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Homologous repair deficiencies and current insights in clinical evaluation of PARP inhibitors in prostate cancer

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Summary Homologous repair deficiency is a clinically relevant molecular aberration in prostate cancer. The goal of this short review is to summarize the study landscape of treatments targeting these aberrations through discussion of the most relevant clinical trials. Due to its shortness, this review does not claim to be exhaustive and a major focus is being laid on PARP inhibitors in clinical development for prostate cancer.

Keywords Prostate cancer \cdot BRCA1 \cdot BRCA2 \cdot PARP \cdot Homologous repair deficiency \cdot Cancer treatment

Abbreviations

Inductions	
ADT	Androgen deprivation therapy
AR	Androgen receptor
CRPC	Castration-resistant prostate cancer
DDR	DNA damage repair
HRR	Homologous recombination repair
PARPi	Poly(ADP-ribose) polymerase inhibitors
PCa	Prostate cancer
PFS	Progression-free survival

Introduction

Prostate cancer (PCa) is the most frequently diagnosed noncutaneous cancer in men and was ranked third regarding incidence among all cancers in the European Union in 2018 [1]. Although prostate-specific antigen (PSA) screening has facilitated early detection, treatment of advanced metastatic PCa remains challeng-

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T. Topakian, MSc thais.topakian@meduniwien.ac.at ing as the disease is highly heterogeneous, exhibiting strong histopathological, genetic and clinical diversity [2]. Since PCa growth is mainly dependent on androgen receptor (AR) expression, androgen deprivation therapy (ADT) targeting AR or AR ligand binding remains one of the standards of care to date. However, despite initial response, patients often develop resistance to ADT, leading to incurable castration-resistant prostate cancer (CRPC).

The endeavor of profiling the molecular and genetic landscape of PCa and searching for novel predictive biomarkers that specify and optimize therapy has recently led to the discovery of multiple genomic alterations. Next generation sequencing studies of prostate tumors revealed a high number of recurrent gene mutations interfering with homologous recombination repair (HRR). Among others, the Cancer Genome Atlas (TCGA) Research and American Association for Cancer Research networks have identified HRR gene alterations in 19-23% of metastatic CRPC (mCRPC) samples [3, 4]. Analyzing the TCGA prostate cancer database, Kim et al. found mutations and copy-number variations of DNA damage repair (DDR) genes in 30% of localized (nonmetastatic) PCa indicating that DDR alterations might occur early during disease progression [5]. Reported prevalence of mutations included HRR genes such as cyclin-dependent kinase 12 (CDK12), RAD51C, breast cancer 1 (BRCA1) or ataxia telangiectasia mutated (ATM). Interestingly, all studies found breast cancer 2 (BRCA2) to be the most commonly mutated gene.

Germline BRCA2 mutations are associated with a high risk of developing cancer across many entities with a recently reported risk of 19–61% for PCa by age 80 [6]. The concept of "synthetic lethality", initially described in drosophila by Dobzhansky as a state where simultaneous loss-of-function of two genes results in a nonviable phenotype, led to the

discovery of the synthetically lethal interplay between BRCA1 or BRCA2 with poly(ADP-ribose) polymerase (PARP1). In this case, the DDR deficiencies caused by BRCA1 or 2 loss-of-function lead to cell death if the compensatory DNA repair pathway via PARP1 is inhibited. PARP inhibitors (PARPi) were, thus, developed and approved for the treatment of breast and ovarian cancers. Furthermore, emerging evidence indicated a strong relationship between BRCA1/2 deficiencies with sensitivity to DNA-damage causing platinum-based chemotherapy in breast and ovarian, but also in PCa [7]. A synthetic lethal mode-of-action was also claimed for radium-223 treatment in mCRPC with bone metastasis and homologous repair gene defects (HRD) [8]; however the mechanistic explanation remained most reasonable for PARP, which, as components of base excision repair (BER), restore damaged DNA in case of HRR gene loss.

Beside the synthetic lethality of PARPi in the presence of HRR aberrations, numerous studies highlight further effects of PARPi activity in PCa. Gui et al. demonstrated that selectively targeting PARP2 blocked its interaction with FOXA1 and attenuated AR-signaling [9]. Conversely, Li et al. identified an AR and c-myb upregulated HRR gene signature that correlated with castration resistance, metastasis, relapse and reduced survival and demonstrated that targeting AR by enzalutamide led to downregulation of this specific gene set [10]. These findings suggest PARPi as an alternative therapeutic approach for ADT irrespective of HRR status and indicate a potential combinatory effect with antiandrogens such as apaor enzalutamide. Several PARPi have been or are currently evaluated in clinical trials for the treatment of PCa, including olaparib, rucaparib, talazoparib, veliparib and niraparib. While none of them are currently approved within Europe, numerous trials already show promising activity.

The single-arm phase II trial TOPARP-A recruited 50 men with mCRPC, who had previously undergone chemotherapy and/or ADT, for the treatment olaparib irrespective of their HRR status. Response was defined according to RECIST, with a decline in PSA levels by at least 50% or a reduction of counted circulating tumor cells to less than 5 per 7.5 mL blood. Of 16 patients who responded to olaparib, 88% were identified to carry HRR mutations. Within responders, HRR mutation carriers exhibited significantly longer radiologic progression-free survival (PFS, 9.8 vs. 2.7 months). Interestingly, durable responses for more than a year were achieved for all 8 patients with BRCA1/2 aberrations [11].

To further evaluate antitumor activity of olaparib stratified by specific DDR mutations, the TOPARP-B phase II trial analyzed responses of 98 men genomically preselected for putative pathogenic mutations in DDR pathways. All patients had received prior docetaxel and the majority prior enzalutamide or abiraterone. Striking for this pretreated collective were the response rates of 57 and 41% in the 400 and 300 mg treatment arms, respectively. Again, BRCA1/2-mutated patients represented the subgroup with strongest response and longest median radio-graphic PFS (8.3 months) [12].

The still ongoing phase III trial PROfound is investigating 387 men with mCRPC and at least one mutation in DDR genes subjected to treatment with olaparib or physician's choice of enzalutamide or abiraterone. Two cohorts were defined based on mutations with BRCA1/2 and ATM (cohort A) and other alterations (cohort B). In an interim analysis presented at ESMO congress last year, olaparib treatment resulted in significantly prolonged radiographic progression-free survival in cohort A (HR 0.34) and the overall population (cohort A+B, HR 0.49) as well as a trend for prolonged overall survival not reaching the previously determined levels of statistical significance for cohort A. Most likely, heavy cross-over between treatment arms hindered the observation of a statistically significant difference in OS for cohort A in the interim analysis (HR 0.64, n.s.), but could be met in the final analysis [13].

Besides olaparib monotherapy, combinatorial effects of olaparib with antihormonal agents are being investigated. Due to its brevity, this short review cannot cover all currently ongoing trials. Of note, however, Clarke et al. assessed the efficacy and tolerability of abiraterone plus olaparib and compared it to abiraterone with placebo. A total of 71 patients were assigned to each cohort, of which 14 and 11% were identified to have HRR deficiencies in the olaparib and placebo arm, respectively. Radiographic PFS was significantly prolonged in the abiraterone plus olaparib group regardless of HRR status (13.8 vs 8.2 months), again with more adverse effects in the olaparib arm [14].

Another promising PARP inhibitor, talazoparib, is being evaluated for CRPC patients with DDR defects. Described as being the most potent inhibitor in trapping PARP enzymes at single-strand breaks in vitro, talazoparib provides high metabolic stability in mice and is orally bioavailable [15]. The phase II study TALAPRO-1 presently examines efficacy and safety of talazoparib (1mg/day) stratified by DDR mutations of 11 genes. Interim analyses showed highest efficacy of talazoparib treatment for BRCA1/2 and PALB2 deficient CRPC patients with ORR of 50 and 33%, respectively. Interestingly, the BRCA1/2 cohort had the longest radiographic PFS with 8.2 months compared to patients with other DDR alterations [16, 17]. In addition, the currently ongoing TALAPRO-2 phase III study is examining the combination therapy of talazoparib and enzalutamide. Recently presented intermediate results of TALAPRO-2 indicated that 0.5 mg talazoparib per day in combination with background enzalutamide treatment in therapy-naïve patients had a manageable safety profile and led to a prominent decline in PSA levels after a median treatment duration of 11 weeks [18].

The high expectations for PARPi efficacy in CRPC treatment are further reflected by 10 currently recruiting clinical trials with the aim to examine rucaparib in combination with other anticancer treatments or as monotherapy. Preliminary results of the phase II study TRITON2 demonstrated high ORR (43.9%) with durable response over 24 weeks for the majority of mCRPC patients carrying BRCA mutations. However, only poor response could be achieved for patients with other mutations than BRCA [19]. Similar interim results were presented for the GALAHAD phase II study with highest ORR (41%) and rPFS of 8.2 months for BRCA mutated CRPC patients treated with the PARPi niraparib [20].

Besides combinatorial therapies of PARPi, antihormonal agents and chemotherapy, the clinical research focus has recently been placed on the question whether PARP inhibition together with immune checkpoint inhibition would show antitumor activity. Several preclinical studies confirmed that PARP inhibition triggers STING activation and promotes type I immunity, which may benefit in increased inflammation and tumor immune infiltration [21]. To our knowledge, there are two currently recruiting phase II trials investigating this topic in CRPC. With final results pending, these two studies are currently assessing the combination of PD-L1 inhibitors avelumab with talazoparib and durvalumab with olaparib, respectively. The latter combination was tested in 17 CRPC patients with preliminary response rates of 53% for those with DDR gene mutations. The treatment increased immune response in some patients by enhancing dendritic cell maturation, CD4⁺ and CD8⁺ T-cell activity, which correlated with prolonged rPFS [22].

Take home message

PARP inhibitors are promising novel targeting agents for multiple cancer entities. All PARP inhibitors under development have shown significant clinical benefits for BRCA1/2 mutation carriers suffering from prostate cancer and partially for patients with other homologous recombination repair (HRR) defects. Altogether, data from currently ongoing trials are very likely to change the landscape and treatment standard of HRR-deficient prostate cancer and make DNA-damage repair gene mutation status testing required for all prostate cancer patients in the future.

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Conflict of interest M. Marhold and T. Topakian declare that they have no competing interests.

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