 300mg PO BID
  - Placebo PO BID
  - Until disease progression

C. **SOLO3 (phase III)**

Comparison between olaparib and physician’s choice single agent chemotherapy

- Recurrent platinum sensitive high grade serous ovarian cancer – completed ≥2 prior lines of platinum-based therapy +BRCA 1/2 mutation
  - Olaparib 300mg PO BID
  - Single agent paclitaxel OR Topotecan OR Pegylated liposomal doxorubicin OR Gemcitabine
  - Until disease progression

D. **ARIEL2 (phase II)**

Identification of a HRD molecular signature that correlates with RR, to be applied in ARIEL3

- Recurrent platinum sensitive high grade serous ovarian cancer – completed ≥1 prior lines of platinum-based therapy, disease progression >6mo after penultimate regimen
  - Rucaparib 600mg PO BID
  - Until disease progression

E. **ARIEL3 (phase III)**

For platinum-based chemotherapy immediately prior to randomization, stratified based on 3 HRD groupings

- Recurrent platinum sensitive high grade ovarian cancer – completed ≥2 prior lines of platinum-based therapy, no more than 1 non-platinum agent, disease progression >6mo after penultimate regimen
  - Rucaparib 600mg PO BID
  - Placebo PO BID
  - Until disease progression

F. **NOVA (phase III)**

Enrolled <8 weeks after last platinum-based therapy, food-effect sub-study on the effect of high-fat meal on the PK of niraparib

- Recurrent platinum sensitive high grade serous ovarian cancer or known gBRCA mutation – completed ≥2 prior lines of platinum-based therapy +/- BRCA 1/2 mutation
  - Niraparib 300mg PO QD
  - Placebo PO QD
  - Until disease progression

**Fig. 3.** Schema of phase II and phase III trials of PARP inhibitors in patients with ovarian cancer. **ARIEL** Assessment of Rucaparib in ovarian cancEr triaL. **BRCA1** breast cancer 1 gene. **BRCA2** breast cancer 2 gene. **GOG** gynecologic oncology group. **HRD** homologous recombination deficicncy. **NOVA** niraparib ovarian trial. **PARP**, poly(ADP-ribose) polymerase. **SOLO** study of OLaparib in ovarian cancer.
in the phase I setting with 12 ovarian cancer patients. The MTD of olaparib was 400 mg BID, and toxicities were mild, limited to grade 1/2 nausea and fatigue [51]. Trials of olaparib and cediranib, an oral vascular endothelial growth factor receptor 2 inhibitor, have shown encouraging results. The initial phase I trial established the MTD of olaparib to be 200-mg BID in combination with cediranib 30 mg daily [52, 53]. This dose was tested in a randomized phase II study of 90 women with recurrent platinum-sensitive high-grade serous or endometrioid ovarian cancer [54]. Combination therapy was associated with significantly improved PFS (9 months versus 17.7 months) and ORR (48 % versus 80 %). OS data is not mature, but at 24 months, 81 % of combination patients are alive compared to 65 % of olaparib only patients. Grade 3/4 toxicities more common in the combination group were fatigue, diarrhea, and hypertension. These promising results have led to the development of two phase III trials using combination olaparib and cediranib therapy: a direct comparison to standard chemotherapy in the platinum-resistant or -refractory setting (NCT02502266) and as compared to olaparib monotherapy or standard platinum-based chemotherapy in the platinum-sensitive setting (NCT02446600).

**Veliparib**

Veliparib (ABT-888) is also an oral PARP-1 and PARP-2 inhibitor made by AbbVie that has been extensively studied as a single agent as well as in combination with chemotherapy in gynecologic cancers. Mechanistically, it appears to have an inferior ability to trap PARP-1 and PARP-2 at the site of SSB compared to both olaparib and niraparib [55]. The toxicity profile of veliparib is similar to olaparib, with nausea, fatigue, and myelosuppression seen most commonly with monotherapy [56]. Velaparib has been studied in combination with topotecan, doxorubicin, and cyclophosphamide with variable response [57–59].

In GOG 280, a phase II study on veliparib monotherapy in the treatment of persistent or recurrent BRCA-related ovarian cancer, patients were enrolled to receive veliparib 400 mg BID [56]. Fifty patients with three or fewer prior regimens were enrolled; ORR was 26 % with median PFS of 8.2 months. When stratified on platinum sensitivity, platinum-resistant patients had ORR of 20 % compared to 35 % in platinum-sensitive disease. Grade 3/4 toxicities were limited to one case of grade 4 thrombocytopenia and the following grade 3 events: fatigue (n = 3), nausea (2), leukopenia (1), neutropenia (1), dehydration (1), and increased ALT (1). Almost 50 % of patient experienced grade 2 nausea and 25 % had grade 2 fatigue.

Veliparib is currently undergoing investigation in the phase III arena through GOG PARTNERS 3005 (NCT02470585, Fig. 3f). In a trial schema identical to GOG 218 with bevacizumab [60, 61], veliparib will be studied in the primary setting, in a 1:1:1 randomized, double-
blinded trial given concurrently with standard carboplatin/paclitaxel chemotherapy with and without veliparib maintenance therapy. The trial began accrual in October 2015. The primary endpoint is PFS, which was be explored in the general population as well as those with BRCA mutations.

As one of the few PARPi currently under investigation in cervical cancer, veliparib is being used in combination with topotecan in persistent or recurrent cervical cancer (NCT01266447), as well as a phase 1/2 trial in combination with cisplatin and paclitaxel in advanced or recurrent disease (NCT01281852).

Niraparib
Niraparib (MK4827) is also an oral inhibitor of PARP-1 and PARP-2, the first whose pharmacokinetics allows for once daily dosing. Niraparib entered clinical studies in 2008 with a phase I study of advanced solid tumors, including high-grade serous ovarian cancer, with enrichment for those with BRCA mutations [62]. A modified 3 + 3 design was utilized for dose escalation in 100 patients to determine the 300-mg/day MTD. The most common toxicities were predominantly grade 1/2 anemia (48 %), nausea (42 %), fatigue (42 %), thrombocytopenia (35 %), anorexia (26 %), neutropenia (24 %), and vomiting (20 %). The ORR of BRCA-related ovarian cancer cases (\(n = 20\)) was 40 %, with a median response duration of 387 days. Among the BRCA-deficient ovarian cancer cases, 10 had platinum sensitive disease, and these patients had an ORR of 50 % and median response duration of 431 days. Currently, there are no completed phase II studies of this drug in ovarian cancer.

Tesaro, Inc. is presently recruiting for a phase II trial for women with recurrent high-grade serous ovarian cancer who have completed at least three previous chemotherapy regimens (NCT02354586). It also recently completed accrual for the NOVA trial, a phase III double-blind, placebo-controlled, 2:1 randomized trial of maintenance niraparib versus placebo in patients with recurrent platinum-sensitive high-grade serous ovarian cancer or known to have a germline BRCA mutation (NCT01847274, Fig. 3c) [63•]. NOVA enrolled 490 participants, and its primary objective is PFS. Additionally, the trial is evaluating the effect of a high-fat meal on the pharmacokinetics of a single 300-mg dose of niraparib in patients with ovarian cancer [64]. A 2-treatment (fed versus fasting), 2-way crossover design was used to evaluate the effect of food on PK parameters. Sixteen subjects were enrolled in the food effect cohort, and each subject received two separate 300-mg doses of niraparib, one each in a fasting and a fed state. Data from the NOVA are expected in 2016.

Rucaparib
Rucaparib (CO338, AGO14699, and PF01367338) is another PARP-1 and PARP-2 oral inhibitor that has entered into clinical trial testing [65, 66]. The phase I study of oral rucaparib tested doses of from 40 to 840-mg BID and recommended the phase II dose of single-agent rucaparib to be 600-mg BID. Rucaparib has demonstrated durable responses greater than 6 months in both platinum-sensitive and platinum-resistant ovarian cancer [66]. Preliminary results of the phase II trial in 17 women with germline-BRCA mutations and ovarian cancer reported at the 2014 European Society of Medical Oncology meeting showed an ORR of 82 % and a disease control rate of 93 % at 12 weeks [67]. The most common treatment-related toxicity occurring in 15 % of patients were nausea, asthenia, vomiting, transient transaminitis, and anemia. Rucaparib was granted US FDA Breakthrough Therapy designation on April 6, 2015, based on the interim results of the Kristeleit phase II trial and the ARIEL2 study from the Assessment of Rucaparib in Ovarian Cancer Trial (ARIEL group) by Clovis Oncology, Inc.

The ARIEL2 study evaluated rucaparib in a pivotal phase II prospective biomarker trial in high-grade ovarian cancer focused on identification of a molecular signature to predict clinical benefit for patients with platinum-sensitive disease with at least one prior regimen (NCT01891344, Fig. 3c) [20••]. Interim results were presented at the 2015 Annual Meeting on Women’s Cancer of the Society of Gynecologic Oncology [68] with final results available at the 2015 Annual Meeting of American Society of Clinical Oncology [20••]. The primary objective of this single-arm, open-label study, was to identify a molecular HRD signature in ovarian cancer associated with clinical benefit from rucaparib treatment. Known germline BRCA-related ovarian cancers were capped in this study with the 208 patients divided into the following distribution of homologous repair-deficient molecular subgroups: BRCA-mutated 20 %, BRCA-like 40 %, biomarker negative 34 %, and unclassified 6 %. Tumors with RAD51C genetic alterations had high genomic LOH and demonstrated a BRCA-like phenotype. Overall response rate was highest in the BRCA-mutated (82 %), followed by the BRCA-like group (45 %), with 21 % of the biomarker negative group showing response based on RECIST + CA-125. Median PFS was 9.4, 7.1, and 3.7 months, respectively. Grade 3/4 toxicity was mostly limited to anemia (16 %) and transient transaminitis (11 %). An expansion cohort is now currently recruiting in ARIEL2 Part 2 to include patients with three or more prior chemotherapy regimens. The results of ARIEL2 are potentially practice changing as the data provide proof of
concept for the development of biomarker assays for HRD, which are now under development (see below). Using the predictive HRD assay prospectively determined in ARIEL2, the follow-up study, ARIEL3, a randomized, phase III trial, will stratify patients who have received two or more prior platinum regimens with platinum-sensitive disease into the three HRD groupings and investigate the use of rucaparib as maintenance versus placebo (NCT01968213, Fig. 3d).

**Talazoparib (BMN 673)**

Talazoparib is an oral PARP-1 and PARP-2 inhibitor [69] manufactured by BioMarin that has undergone phase I testing in an open-label study of once-daily treatment in patients with advanced or recurrent solid tumors [70]. The MTD of 1000 µg/day was determined in this study of 39 patients. Dose-limiting thrombocytopenia occurred in one out of six patients and two out of five patients at the 900 and 1100 µg/day, respectively. Grade 3/4 adverse events included anemia, neutropenia, and thrombocytopenia. Twenty-three patients were enrolled with either ovarian or primary peritoneal cancer, and 17 of these patients had a germline BRCA mutation. RECIST and/or CA-125 responses occurred at doses ≥ 100 µg/day in 11 out of 17 BRCA-related ovarian or peritoneal cancers. Currently, there is a phase III study testing Talazoparib in patients with metastatic breast cancer and a phase II trial sponsored by the National Cancer Institute for women with deleterious BRCA 1/2 mutation-associated ovarian cancer who had had prior PARP inhibitor treatment (NCT02326844).

Talazoparib is also being studied in uterine cancer in the PARP inhibitor for Inoperable Advanced eNDometrial cAn-cer phase II trial (PANDA, NCT02127151). This study is not yet open for recruitment. Patients with inoperative advanced or recurrent endometrial cancer with no more than one prior line of systemic cancer therapy will be given daily BMN 673 to determine primary outcome measure PFS.

**Future Directions: Development of a Homologous Recombination Deficiency Assay**

The role homologous recombination assays will continue to play a role in the treatment of women with gynecologic malignancies, especially in those with ovarian cancer because of the therapeutic advantage generated by PARP inhibition in tumors with HRD. The ARIEL2 trial incorporated the translation piece through studying LOH, a marker of HRD and measurable using next-generation sequencing. The investigators developed an algorithm for LOH and confirmed correlation with response rates and PFS. Other HRD assays are in development. At the 2014 EORTC-NCI-AACR Symposium on Molecular Targets and Cancer Therapies Meeting, Haluska et al. presented a HRD score obtained from performing HRD analysis on 106 high-grade ovarian tumors with known responses to niraparib [71]. Using genome-wide SNP data, tumor xenografts were analyzed using three algorithms: LOH, telomeric allelic imbalance (TAI), and large-scale state transitions (LST). The xenografts were again treated with niraparib, and the HRD score was calculated as the sum of the LOH + TAI + LST scores. High HRD scores were correlated with in vivo response to niraparib and BRCA deficiency, and low HRD scores were associated with niraparib resistance or lack of in vivo efficacy. Both ARIEL3 and the NOVA study are developing HRD assays as part of the phase III clinical trials.

Genetic sequencing technology through the use of microarrays and next-generation sequencing, as well the data from The Cancer Genome Project, has made these analyses possible. They provide methods to comprehensively capture the diverse ways HRD may manifest itself outside of
traditional BRCA 1/2 mutation analyses to incorporate copy number variability to LOH, TAI, and LST scores [72–75]. Others have suggested a diagnostic assay integrating a genomic scar-based biomarker with a marker of PARPi resistance because tumors can undergo events that restore HR [72]. Once HR is restored, the tumors can then be misclassified as HRD and thereby inaccurately sensitive to PARP inhibitors and other anti-cancer therapies. There are limitations of these scores, however, as our understanding of genomics and targetable deficiencies in DNA repair mechanisms has been only recently bolstered by advancements in single nucleotide polymorphism microarrays and high-throughout sequencing technology.

The incorporation of prospective assay data into clinical trials is the first step towards utilizing the HRD phenotype in personalized treatment plans for patients. Furthermore, the FDA has recently begun requiring that all new drug applications be accompanied to the market by a biomarker that predicts its effectiveness. Validation studies will undoubtedly be required and forthcoming; however, this represents an exciting area of research with significant practice changing potential.

Conclusion

PARP inhibitors represent an exciting new targeted treatment option in the management of BRCA-related ovarian cancer and may have significant role in the treatment of other gynecologic cancers. The approval of olaparib as fourth-line therapy for women with germline BRCA mutations and ovarian cancer is a perfect example of applying translational research to drive forward another method by which patients may potentially achieve a durable remission from disease. For patients with BRCA mutations especially, ovarian cancer can become a life-long chronic disease that will require multiple different treatment regimens. The genomic science used in studies like ARIEL2 also provides a window into the future of research in personalized cancer care and may eventually determine how clinicians will triage patients to therapy. PARP inhibitors have already demonstrated impressive responses in phase II clinical trials and provide a necessary option for patients with recurrent disease and could also possibly serve as a viable option for newly diagnosed disease.

Compliance with Ethical Standards

Conflict of Interest
The authors declare that they have no competing interests.

Human and Animal Rights and Informed Consent
This article does not contain any studies with human or animal subjects performed by any of the authors.
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