

Deep exploration of PARP inhibitors in breast cancer: monotherapy and combination therapy Journal of International Medical Research 49(2) 1–15 © The Author(s) 2021 Article reuse guidelines: sagepub.com/journals-permissions DOI: 10.1177/0300060521991019 journals.sagepub.com/home/imr



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Abstract

Objective: Nearly 5% of patients with breast cancer carry germline *BRCA* mutations, which are more common in triple-negative breast cancer (TNBC). Previous clinical trials demonstrated the therapeutic efficacy of poly (ADP-ribose) polymerase inhibitors (PARPis) against *BRCA*-mutated metastatic breast cancer. The current study conducted a systemic review and meta-analysis of the clinical efficiency and safety of PARPis, either alone or combined with chemotherapy, in patients with TNBC.

Methods: We searched PubMed, EMBASE, and ClinicalTrials.gov to identify randomized controlled trials comparing PARPi therapy with chemotherapy, and comparisons of chemotherapy plus PARPis with chemotherapy alone were included. The study endpoints included the clinical response, progression-free survival, and adverse event rates.

Results: PARPi therapy was revealed to improve progression-free survival in patients with advanced breast cancer, either alone or in combination with chemotherapy. Subgroup analysis illustrated that patients with mutant *BRCA1* and mutant *BRCA2* and those who had not been treated with platinum-based agents could specifically benefit from PARPis.

Conclusion: PARPi monotherapy can significantly improve clinical outcomes in patients with advanced breast cancer, especially those with TNBC, those who had not previously received platinum therapy, and those with mutant *BRCA1/2*. PARPis combined with chemotherapy represent new treatment options for patients with advanced cancer.

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Introduction

Breast cancer is one of the most common malignant tumors in women. According to National Cancer Institute statistics, nearly 252,710 women were diagnosed with breast cancer in 2018, and 40,610 women died of the disease.¹ Advanced metastasis is an important factor threatening the lives of patients. Chemotherapy, endocrine therapy, radiotherapy, and targeted therapy are the primary treatments for patients with advanced breast cancer. Currently, a widely used targeted therapy in clinical practice is anti-HER2 therapy, including HER2 antibodies and tyrosine kinase inhibitors (TKIs).² However, triple-negative breast cancer (TNBC), which accounts for approximately 15% of breast cancers, lacks therapeutic targets.³ Chemotherapy has always been the main systemic treatment for TNBC. Because of the lack of targets for the three causes of breast cancer and the lack of targeted drugs, patient prognosis is poor, prompting clinicians to develop significant efforts to discovering treatable molecular targets.³ Interestingly, patients **TNBC** often germline with carry BRCA (gBRCA) mutations. $\overset{4,5}{\text{Research}}$ data from Chinese patients with breast cancer revealed that the frequency of BRCA gene mutation in TNBC was approximately 11%.⁶

BRCA1 and *BRCA2* are key tumor suppressor genes for homologous recombination (HR) repair. The proteins encoded by these genes are involved in the repair of DNA double-strand breaks, cell growth,

and prevention of the abnormal cell division that leads to the occurrence of tumors. Poly (ADP-ribose) polymerase (PARP), as a DNA break sensor, is activated after DNA damage, and it recognizes and binds to the DNA break site and participates in the repair of DNA single-strand damage in tumor cells. For tumors with abnormal HR repair function, PARP inhibitors (PARPis) suppress PARP enzymatic activity and increase the formation of PARP-DNA complexes, leading to the repair of DNA damage in tumor cells and promoting apoptosis.⁷ In prior research, PARPis enhanced the efficacy of radiotherapy, alkylating agents, and platinum-based chemotherapy by inhibiting the repair of DNA damage in tumor cells and promoting apoptosis.8 Since 2003, clinical studies on the utilization of PARPis in solid malignancies have been increasingly reported. Breast and ovarian cancers, which are most frequently associated with BRCA mutations. were demonstrated to respond to PARPis.^{9,10} Several PARPis, such as olaparib, rucaparib, and niraparib, have been approved by the US Food and Drug Administration (FDA) as maintenance therapies for recurrent ovarian cancer.^{11–14} A meta-analysis of the efficacy of PARPis as maintenance treatments for platinumsensitive recurrent ovarian cancer suggested that these drugs were effective regardless of BRCA mutation status, and substantial improvements of progression-free survival (PFS) were observed for patients with germline mutations.¹⁵ Currently, several clinical studies on PARPis for advanced breast cancer are underway. The randomized phase III trials OlympiAD¹⁶ and EMBRACA¹⁷ compared the effects of olaparib and talazoparib with doctor-selected chemotherapy in patients with gBRCAmutant, HER2-negative breast cancer. PFS, as the primary endpoint, was significantly prolonged in the PARPi group. The FDA approved the two drugs for the clinical treatment of patients with gBRCA1/2mutant, HER2-negative metastatic breast cancer who had previously received chemotherapy. BRCA mutation is an important therapeutic target for TNBC. Preclinical studies confirmed that PARPis could induce synergistic lethal effects in BRCAmutant tumors. The results of OlympiAD further confirmed the role of PARPi therapy in patients with gBRCA-mutant, HER2negative metastatic breast cancer from a clinical perspective. However, the results from phase II and III randomized controlled trials (RCTs) have been inconsistent.

In this study, we assessed the efficacy of PARPi therapy in patients with advanced breast cancer through a meta-analysis, including subgroup analyses.

Methods

Study search strategy

We first initiated a systemic review of the literature according to the Cochrane and Preferred Reporting Items for Systematic Reviews and Meta-Analyses guidelines.¹⁸ We used several methods to screen the final studies for our research. PubMed, Embase. and ClinicalTrials.gov were searched according to the keywords represented in the titles and abstracts, including "breast "PARP." cancer," "BRCA." "chemotherapy," and This study was not registered with PROSPERO. In total, 1499 articles were screened

In total, 1499 articles were screened using the online databases. Among them,

146 were identified via a manual search of article references. Figure 1 presents the details of the search results. Of the searched studies, 340 were duplicates, and 1100 did not match the study aim based on a comprehensive reading of the titles and abstracts. After removing these articles, we further screened the remaining 69 studies by reading the full text intensively. Consequently, four clinical trials providing sufficient data were included in the metaanalyses.^{16,17,19,20} The characteristics of the included studies are summarized in Table 1.

Study criteria

Studies were eligible for inclusion if they were multicenter phase II or phase III RCTs. The included patients were diagnosed with advanced breast cancer, and they were randomly assigned to treatment with chemotherapy, PARPis, or both. The included studies reported at least one of the following clinical outcomes: response rate, PFS, overall survival (OS), and toxicity. Studies that had only one arm, those designed for neoadjuvant therapy, and those using other targeted therapies were excluded.

Study evaluation and data extraction

The Jadad score was used to assess the quality of each included study. The scoring criteria include the generation of random sequences, blinding procedure, and adscription of withdrawals and dropouts.²¹ We extracted the following data from the included articles: number of patients enrolled, chemotherapy regimen, treatment group, *BRCA1* and *BRCA2* status, hormone receptor and HER2 status, toxicity, and efficacy.



Figure 1. Flow chart of study selection.

Statistical analysis

The clinical data, including the number of patients, clinical efficacy, and toxicity, were extracted from the included articles. PFS was defined as the time from randomization to objective radiologic disease progression. According to the modified Response Evaluation Criteria in Solid Tumors, version 1.1, the clinical response (CR) rate was defined as the sum of the stable disease. complete response, and partial response rates. Quantitative statistical combinations were calculated using Review Manager (version 5.3, Copenhagen: The Nordic Cochrane Centre. The Cochrane Collaboration, 2014) using fixed-effects or random-effects modeling considering the existing study variations. Heterogeneity was quantified using the I^2 statistic. The fixed-effects or random-effects model selection principles were based on the value of I^2 . For $I^2 < 40\%$, which indicated low

heterogeneity, the fixed-effects model was chosen. For $I^2 \ge 40\%$, the random-effects model was chosen. The integrative results were represented as the odds ratio (OR) between groups and the 95% confidence interval (CI). In all analyses, P < 0.05 indicated statistical significance.

Results

All four included trials were open-label, multicenter phase II or III RCTs. All recruited patients were diagnosed with advanced breast cancer. Two studies compared PARPis with standard chemotherapy,^{16,17} and the other two studies primarily evaluated PARPis combined with chemotherapy.^{19,20} The methodological quality of the trials was assessed using the Jadad score (Table 2). The quality scores ranged from 4 to 5, indicating good

Clinical trials	Recruited patients	No. of patients	Design	Studied PARPi	Treatment/arms
Kummar et al., 2016	Adult patients with refractory/metastatic TNBC	45	Open-label, multicenter, randomized phase II study	Veliparib (ABT-888)	A: Cyclophosphamide 50 mg once daily (n = 18) B: Cyclophosphamide 50 mg once daily + veliparib 60 mg dailv in 21-dav cycles (n = 21)
O'Shaughnessy et al., 2014	Adult patients with refractory/metastatic TNBC	519	Open-label, multicenter, randomized phase III study	Iniparib	A: GC alone (n = 258) B: GC + iniparib (at a dose of 5.6 mg/kg body weight) on days 1, 4, 8, and 11 of each 21-day
Litton <i>et al.</i> , 2018	Patients with advanced breast cancer and a germline <i>BRCA1/2</i> mutation	431	Open-label, multicenter, randomized phase III study	Talazoparib	A: Standard single-agent therapy* (n = 287) B: Talazoparib (1 mg once daily) (n = 144)
Robson et <i>al.</i> , 2017	Patients with advanced breast cancer and a germline <i>BRCA1/2</i> mutation	302	Open-label, multicenter, randomized phase III study	Olaparib	A: Standard single-agent therapy* $(n = 97)$ B: Olaparib tablets (300 mg twice daily, $n = 205$)
TNBC. triple-negative	e breast cancer; GC. gemcitabine (10	00 mg/m ² body s	urface area) and carboplatin (at a dos	e equivalent to an area	under the concentration-time curve of

וואסר, נדוףפ-הפצעועים הרפוז כמורפרן סר. פנותנומטוופ (וטטט וווצווו טטט אטוומר מופא מוט גמו טרמי מיטס בקעועמובוו גט מו מופמ עוואסר שיר בטורכווע 2) on days 1 and 8. *Standard single-agent therapy indicates the physician's choice, including capecitabine, eribulin, gemcitabine, or vinorelbine, in continuous 21-day cycles.

Table 1. Characteristics of the included studies.

Clinical trials	Kummar et <i>al</i> ., 2016	O'Shaughnessy et al., 2014	Litton et al., 2018	Robson et al., 2017
Randomization	2	2	2	2
Concealment of allocation	2	2	2	2
Double blinding	0	0	0	0
Withdrawals and dropouts	0	I	I	I
Jadad score ^a	4	5	5	5

Table 2. Jadad scale.

^aMethodological quality of meditative movement studies reviewed using Jadad scoring criteria. The maximum score is 7. Scores of 1 to 3 indicated low quality, whereas scores of 4 to 7 indicated high quality.

quality despite the lack of double-blind studies.

Clinical efficacy of PARPis combined with chemotherapy

Two earlier reports evaluated the clinical efficacy of PARPis plus chemotherapy in patients with advanced breast cancer.^{19,20} We combined the results of these studies and chose PFS and CR rates as the endpoints.

The meta-analysis illustrated that the addition of PARPis to chemotherapy did not increase the CR rate as expected (OR = 0.80, 95% CI = 0.56–1.15, P = 0.22). However, the combination regimens were linked to significantly improved PFS rates (OR = 0.72, 95% CI = 0.62–0.89, P = 0.001). The forest plot illustrated that the addition of PARPis to chemotherapy improved the long-term survival of patients (Figure 2).

Clinical efficacy of PARPi versus chemotherapy

Two included trials reported the result of single-agent PARPi therapy versus standard therapy in patients with advanced breast cancer and gBRCA1/2 mutations.^{16,17}

The results demonstrated that PARPi treatment (olaparib¹⁶ or talazoparib¹⁷) was statistically associated with better CR rates

(OR = 0.44, 95% CI = 0.30–0.66, P < 0.0001) and increased PFS rates (OR = 0.40, 95% CI = 0.35–0.46, P < 0.0001, Figure 3). The analysis of PFS rates from 2 to 12 months in the forest plot illustrated the continuous effects of PARPis. Statistical heterogeneity was not obvious (Cochran's Q test, P = 0.16, $I^2 = 30\%$).

We conducted subgroup analyses based on the BRCA status, receptor status (hormone receptor-positive or triple-negative), and receipt of platinum treatment. Among all subgroups analyzed, the PARPi arm was significantly preferred in terms of PFS in four subgroups: TNBC (OR = 0.39, 95%) CI = 0.24 - 0.63, P = 0.0001), mutant BRCA1 (OR = 0.36, 95% CI = 0.23-0.57, P < 0.0001), mutant BRCA2 (OR = 0.54, 95% CI = 0.34-0.85, P = 0.007), and no prior platinum treatment (OR = 0.48, 95%CI = 0.34 - 0.69, P < 0.0001, Figure 4). Patients with hormone receptor-positive cancer and those who previously received platinum treatment did not significantly benefit from PARPis.

Assessment of serious adverse events

Adverse events related to the treatment were recorded in all of the included clinical trials. The main toxic effects were reflected in the blood and digestive systems. First, we compared the risks of grade 3 and 4 side

	chemotherapy+ PAR	Pi	chemothe	erapy		Odds Ratio	Odds Ratio
Study or Subgroup	Events	Total	Events	Total	Weight	M-H, Fixed, 95% Cl	M-H, Fixed, 95% Cl
Clinical Response							
Joyce O' Shaughnessy 2014	173	261	180	258	21.4%	0.85 [0.59, 1.23]	
Shivaani Kummar 2016	27	38	16	18	2.2%	0.31 [0.06, 1.56]	
Subtotal (95% CI)		299		276	23.6%	0.80 [0.56, 1.15]	•
Total events	200		196				
Heterogeneity: Chi ² = 1.44, df =	1 (P = 0.23); P = 31%						
Test for overall effect: Z = 1.22 (F	P = 0.22)						
PFS (2 months)							
Joyce O' Shaughnessy 2014	74	261	87	258	22.0%	0.78 [0.54, 1.13]	
Shivaani Kummar 2016	6	24	9	18	2.7%	0.33 [0.09, 1.23]	
Subtotal (95% CI)		285		276	24.7%	0.73 [0.51, 1.04]	•
Total events	80		96				
Heterogeneity: Chi ² = 1.49, df =	1 (P = 0.22); I ² = 33%						
Test for overall effect: Z = 1.73 (F	P = 0.08)						
PFS (4 months)							
Joyce O' Shaughnessy 2014	123	261	142	258	26.5%	0.73 [0.52, 1.03]	
Shivaani Kummar 2016	12	24	11	18	2.2%	0.64 [0.18, 2.20]	
Subtotal (95% CI)		285		276	28.7%	0.72 [0.52, 1.01]	•
Total events	135		153				
Heterogeneity: Chi ² = 0.04, df =	1 (P = 0.84); I ² = 0%						
Test for overall effect: Z = 1.93 (F	P = 0.05)						
PFS (6 months)							
Joyce O' Shaughnessy 2014	178	261	195	258	21.9%	0.69 [0.47, 1.02]	
Shivaani Kummar 2016	18	24	12	18	1.2%	1.50 [0.39, 5.77]	
Subtotal (95% CI)		285		276	23.1%	0.73 [0.51, 1.06]	•
Total events	196		207				
Heterogeneity: Chi ² = 1.17, df =	1 (P = 0.28); I ² = 14%						
Test for overall effect: Z = 1.63 (F	P = 0.10)						
Total (95% CI)		1154		1104	100.0%	0.75 [0.62, 0.89]	•
Total events	611		652				
Heterogeneity: Chi ² = 4.40, df =	7 (P = 0.73); I ² = 0%						
Test for overall effect: Z = 3.26 (F	P = 0.001)						Favours [Cham+PARPi] Favours [Cham]
Test for subaroup differences: C	chi² = 0.21. df = 3 (P =	0.98). F	² = 0%				arours (chemistrating Pavours (chemi

Figure 2. Forest plot of the pooled relative risk of clinical efficacy from the included studies reporting the clinical outcome associated with the combination of PARPis and chemotherapy. Horizontal lines represent 95% Cls.

M-H, Mantel-Haenszel; df, degrees of freedom; chem., chemotherapy; PARPi, poly ADP-ribose polymerase inhibitor; CI, confidence interval.

effects between PARPi therapy and chemotherapy (Figure 5). Robson et al.¹⁶ reported fewer adverse effects in the single-agent PARPi arm. The current study revealed no differences in terms of serious side effects (\geq Grade 3) between the PARPi and standard chemotherapy arms (OR = 0.76, 95% CI = 0.54–1.08, P = 0.12).

The other two studies reported data for the specific adverse reactions of PARPis in combination with chemotherapy.^{19,20} The most common side effects were neutropenia, anemia, thrombocytopenia, leucopenia, and fatigue or asthenia. From the forest graph (Figure 6), the incidence of all of the aforementioned adverse events were similar between the PARPi monotherapy and combination treatment groups (OR = 1.09, 95% CI = 0.88–1.35).

Publication bias

All meta-analyses in our study were divided into two parts: PARPi alone or combined with chemotherapy versus chemotherapy alone. Therefore, the publication bias assessments were divided into two parts. The funnel plot (Figure 7) presented no evidence of remarkable asymmetry in the monotherapy and combination arms (P=0.5 and P=0.98, respectively).

Discussion

The current meta-analysis assessed the efficiency, safety, and benefits of PARPis. Relative to standard chemotherapy, PARPi monotherapy appeared to be effective and safe for patients with advanced breast cancer. Subgroup analysis illustrated

	PAR	Pi	chemot	therapy		Odds Ratio	Odds Ratio
Study or Subgroup	Events	Total	Events	Total	Weight	M-H, Fixed, 95% Cl	M-H, Fixed, 95% Cl
Clinical response							
Jennifer K. 2018	36	219	28	114	4.6%	0.60 [0.35, 1.05]	
Mark Robson 2017	67	167	45	66	5.8%	0.31 [0.17, 0.57]	
Subtotal (95% CI)		386		180	10.4%	0.44 [0.30, 0.66]	•
Total events	103		73				
Heterogeneity: Chi ² = 2.	48, df = 1 (P	= 0.12);1	² = 60%				
Test for overall effect: Z	= 3.97 (P < 0	0.0001)					
PFS (2 months)							
Jennifer K. 2018	29	287	38	144	6.8%	0.31 (0.18, 0.53)	
Mark Robson 2017	28	205	34	97	5.9%	0 29 [0.16, 0.52]	
Subtotal (95% CI)		492		241	12.7%	PARPi [0.21, 0.45]	◆
Total events	57		72				
Heterogeneity: Chi ² = 0.	03, df = 1 (P	= 0.87); 1	² = 0%				
Test for overall effect: Z	= 5.96 (P < 0	0.00001)					
DES (4 months)							
Ionnifer 1/ 2018	63	297	57	144	0.004	0.43 (0.38, 0.66)	
Mark Roheon 2017	51	207	53	07	9 1 %	0.43 [0.28, 0.00]	
Subtotal (95% CI)	51	492	33	241	16.9%	0.36 [0.26, 0.49]	•
Total events	114		110				1
Heterogeneity: Chi ² = 1.	70. df = 1 (P	= 0.19); [= 41%				
Test for overall effect: Z	= 6.14 (P < 0	0.00001)					
	•						
PFS (6 months)							
Jennifer K. 2018	109	287	82	144	10.1%	0.46 [0.31, 0.70]	
Mark Robson 2017	98	205	72	97	7.6%	0.32 [0.19, 0.54]	
Subtotal (95% CI)		492		241	17.7%	0.40 [0.29, 0.55]	•
Total events	207		154				
Heterogeneity: Chi* = 1.	21, df = 1 (P	= 0.27);1	*=17%				
lest for overall effect. Z	= 5.57 (P < l	J.UUUU1)					
PFS (8 months)							
Jennifer K. 2018	152	287	89	144	8.3%	0.70 [0.46, 1.05]	
Mark Robson 2017	111	205	76	97	7.1%	0.33 [0.19, 0.57]	
Subtotal (95% CI)		492		241	15.4%	0.53 [0.38, 0.73]	•
Total events	263		165				
Heterogeneity: Chi ² = 4.	64, df = 1 (P	= 0.03); 1	² = 78%				
Test for overall effect: Z	= 3.87 (P = 0	0.0001)					
PFS (10 months)							
Jennifer K. 2018	167	287	106	144	8.8%	0.50 [0.32, 0.77]	
Mark Robson 2017	136	205	86	97	5.9%	0.25 [0.13, 0.50]	
Subtotal (95% CI)		492		241	14.7%	0.40 [0.28, 0.58]	•
Total events	303		192				
Heterogeneity: Chi ² = 2.	69, df = 1 (P	= 0.10); 1	² = 63%				
Test for overall effect Z	= 4.90 (P < (0.00001)					
PFS (12 months)							
Jennifer K. 2018	178	287	115	144	8.7%	0.41 [0.26, 0.66]	
Mark Robson 2017	165	205	89	97	3.5%	0.37 [0.17, 0.83]	
Subtotal (95% CI)		492		241	12.2%	0.40 [0.27, 0.60]	◆
Total events	343		204				
Heterogeneity: Chi ² = 0.	05, df = 1 (P	= 0.82); 1	² = 0%				
Test for overall effect: Z	= 4.42 (P < (0.0001)					
Total (95% CI)		3338		1626	100.0%	0.40 (0.35, 0.46)	•
Total events	1390	3330	970	1020	.00.070	0.40 [0.00, 0.40]	
Heterogeneity: Chi2 = 19	3.34 df = 13	(P = 0.15)): P= 29%				
Test for overall effect: Z	= 13.08 (P <	0.00001)				0.01 0.1 1 10 100
Test for subaroup differ	ences: Chi ²	= 5.33. df	f= 6 (P = 0.	50). I ² =	0%		Favours [PARPi] Favours [enem]

Figure 3. Forest plot of pooled relative risk of clinical efficacy from the included studies reporting clinical outcomes associated with PARPi monotherapy compared with chemotherapy. Horizontal lines represent 95% Cls.

M-H, Mantel–Haenszel; df, degrees of freedom; chem., chemotherapy; PARPi, poly ADP-ribose polymerase inhibitor; CI, confidence interval.

that patients with TNBC, *BRCA1* mutation, and no prior history of platinum therapy more strongly benefited from PARPi treatment. In addition, the combination of PARPis and chemotherapy significantly improved the survival of patients with TNBC.

Chemotherapy remains the primary treatment for patients with metastatic breast cancer at present. In recent years, several

	PARE	Pi	chen	aothera	ipy	Odds Ratio	Odds Ratio
Study or Subgroup	Events	Total	Events	Total	Weight	M-H, Fixed, 95% Cl	M-H, Fixed, 95% Cl
All patients							
Jennifer K. 2018	111	287	78	144	60.3%	0.53 [0.36, 0.80]	
Mark Robson 2017	92	205	56	97	39.7%	0.60 (0.37, 0.97)	
Subtotal (95% CI)		492		241	100.0%	0.56 [0.41, 0.76]	•
Total events	203		134				
Heterogeneity: Chi ² =	0.12 df = 1	(P = 0)	73): 12 = 0	196			
Test for overall effect:	7 = 3 67 (P	= 0.00	02)				
restion overall ellect.	2 - 5.01 (1	- 0.00	02/				
Hormone-receptor po	sitive						
Jennifer K. 2018	50	157	40	84	74.6%	0.51 [0.30, 0.89]	
Mark Robson 2017	75	100	36	49	25.4%	1.08 [0.50, 2.36]	
Subtotal (95% CI)		257		133	100.0%	0.66 [0.42, 1.03]	•
Total events	125		76				C79944
Heterogeneity: Chi ² =	2.36 df = 1	(P = 0)	12): 12 = 5	58%			
Test for overall effect:	Z = 1.83 (P	= 0.07)				
Triple negative							
Jennifer K. 2018	47	130	36	60	59.3%	0.38 [0.20, 0.71]	
Mark Robson 2017	25	102	21	48	40.7%	0.42 [0.20, 0.86]	
Subtotal (95% CI)		232		108	100.0%	0.39 [0.24, 0.63]	◆
Total events	72		57				1.51
Heterogeneity: Chi ² =	0.04 df = 1	(P = 0)	84): 12 = 0	196			
Test for overall effect:	7 = 3.84 (P	= 0.00	01)				
restion overall ellect.	2 - 5.04 (1	- 0.00	01)				
Mutated BRCA1							
Jennifer K. 2018	46	133	37	63	55.2%	0.37 10.20, 0.691	
Mark Robson 2017	33	114	27	50	44.8%	0.35 (0.17 0.69)	
Subtotal (95% CI)		247		113	100.0%	0.36 [0.23, 0.57]	•
Total events	79		64				
Heterogeneity Chi ² =	0.02 df=1	(P = 0)	88) 12=1	196			
Test for overall effect:	7 = 4 36 (P	< 0.00	01)				
restion overall ellect.	2 - 4.00 (0.00	0.7				
Mutated BRCA2							
Jennifer K. 2018	46	154	38	81	68.1%	0.48 [0.28, 0.84]	
Mark Robson 2017	50	84	31	45	31.9%	0.66 (0.31, 1.43)	
Subtotal (95% CI)		238	100	126	100.0%	0.54 [0.34, 0.85]	◆
Total events	96		69				Londer M
Heterogeneity: Chi ² =	0.44 df = 1	(P = 0)	51): I ² = (1%			
Test for overall effect:	7 = 2.68 (P	= 0.00	7)				
		0.00	· ·				
Previous platinum tre	eatment(YE	S)					
Jennifer K. 2018	33	46	23	30	44.3%	0.77 [0.27, 2.23]	
Mark Robson 2017	35	60	17	26	55.7%	0.74 [0.28, 1.93]	
Subtotal (95% CI)		106		56	100.0%	0.76 [0.37, 1.54]	-
Total events	68		40				
Heterogeneity: Chi ² =	0.00. df = 1	(P = 0)	95); ² = (1%			
Test for overall effect:	Z = 0.77 (P	= 0.44)				
Previous platinum tre	eatment(NO)					
Jennifer K. 2018	88	241	59	114	59.2%	0.54 [0.34, 0.84]	
Mark Robson 2017	66	145	48	71	40.8%	0.40 [0.22, 0.73]	
Subtotal (95% CI)		386		185	100.0%	0.48 [0.34, 0.69]	•
Total events	154		107				
Heterogeneity: Chi2=	0.59, df = 1	(P = 0)	44); 12 = 0	9%			
Test for overall effect	Z = 4.00 (P	< 0.00	01)				
		0.00	/				
							U.U1 U.1 1 10 100
Test for subgroup diff	erences: Ch	$ni^2 = 6.2$	25 df = 6	(P = 0)	40) P = 4	0%	Pavours [PARPi] Pavours [cnem]

Figure 4. Pooled subgroup analysis of the relative risk of survival from the included studies reporting the disease-free survival of specific patients (e.g., triple-negative breast cancer, *BRCA1/2* mutation, receipt of previous platinum therapy). Horizontal lines represent 95% Cls.

M-H, Mantel–Haenszel; df, degrees of freedom; chem., chemotherapy; PARPi, poly ADP-ribose polymerase inhibitor; Cl, confidence interval.

studies demonstrated the efficacy of platinum drugs against TNBC. Platinum drugs cause DNA cross-linking, hinder DNA synthesis, and inhibit tumor growth.²² However, resistance to chemotherapy, which is the main cause of treatment failure in patients advanced breast cancer and poor prognosis, is extremely common. Therefore, preclinical and clinical trials have been devoted to identifying new therapeutic targets.



Figure 5. Pooled analysis of side effects comparing PARPis with chemotherapy. Horizontal lines represent 95% Cls.

M-H, Mantel–Haenszel; df, degrees of freedom; chem., chemotherapy; PARPi, poly ADP-ribose polymerase inhibitor; CI, confidence interval.

	chemotherapy+PA	RPi	chemo	therapy		Odds Ratio	Odds Ratio
Study or Subgroup	Events	Total	Events	Total	Weight	M-H, Fixed, 95% Cl	M-H, Fixed, 95% Cl
Neutropenia							
Joyce O' Shaughnessy 2014	207	255	201	244	23.7%	0.92 [0.59, 1.45]	
Shivaani Kummar 2016	1	37	0	18	0.4%	1.52 [0.06, 39.18]	
Subtotal (95% CI)		292		262	24.1%	0.93 [0.59, 1.46]	◆
Total events	208		201				
Heterogeneity: Chi ² = 0.09, df = 1	(P = 0.77); I ² = 0%						
Test for overall effect: Z = 0.31 (P	= 0.76)						
Anemia							
Joyce O' Shaughnessy 2014	48	255	53	244	26.9%	0.84 [0.54, 1.29]	
Shivaani Kummar 2016	4	37	1	18	0.7%	2.06 [0.21, 19.91]	
Subtotal (95% CI)		292		262	27.6%	0.87 [0.57, 1.33]	•
lotal events	52		54				
Heterogeneity: Chi ² = 0.59, df = 1	(P = 0.44); $P = 0%$						
Test for overall effect: Z = 0.65 (P	= 0.52)						
Thrombocytopenia							
Joyce O' Shaughnessy 2014	74	255	58	244	25.8%	1.31 [0.88, 1.96]	+
Shivaani Kummar 2016	1	37	0	18	0.4%	1.52 (0.06, 39,18)	
Subtotal (95% CI)		292		262	26.1%	1.31 [0.88, 1.95]	•
Fotal events	75		58				184 1
Heterogeneity Chi ² = 0.01, df = 1	(P = 0.93); $P = 0.96$						
Test for overall effect: Z = 1.35 (P	= 0.18)						
Leukopenia							
Joyce O' Shaughnessy 2014	41	255	37	244	19.4%	1.07 (0.66, 1.74)	
Shivaani Kummar 2016	1	37	0	18	0.4%	1.52 (0.06, 39,18)	
Subtotal (95% CI)		292		262	19.8%	1.08 [0.67, 1.74]	◆
Fotal events	42		37				
Heterogeneity Chi ² = 0.04 df = 1	(P = 0.83); IF = 0%						
Test for overall effect: Z = 0.32 (P	= 0.75)						
Fatigue or asthenia							
loyce O' Shaughnessy 2014	11	57	4	59	1.9%	3.29 [0.98, 11.02]	
Shivaani Kummar 2016	1	37	0	18	0.4%	1.52 (0.06, 39.18)	
Subtotal (95% CI)		94		77	2.3%	2.99 [0.96, 9.31]	-
Total events	12		4				
Heterogeneity Chi ² = 0.19 df = 1	(P = 0.66); P = 0%		÷.				
Test for overall effect: Z = 1.89 (P	= 0.06)						
Total (95% CI)		1262		1125	100.0%	1.09 [0.88, 1.35]	+
Total events	389		354				
Heterogeneity: Chi ² = 6.43. df = 9	$(P = 0.70); I^2 = 0\%$						
Test for overall effect: Z = 0.81 (P	= 0.42)						0.01 0.1 1 10 10
Test for subgroup differences: Ch	ni2 = 5.44. df = 4 (P =	0.24)	² = 26.5%				Favours [chem+PARPi] Favours [chem]

Figure 6. Pooled analysis of specific side effects of the combination of PARPis with chemotherapy. Horizontal lines represent 95% Cls.

M-H, Mantel–Haenszel; df, degrees of freedom; chem., chemotherapy; PARPi, poly ADP-ribose polymerase inhibitor; CI, confidence interval.

Studies have revealed that 5% to 10% of patients with breast cancer have a clear genetic mutation, called hereditary breast cancer, 23,24 in which *BRCA1/2* gene

mutation accounts for 15% of such lesions.^{25,26} Most *BRCA*-associated breast cancers are triple-negative.²⁷ Interestingly, *BRCA1* is the most studied gene associated



Figure 7. Funnel plot for publication bias. (a) Funnel plot of the relationship between PARPi combination therapy and clinical efficacy. (b) Funnel plot of the relationship between PARPi monotherapy and clinical efficacy.

PARPi, poly ADP-ribose polymerase inhibitor.

with platinum resistance, and *BRCA1*-deficient tumor cells are more sensitive to cisplatin and other platinum drugs.²⁸ A previous study also revealed that breast cancer cells with *BRCA1/2* mutations are more sensitive to DNA cross-linking agents, such as cisplatin, carboplatin, and mitomycin.²⁹

PARPis induce DNA single-strand breaks by blocking the repair of single-DNA breakpoints, stranded whereas BRCA mutants cannot initiate HR to repair DNA duplexes. PARP and BRCA are genes with synergistic lethal effects against tumor cells. Therefore BRCA mutants are sensitive to PARPis, leading to satisfactory clinical effects.^{16,30} The Clinical 2017 American Society of Oncology (ASCO) meeting reported the results of Phase III clinical trials of olaparib for patients with metastatic breast cancer. In January 2018, the FDA approved olaparib for the treatment of HER2-negative metastatic breast cancer carrying the BRCA gene mutation. In 2020, ASCO, the American Society of Radiation Oncology, and the Society of Surgical Oncology convened expert teams to formulate recommendations for the treatment of patients with breast cancer and susceptible germline mutations based on systematic reviews of the literature. The teams proposed that for HER2-negative breast cancer with BRCA1/ 2 mutations, olaparib or talazoparib should be used instead of chemotherapy in the first three lines of treatment. For BRCA1/2 mutation carriers with metastatic HER2negative breast cancer, there are no data directly comparing the efficacy of PARPis and platinum-based chemotherapy.³¹ The current meta-analysis integrated data for olaparib and talazoparib, and the clinical effects were consistent with the reported phase III studies.^{16,17} Furthermore, patients with BRCA1/2 mutations and those who have not received platinum therapy can significantly benefit from PARPis. The results from the subgroup analysis were not completely consistent with the previously reported results, which revealed that the clinical outcomes of BRCA1 mutation carriers were worse than those of BRCA2 carriers.³² mutation The correlation between BRCA1/2 gene mutation and the prognosis of breast cancer is unclear.^{33,34} We speculated that mutant BRCA1 and BRCA2 played different roles in the response to PARPis. It is possible that the previously reported worse survival of BRCA2-mutant cancer was related to the

different response to PARPis analyzed in our study. However, additional prospective studies are needed for confirmation.

Although PARPis are extremely effective BRCA-mutant and platinumagainst resistant ovarian cancer, we found that patients with breast cancer who had previously received platinum-based treatment did not benefit from PARPis compared with the effects of chemotherapy. The specific mechanism is unclear at present. PARPis are currently approved for HER2-negative metastatic breast cancer carrying BRCA germline mutations. According to the latest guideline. platinum-based chemotherapy is recommended preferentially.³⁵ However, our study suggested that patients who did not receive platinum treatment could benefit from PARPis, which means that PARPis would have greater utility in the adjuvant treatment stage. Clinical trials using PARPis in the adjuvant and neoadjuvant phases are underway. **BrighTNess** (NCT02032277) was a phase III trial assessing the combination of veliparib and chemotherapy in the neoadjuvant setting;³⁶ BRE09-146 (NCT01074970) was a phase II trial that evaluated rucaparib combined with cisplatin in the adjuvant phase.³⁷ Results from BrighTNess illustrated that a grossly subtherapeutic dose of PARPis in combination with standard doses of chemotherapy did not significantly improve clinical outcomes. We must await additional clinical trials and other research.

Although the original purpose of PARPis was to increase the sensitivity of tumor cells to chemotherapy by causing DNA damage, the clinical outcomes of combination studies of chemotherapy and PARPis were heterogeneous. The main reason is that the side effects of chemotherapy on normal healthy cells tend to limit the drug dosage, and combined usage with PARPis will increase side effects in healthy cells. Preclinical studies illustrated that high doses of PARPis combined with relatively low doses of chemotherapy could inhibit the proliferation of tumor cells.^{38,39} The safety and efficacy of this combination therapy are being tested in clinical trials. A phase III clinical trial reported the results of iniparib plus chemotherapy compared with chemotherapy alone in metastatic TNBC. Unfortunately, the trial did not meet the expected primary endpoints of PFS and OS.²⁰ In our meta-analyses, the combination of PARPis with chemotherapy provided survival benefits for patients.

Clinical scientists are also working to further improve the clinical remission rate of TNBC and overcome the occurrence of drug resistance. PARPi combination treatments are worthy of further study. Clinical trials of PARPis combined with immune checkpoint inhibitors are also undergoing. The rationale for these combinations is that tumors with HR defects usually carry more genetic mutations, which may lead to the production of more new antigens and induction of stronger anti-tumor immune responses.⁴⁰ Several studies, including the TOPACIO (NCT02657889), MEDIOLA (NCT02734004), and NCT02849496 trials, combined PARPis with immune checkpoint inhibitors.^{41–43} In the setting of ovarian cancer, the results of one Phase I study (TOPACIO/Keynote-162)⁴⁴ demonstrated that niraparib combined with pembrolizumab was feasible and safe, with no expected toxicity observed. Strategies using immune checkpoint inhibitors are generally not hindered by additive toxicities in breast cancer, but the utility of combining PARPis with immunotherapy has not been particularly effective to date.

Conclusion

PARPi monotherapy could obviously improve the clinical outcomes of patients with advanced breast cancer, especially those with TNBC, *BRCA1/2* mutations, and no prior history of platinum therapy. The combination of PARPis with chemotherapy represents a novel option for such patients. In patients with advanced TNBC who responded to previous platinum therapy, PARPis can be considered. Future PARPi studies should cover the following points: the selection of the most suitable patients for PARPi therapy, the development of drug resistance, and the optimum combination therapy.

Author contributions

ZC and KC contributed to study conception and design. ZC, XW, and YZ reviewed the literature and designed the article structure. ZC, KC, and YZ contributed to the acquisition and analysis of data. ZC and XW participated in data interpretation. ZC and KC were major contributors to writing the manuscript. XW, XL, and YZ revised and edited the manuscript critically for important intellectual content. ZC, KC, YZ, and XL gave final approval of the version to be published.

Declaration of conflicting interest

The authors declare that there is no conflict of interest.

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