

Review Article



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Conflict of Interest

The authors declare that they have no competing interests.

Author Contributions

Conceptualization: Xu X; Data curation: Liu S, Sun X, Xu X, Lin F; Formal analysis: Liu S, Sun X, Lin F; Funding acquisition: Sun X; Investigation: Liu S, Xu X, Lin F; Methodology: Liu S, Sun X; Project administration: Liu S;

Comparison of Endocrine Therapies in Hormone Receptor-Positive and Human Epidermal Growth Factor Receptor 2-Negative Locally Advanced or Metastatic Breast Cancer: A Network Meta-Analysis

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ABSTRACT

We aimed to explore what kind of endocrine treatments are optimal for hormone receptor-positive and human epidermal growth factor receptor 2-negative locally advanced or metastatic breast cancer in some specific clinical situations. We searched randomized controlled trials in Embase, Medline, the Cochrane library, and PubMed from inception to April 1, 2020 and performed a network meta-analysis based on a Bayesian fixed-effects model. Progression-free survival (PFS) with hazard ratios and corresponding 95% confidence interval was defined as the primary endpoint, while overall survival (OS), objective response rate (ORR), clinical benefit rate and serious adverse events were used as secondary endpoints. A total of 35 studies involving 12,285 patients and 24 treatment options were included. In general, most co-treatment options prolonged PFS compared to single-agent therapy, of which aromatase inhibitor (AI) plus everolimus and fulvestrant plus palbociclib were probably the most effective agents, and the latter had the best safety record. However, despite the superior efficacy of fulvestrant plus capecitabine for PFS and OS, palpable toxic effects have been demonstrated for this treatment, so its application must be scrupulously considered. The results of subgroup analysis indicated that fulvestrant combined with palbociclib improved prognosis for phosphatidylinositol 3-kinase (PI3K)-mutated patients, PI3K-unmutated patients, patients with endocrine therapy resistance, and visceral metastatic patients, while no obvious improvement was detected in OS. Moreover, the efficacy of fulvestrant plus cyclin-dependent kinase 4/6 (CDK4/6) inhibitors was slightly better than that of AI plus CDK4/6 inhibitors, while AI plus everolimus was more efficacious than fulvestrant combined with everolimus in terms of PFS, OS, and ORR. In conclusion, our results provide moderate evidence that fulvestrant plus palbociclib and AI plus everolimus were the most effective treatments, while the efficacy and safety of fulvestrant plus palbociclib was obviously superior in some specific clinical situations.

Keywords: Breast; Neoplasms; Prognosis; Safety

Resources: Liu S; Software: Liu S, Sun X, Xu X; Supervision: Xu X, Lin F; Validation: Liu S, Xu X, Lin F; Visualization: Liu S, Xu X; Writing - original draft: Liu S; Writing - review & editing: Liu S, Sun X, Xu X, Lin F.

INTRODUCTION

Approximately 60%–70% of breast cancer patients present as hormone receptor-positive and human epidermal growth factor receptor 2 (HER2)-negative, with their treatment being based on endocrine therapy (ET) against the estrogen axis [1,2]. Although hormone receptor-positive/HER2-negative patients have a superior prognosis compared with triple-negative and HER2-overexpression breast cancer, the vast majority of patients will eventually progress to metastatic disease with or without ET resistance [3,4].

The activation of the phosphatidylinositol 3-kinase (PI3K)—protein kinase B (PKB/AKT)—mammalian target of rapamycin (mTOR) axis has emerged as a promotor of both tumor cell growth and ET resistance [5], and this pathway is considered to be a reasonable therapeutic target to inhibit tumor growth and revert ET resistance. Furthermore, another therapeutic target, serine/threonine protein kinase mTOR, plays an important role in many solid tumors, including breast cancer, and can regulate cell proliferation, differentiation and apoptosis [6]. In patients who had previously received ET, SOLAR-1 demonstrated that cotreatment with alpelisib, a PI3K α -specific inhibitor, improved progression-free survival (PFS) among patients with PIK3CA mutations, which are detected in 40% of hormone receptor-positive/HER2-negative breast cancer [7]. The mTOR inhibitor everolimus also shows an anti-tumor effect and can reverse endocrine sensitivity in combination with ET for hormone receptor-positive/HER2-negative advanced breast cancer patients [8].

Dysregulation of the cyclin D–cyclin-dependent kinase (CDK)—retinoblastoma (Rb)—E2F pathway results in cell cycle progression through regulation of the G1-S checkpoint, and is detected in many cancers, especially in hormone receptor-positive breast carcinoma [9,10]. Components of the CDK4/6-Rb-E2F axis are abnormally mutated in 50%–70% of breast cancers, which contributes ET resistance [11]. Currently, in addition to traditional endocrine monotherapy including selective estrogen receptor modulators and aromatase inhibitors (AIs), many phase III randomized controlled trials (RCTs) such as MONALEESA-2 and MONARCH-3, have also demonstrated that cotreatment with cyclin-dependent kinase 4/6 (CDK4/6) inhibitors can significantly improve the prognosis in first-line treatments with manageable toxic effects [12,13]. Additionally, PALOMA-3 and MONARCH-2 showed that fulvestrant combined with CDK4/6 inhibitors was associated with longer PFS in ET-resistant patients [14,15]. Some articles indirectly compared the efficacy and toxicity among three different CDK4/6 inhibitors in the first-line treatment, and they appeared to have a similar benefit without significant difference [16].

In addition to inhibitors described above, there are still many targeted drugs and chemotherapy options that have the potential to improve the prognosis of hormone receptor-positive breast cancer. Notably, it remains unclear which kind of medicine works better under certain clinical conditions and which options are safer. Therefore, the purpose of our systematic review and network meta-analysis was to indirectly compare efficacy and safety between different ET strategies, given the lack of direct comparison for hormone receptor-positive/HER2-negative locally advanced or metastatic breast cancer in different situations.

METHODS

Search strategy

For this network meta-analysis, relevant phase 2 and 3 RCTs published before April 1, 2020 were searched systematically and logically without limitation of language in electronic databases in Embase, Medline, the Cochrane Library, and PubMed. The literature search strategy employed was as follows: breast AND (cancer OR neoplasm OR carcinoma) AND endocrine therapy AND HER2 negative. Meanwhile, related literature and conference abstracts were searched for references to ensure the comprehensiveness of results.

Selection criteria

Two researchers independently examined the literature, screening the title and abstract, and disagreements on eligible studies were resolved by discussion and negotiation or by consulting a third investigator. The inclusion criteria were as follows: 1) the patients had a pathological diagnosis of hormone receptor-positive/HER2-negative locally advanced or metastatic breast cancer; 2) a comparison was reported between different ETs; 3) the literature provided complete data on PFS, overall survival (OS), clinical benefit rate (CBR), objective response rate (ORR), or serious adverse events; 4) the study type was phase 2 or 3 non-single arm RCT. Only the reports with the longest follow-up period or the most comprehensive data were selected in order to omit overlapping populations. Simultaneously, we excluded trials in which the population was hormone receptor-negative or HER2-positive, or had early-stage breast cancer, and unsuitable study types were also deleted, such as reviews, letters, and retrospective studies.

Data extraction and quality assessment

Two researchers independently extracted and tabulated detailed data from eligible studies, including study name, intervention, sample size, patient clinical characteristics, and outcomes. In this network meta-analysis, the primary endpoint was PFS with corresponding hazard ratios (HRs), which was assessed based on the time from randomization to tumor progression or death based on RECIST (version 1.1). OS (time from randomization to death due to any cause), CBR (patients with complete response, partial response and stable disease ≥ 6 months), ORR (patients with complete response and partial response), and serious adverse events were defined as secondary endpoints. The risk of bias in the included literature was also assessed by two researchers independently utilizing the Cochrane collaboration's tool, which consisted of seven elements: random sequence generation, allocation concealment, blinding of participants and personnel, blinding of outcome assessment, incomplete outcome data, selective outcome reporting, and other sources of bias [17].

Statistical analysis

In this network-meta analysis, HR with corresponding 95% confidence interval (CI) was applied to assess PFS and OS, while relative risk and corresponding 95% CI were used to assess dichotomous data. The entire process was divided into 2 parts: direct comparison and indirect comparison. For direct comparison, the I-squared test was performed to judge heterogeneity originating from data combination. A fixed-effects model was used when I-squared $< 50\%$, indicating slight heterogeneity, otherwise a random-effects model was used. Indirect comparison was performed utilizing a Bayesian fixed-effects model to pool extracted data and evaluate various treatment options. The surface under cumulative ranking curve (SUCRA) was derived to evaluate the ranking possibility of each treatment strategy, and the closer the value of SUCRA was to 1, the better the efficacy and safety of intervention [18]. Moreover, we grouped patients into multiple

subtypes based on PIK3CA-mutation status, ET resistance, and metastatic setting to assess the superior treatment in different situations. For all statistical results, *p*-values of 0.05 or lower were considered to indicate statistical significance. Statistical analysis was conducted using STATA 15.0 (StataCorp LLC, College Station, TX, USA), WinBUGS 14.0 (Imperial College School of Medicine, London, UK), GeMTC 0.14.3 (University of Groningen, Groningen, Netherlands).

Ethics statement

This article does not contain any studies with human participants or animals performed by any of the authors.

RESULTS

Search results

Without restriction of language, a total of 1,014 studies were discovered through the search strategy, including 122 studies in PubMed, 74 in Medline, 576 in the Cochrane Library, 206 in Embase and 37 additional records. After screening the titles and abstracts, 124 remaining records were considered to be eligible for a full text review. Ultimately, 35 articles were selected which met all the inclusion criteria, giving a total of 12,285 patients involved in this network meta-analysis, as shown in [7,8,12-15,19-47]. The search process based on the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) technique is shown in in **Figure 1** [48].

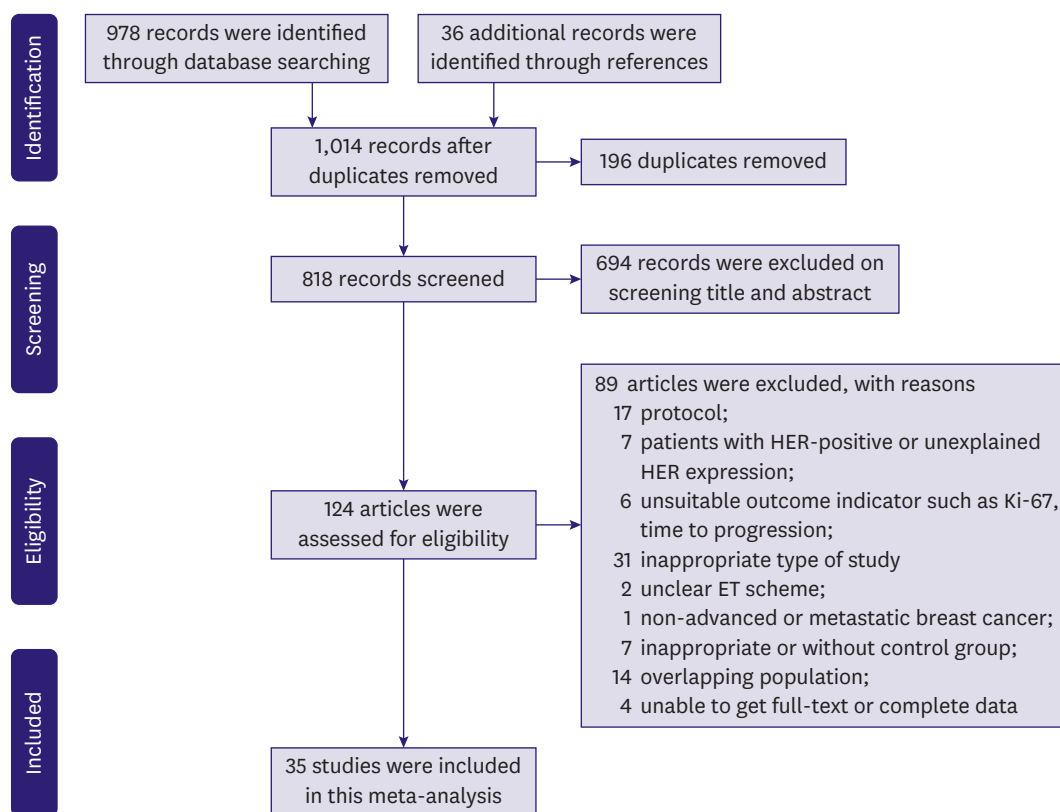


Figure 1. Flow diagram summarizing the selection process.

HER = human epidermal growth factor receptor; ET = endocrine therapy.

Characteristics and quality of study

The clinical characteristics of enrolled studies were similar in that all participations had histologically confirmed hormone receptor-positive/HER2-negative locally advanced or metastatic breast cancer based on the results of biopsy (primary tumor or metastatic disease) and about 85% of patients were post-menopausal. The median age of included patients ranged from 43 to 67 years. The interventions reported consisted of three CDK4/6 inhibitors in 10 reports, PI3K inhibitors in 5, mTOR inhibitor in 4, and anti-vascular endothelial growth factor (VEGF) drug in 2, and the control groups were generally treated with monotherapy, such as AI or fulvestrant. The dosage of fulvestrant in most studies was 500 mg per cycle (28 days), while the dosage in SoFEA and SWOG0226 was half as much. A total of 24 treatment options were involved in this network meta-analysis. In terms of outcomes, all studies, except for that of Howell et al. [19] and KCSG BR10-04, which only provided data on CBR or ORR, provided PFS data in detail. BOLERO-6 was a three-arm study, and all the others are two-arm studies, and all the studies which qualified for inclusion were phase 2 or 3 RCTs. More detailed features are listed in **Table 1**. A network plot displaying direct comparisons between different treatments is shown in **Figure 2**.

Table 1. Characteristics of the outcomes of the studies included in this network meta-analysis

Study	Treatment	No. of patients	Characteristic	PFS (HR and 95% CI)	OS (HR and 95% CI)	ORR	CBR
PALOMA-1 [36]	AI+PALBO	84	First-line therapy for postmenopausal women with advanced HR-positive/HER2-negative breast cancer	0.49 (0.32–0.75)	0.81 (0.49–1.35)	36/84	68/84
	AI	81				27/81	47/81
PALOMA-2 [37]	AI+PALBO	444	First-line therapy for postmenopausal women with advanced HR-positive/HER2-negative breast cancer	0.58 (0.46–0.72)	NA	187/444	377/444
	AI	222				77/222	156/222
PALOMA-3 [14,42,46]	FUL+PALBO	347	ET resistant women with HR-positive/HER2-negative advanced or metastatic breast cancer	0.46 (0.36–0.59)	0.81 (0.64–1.03)	66/347	231/347
	FUL	174				15/174	69/174
MONALEESA-2 [13,43]	AI+RIBO	334	First-line therapy postmenopausal women with HR-positive/HER2-negative metastatic breast cancer	0.56 (0.43–0.72)	0.75 (0.52–1.08)	135/334	205/334
	AI	334				97/334	176/334
MONALEESA-3 [24]	FUL+RIBO	484	Postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.59 (0.48–0.73)	0.72 (0.57–0.92)	157/484	340/484
	FUL	242				52/242	152/242
MONALEESA-7 [41,44]	AI+RIBO	248	Premenopausal women with HR-positive/HER2-negative advanced breast cancer	0.57 (0.44–0.74)	0.70 (0.50–0.98)	NA	NA
	AI	247					
MONARCH-2 [15,45]	FUL+ABEMA	446	ET resistant women with HR-positive/HER2-negative advanced breast cancer	0.55 (0.45–0.68)	0.76 (0.61–0.95)	157/446	322/446
	FUL	223				36/223	125/223
MONARCH-3 [12]	AI+ABEMA	328	First-line therapy postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.54 (0.41–0.72)	NA	158/328	256/328
	AI	165				57/165	118/165
MONARCH-plus [27]	AI+ABEMA	207	ET sensitive postmenopausal women with HR-positive/HER2-negative	0.50 (0.35–0.72)	NA	116/176	145/176
	AI	99				30/83	51/83
	FUL+ABEMA	104				40/80	62/80
SOLAR-1a [7]	FUL+ALP	169	First-line therapy men or postmenopausal women with HR-positive/HER2-negative advanced breast cancer with PI3K mutation	0.65 (0.50–0.85)	NA	45/169	104/169
	FUL	172				22/172	78/172
SOLAR-1b [7]	FUL+ALP	115	First-line therapy men or postmenopausal women with HR-positive/HER2-negative advanced breast cancer without PI3K mutation	0.85 (0.58–1.25)	NA	NA	NA
	FUL	116					
BELLE-2 [21]	FUL+BUP	576	Postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.78 (0.67–0.89)	0.87 (0.74–1.02)	68/576	252/576
	FUL	571				44/571	240/571
BELLE-3 [23]	FUL+BUP	289	Postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.67 (0.53–0.84)	NA	22/289	71/289
	FUL	143				3/143	22/143
SANDPIPER a [20]	FUL+TAS	340	Postmenopausal women with HR-positive/HER2-negative advanced breast cancer with PI3K mutation	0.70 (0.56–0.89)	NA	74/264	136/264
	FUL	176				16/134	50/134
SANDPIPER b [20]	FUL+TAS	77	Postmenopausal women with HR-positive/HER2-negative advanced breast cancer without PI3K mutation	0.69 (0.44–1.08)	NA	NA	NA
	FUL	38					
ACE [28]	AI+TUC	244	Postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.75 (0.58–0.98)	NA	45/244	114/244
	AI	121				11/121	43/121

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Table 1. (Continued) Characteristics of the outcomes of the studies included in this network meta-analysis

Study	Treatment	No. of patients	Characteristic	PFS (HR and 95% CI)	OS (HR and 95% CI)	ORR	CBR
BOLERO-2 [22,47]	AI+EVE AI	485 239	Postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.38 (0.31–0.48)	0.89 (0.73–1.10)	61/485 5/239	242/485 53/239
BOLERO-6 [8]	AI+ EVE EVE CAP	104 103 102	ET resistant postmenopausal women with HR-positive/HER2-negative advanced breast cancer	AI+EVE vs EVE: 0.73 (0.56–0.97); AI+EVE vs CAP: 1.15 (0.86–1.52)	NA	NA	NA
FALCON [26]	Ful AI	230 232	ET-naïve postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.80 (0.64–1.00)	0.88 (0.63–1.22)	93/230 90/232	180/230 172/232
SoFEA [39]	FUL+AI FUL	122 141	Postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.95 (0.75–1.22)	0.95 (0.76–1.17)	NA	NA
Paul et al. [35]	AI+DAS AI	57 63	Postmenopausal women with HR-positive/HER2-negative breast cancer with 0–1 prior chemotherapy and no prior AI for advanced disease	0.69 (0.43–1.09)	NA	NA	35/55 37/61
SWOG0226 [38]	FUL+AI AI	266 270	Postmenopausal women with HR-positive/HER2-negative untreated advanced breast cancer	0.81 (0.67–0.98)	0.81 (0.65–1.00)	NA	NA
PrE0102 [33]	FUL+EVE FUL	66 65	ET resistant postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.61 (0.40–0.92)	1.31 (0.72–2.38)	12/66 8/65	42/66 27/65
KCSG-BR10-04 [30]	FUL AI	44 47	Premenopausal women with HR-positive/HER2-negative tamoxifen-pretreated advanced breast cancer	NA	0.85 (0.27–2.74)	16/30 10/24	NA
KCSG-BR15-10 [34]	AI+PALBO CAP	92 92	Postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.64 (0.42–1.00)	NA	NA	NA
FERGI [25]	FUL+PIC FUL	88 79	ET resistant postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.74 (0.52–1.06)	NA	NA	NA
FAKTION [29]	FUL+CAP FUL	69 71	ET resistant postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.58 (0.39–0.84)	0.59 (0.34–1.05)	NA	NA
AROBASE [40]	AI+BEV PTX+BEV	58 59	Postmenopausal women with HR-positive/HER2-negative advanced breast cancer	1.00 (0.66–1.51)	NA	NA	NA
Musolino et al. [32]	FUL+DOV FUL	47 50	ET resistant postmenopausal women with HR-positive/HER2-negative advanced breast cancer	0.68 (0.41–1.41)	0.81 (0.39–1.65)	13/47 5/50	31/47 21/50
Howell et al. [19]	FUL TAM	313 274	Postmenopausal women with HR-positive/HER2-negative untreated advanced breast cancer	NA	NA	99/313 93/274	170/313 170/274
Dickler et al. [31]	AI+BEV AI	173 170	Women with HR-positive/HER2-negative advanced breast cancer	0.75 (0.59–0.96)	0.87 (0.65–1.18)	NA	NA

PFS = progression-free survival; HR = hazard ratio; CI = confidence interval; OS = overall survival; ORR = objective response rate; CBR = clinical benefit rate; FUL = fulvestrant; FUL+ABEMA = fulvestrant plus abemaciclib; FUL+AI = fulvestrant plus aromatase inhibitor; FUL+ALP = fulvestrant plus alpelisib; FUL+BUP = fulvestrant plus buparlisib; FUL+CAP = fulvestrant plus capecitabine; FUL+DOV = fulvestrant plus dovitinib; FUL+EVE = fulvestrant plus everolimus; FUL+PALBO = fulvestrant plus palbociclib; FUL+PIC = fulvestrant plus pictilisib; FUL+RIBO = fulvestrant plus ribociclib; FUL+TAS = fulvestrant plus taselisib; PTX+BEV = paclitaxel plus bevacizumab; TAM = tamoxifen; AI = aromatase inhibitor; AI+ABEMA = aromatase inhibitor plus abemaciclib; AI+BEV = aromatase inhibitor plus bevacizumab; AI+DAS = aromatase inhibitor plus dasatinib; AI+EVE = aromatase inhibitor plus everolimus; AI+PALBO = aromatase inhibitor plus palbociclib; AI+RIBO = aromatase inhibitor plus ribociclib; AI+TUC = aromatase inhibitor plus tucidinostat; CAP = capecitabine; EVE = everolimus; NA = not applicable; ET = endocrine therapy.

Two researchers also independently assessed the risk of bias in the final studies included using the Cochrane Collaboration's tool. The judging criteria were divided into 3 levels (low risk, high risk, and unclear), and no apparently low-quality studies were included. The detailed quality evaluation results are shown in **Supplementary Table 1**.

Outcomes

Progression-free survival

In this network meta-analysis, a total of 27 studies provided detailed data on PFS. Firstly, direct comparisons were calculated using a fixed-effects model to observe the efficacy of multiple treatments including AI or fulvestrant plus three CDK4/6 inhibitors and buparlisib, and the results showed that ET combined with targeted drugs was superior to monotherapy for inoperable patients. Next, an indirect comparison produced a similar conclusion, finding that AI or fulvestrant plus CDK4/6 inhibitors, PI3K inhibitors, everolimus, and capecitabine significantly improved the prognosis regardless of the type of CDK4/6 inhibitors, as shown

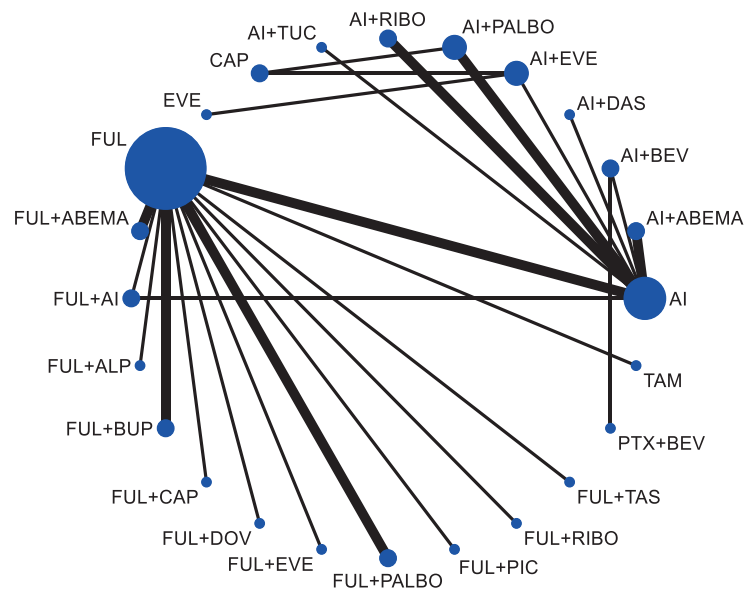


Figure 2. Network meta-analysis plot of direct comparison. Each circle in the plot represents a treatment option, and the size indicates the number of patients. Solid lines represent direct comparisons between different treatment options, and the thickness indicates the number of studies involved.

FUL = fulvestrant; FUL+ABEMA = fulvestrant plus abemaciclib; FUL+AI = fulvestrant plus aromatase inhibitor; FUL+ALP = fulvestrant plus alpelisib; FUL+BUP = fulvestrant plus buparlisib; FUL+CAP = fulvestrant plus capecitabine; FUL+DOV = fulvestrant plus dovitinib; FUL+EVE = fulvestrant plus everolimus; FUL+PALBO = fulvestrant plus palbociclib; FUL+PIC = fulvestrant plus pictilisib; FUL+RIBO = fulvestrant plus ribociclib; FUL+TAS = fulvestrant plus tasiselisib; PTX+BEV = paclitaxel plus bevacizumab; TAM = tamoxifen; AI = aromatase inhibitor; AI+ABEMA = aromatase inhibitor plus abemaciclib; AI+BEV = aromatase inhibitor plus bevacizumab; AI+DAS = aromatase inhibitor plus dasatinib; AI+EVE = aromatase inhibitor plus everolimus; AI+PALBO = aromatase inhibitor plus palbociclib; AI+RIBO = aromatase inhibitor plus ribociclib; AI+TUC = aromatase inhibitor plus tucidinostat; CAP = capecitabine; EVE = everolimus.

in **Figure 3**. However, compared with AI plus CDK4/6 inhibitors, the effect of fulvestrant combined with CDK4/6 inhibitors was superior; and, for everolimus, a combination with AI showed better efficiency than a combination with fulvestrant. Moreover, tucidinostat presented a disappointing result in that no significant difference was found between AI plus tucidinostat and monotherapy. Finally, the SUCRA curve revealed that AI plus everolimus and fulvestrant plus palbociclib were ranked first with similar probabilities (93.5% and 94.3%, respectively) followed by fulvestrant plus abemaciclib or ribociclib. Everolimus alone ranked last, having the worst effect compared to AI, fulvestrant, or capecitabine alone. The results of direct comparison and cumulative probability ranking are shown in **Supplementary Figure 1A** and **Figure 4A**, respectively.

Next, the patients were divided into two subgroups, a PIK3CA-mutated and a PIK3CA-unmutated group. The results indicated that fulvestrant combined with palbociclib or PI3K inhibitors including alpelisib, buparlisib, and tasiselisib had excellent efficacy compared to monotherapy in advanced breast cancer with PIK3CA mutation. However, for patients without PIK3CA mutation, only fulvestrant plus palbociclib or buparlisib prolonged the PFS compared to fulvestrant alone. Detailed results are shown in **Supplementary Figures 2** and **3**. Fulvestrant combined with palbociclib ranked first while fulvestrant alone ranked last, regardless of the status of PIK3CA mutation, as shown in **Supplementary Figure 4A** and **B**.

Similarly, an independent subgroup analysis was performed on ET-resistant participants according to response to prior therapy. Compared to fulvestrant monotherapy, fulvestrant

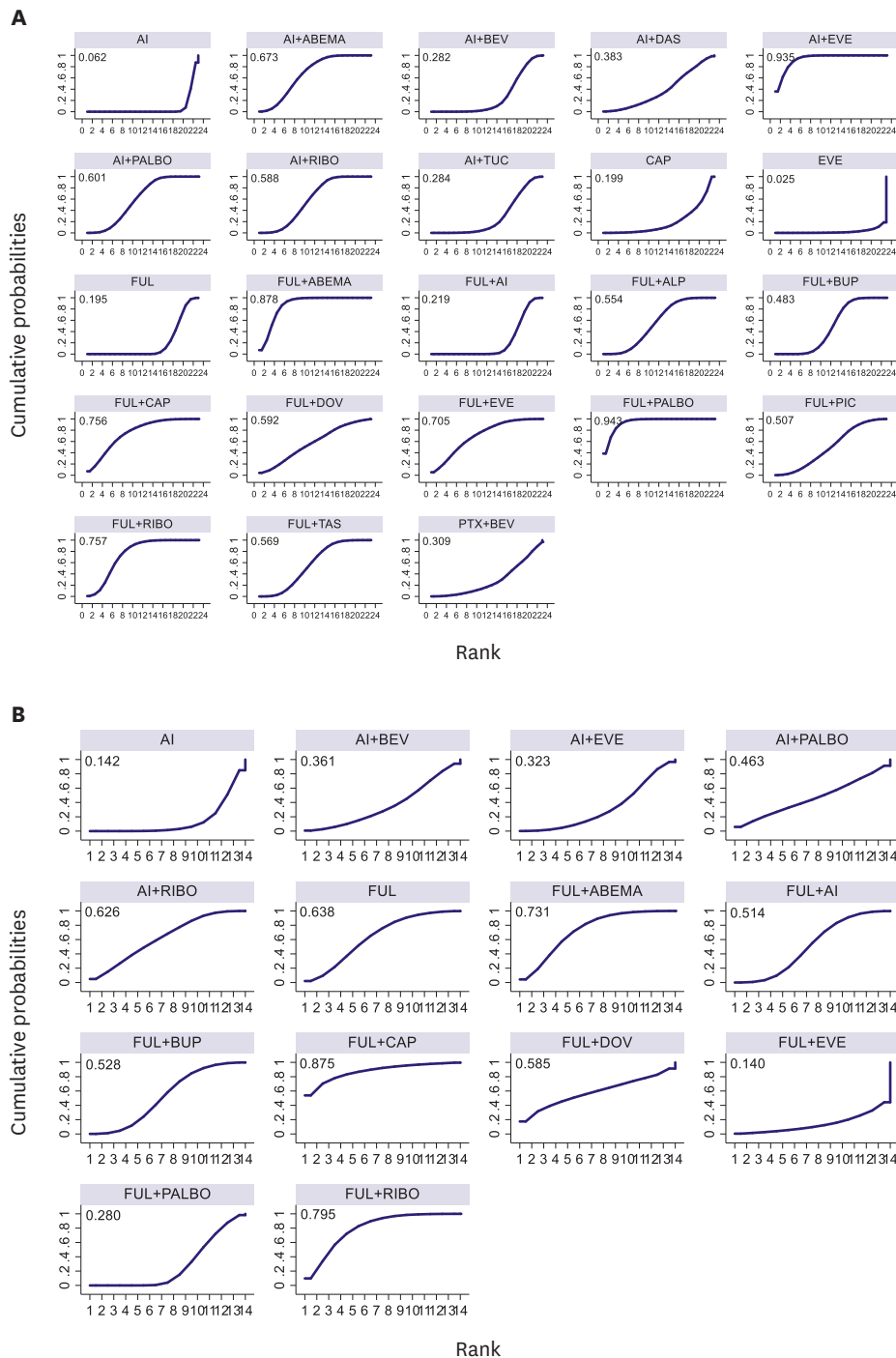


Figure 4. Surface under cumulative ranking curves of different endocrine treatment options. The horizontal axis represents the ranking, and the vertical axis represents the cumulative possibility. The larger the area under the curve, the more effective the treatment. (A) Curve of 23 treatment options for progression-free survival; (B) curve of 14 options for overall survival; (C) curve of 16 treatment options for objective response rate; (D) curve of 17 treatment options for clinical benefit rate; (E) curve of 14 treatment options for serious adverse events.

AI = aromatase inhibitor; AI+ABEMA = aromatase inhibitor plus abemaciclib; AI+BEV = aromatase inhibitor plus bevacizumab; AI+DAS = aromatase inhibitor plus dasatinib; AI+EVE = aromatase inhibitor plus everolimus; AI+PALBO = aromatase inhibitor plus palbociclib; AI+RIBO = aromatase inhibitor plus ribociclib; AI+TUC = aromatase inhibitor plus tucidostat; CAP = capecitabine; EVE = everolimus; FUL = fulvestrant; FUL+ABEMA = fulvestrant plus abemaciclib; FUL+AI = fulvestrant plus aromatase inhibitor; FUL+ALP = fulvestrant plus alpelisib; FUL+BUP = fulvestrant plus buparlisib; FUL+CAP = fulvestrant plus capecitabine; FUL+DOV = fulvestrant plus dovitinib; FUL+EVE = fulvestrant plus everolimus; FUL+PALBO = fulvestrant plus palbociclib; FUL+PIC = fulvestrant plus pictilisib; FUL+RIBO = fulvestrant plus ribociclib; FUL+TAS = fulvestrant plus taseleisib; PTX+BEV = paclitaxel plus bevacizumab; TAM = tamoxifen.

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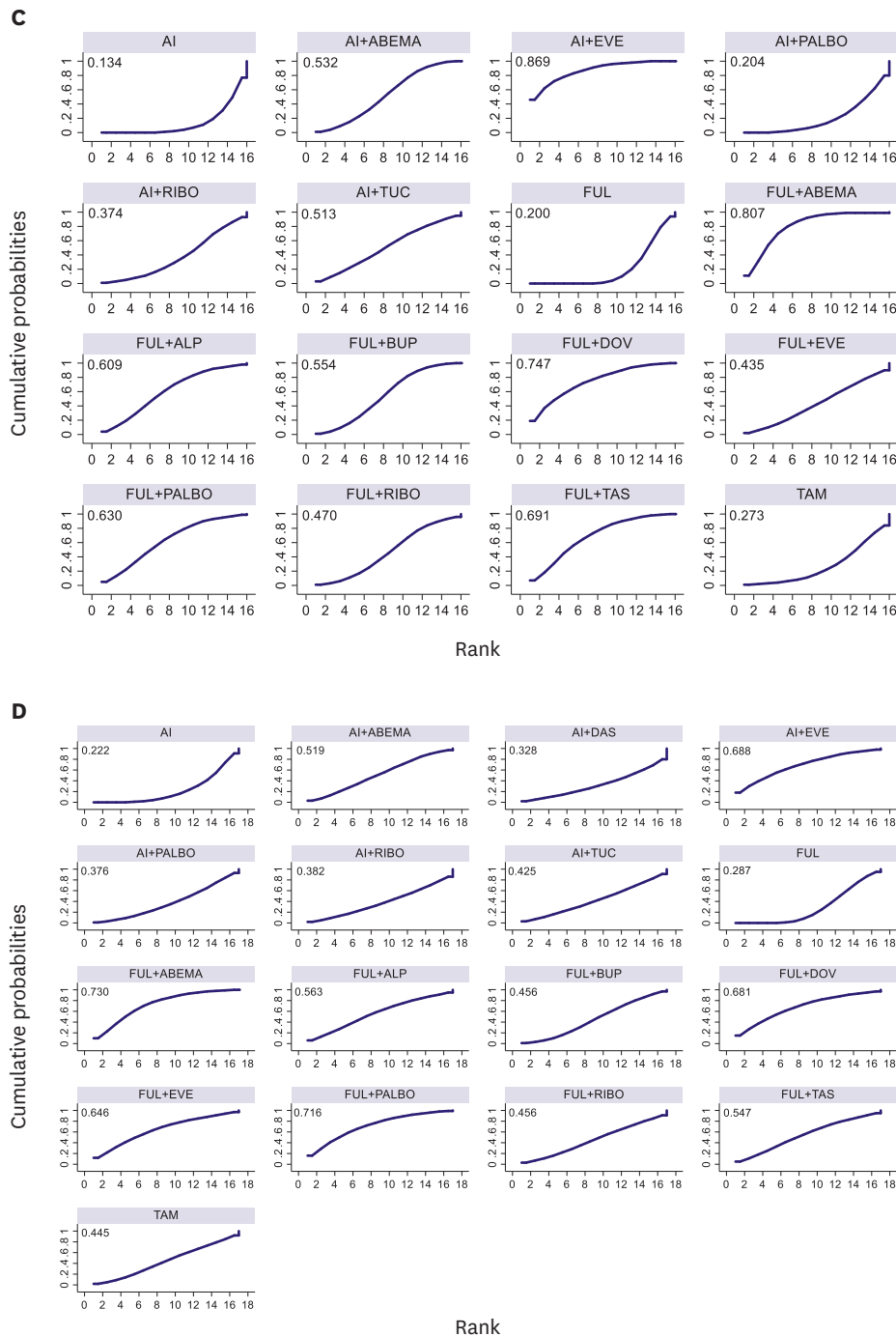


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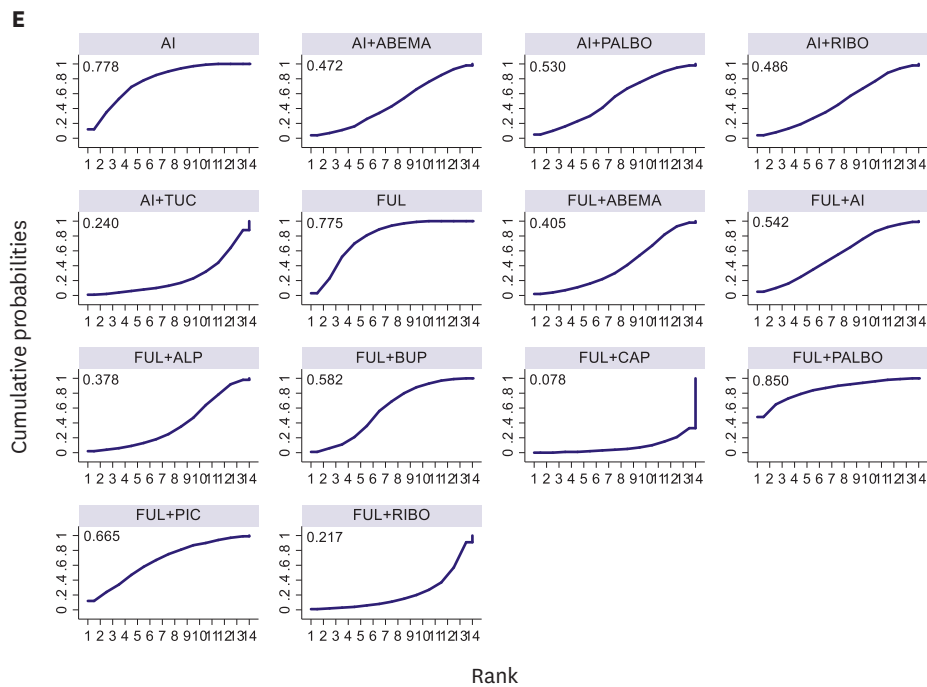


Figure 4. (Continued) Surface under cumulative ranking curves of different endocrine treatment options. The horizontal axis represents the ranking, and the vertical axis represents the cumulative possibility. The larger the area under the curve, the more effective the treatment. (A) Curve of 23 treatment options for progression-free survival; (B) curve of 14 options for overall survival; (C) curve of 16 treatment options for objective response rate; (D) curve of 17 treatment options for clinical benefit rate; (E) curve of 14 treatment options for serious adverse events.

AI = aromatase inhibitor; AI+ABEMA = aromatase inhibitor plus abemaciclib; AI+BEV = aromatase inhibitor plus bevacizumab; AI+DAS = aromatase inhibitor plus dasatinib; AI+EVE = aromatase inhibitor plus everolimus; AI+PALBO = aromatase inhibitor plus palbociclib; AI+RIBO = aromatase inhibitor plus ribociclib; AI+TUC = aromatase inhibitor plus tucidinstat; CAP = capecitabine; EVE = everolimus; FUL = fulvestrant; FUL+ABEMA = fulvestrant plus abemaciclib; FUL+AI = fulvestrant plus aromatase inhibitor; FUL+ALP = fulvestrant plus alpelisib; FUL+BUP = fulvestrant plus buparlisib; FUL+CAP = fulvestrant plus capecitabine; FUL+DOV = fulvestrant plus dovitinib; FUL+EVE = fulvestrant plus everolimus; FUL+PALBO = fulvestrant plus palbociclib; FUL+PIC = fulvestrant plus pictilisib; FUL+RIBO = fulvestrant plus ribociclib; FUL+TAS = fulvestrant plus taselisib; PTX+BEV = paclitaxel plus bevacizumab; TAM = tamoxifen.

plus CDK4/6 inhibitors or PIK3CA inhibitors achieved excellent outcomes for all except pictilisib, as shown in **Supplementary Figure 5**. Similar to the above results, fulvestrant combined with palbociclib still ranked first, followed by fulvestrant plus abemaciclib and fulvestrant plus capecitabine, as shown in **Supplementary Figure 4C**.

Based on the location of the metastatic setting, we extracted PFS data from patients with visceral metastases and presented a subgroup analysis. Compared with endocrine monotherapy, AI or fulvestrant combined with targeted drugs was found to be capable of significantly prolonging PFS in visceral metastasis patients except in the cases of tucidinstat and pictilisib (**Supplementary Figure 6**). In terms of rankings, the results suggested that fulvestrant plus palbociclib or abemaciclib ranked first (86.2% and 84.7%, respectively) followed by fulvestrant plus buparlisib, and monotherapies including AI and fulvestrant ranked last (**Supplementary Figure 4D**).

Overall survival

In addition, a total of 16 clinical studies contributed data on OS involving 14 different treatment options. A fixed-effects model was applied for direct comparison due to the low heterogeneity, and only AI plus ribociclib was linked to an effect of prolonging survival time (**Supplementary Figure 1B**). Similarly, an indirect comparison was performed, and this suggested that fulvestrant plus ribociclib, abemaciclib, or capecitabine, and AI plus ribociclib

can improve OS compared to AI alone, and only the results of fulvestrant combined with ribociclib or abemaciclib were statistically different compared to fulvestrant alone (**Figure 5**). Unexpectedly, fulvestrant plus capecitabine showed the highest probability to rank first, followed by fulvestrant combined with ribociclib or abemaciclib (**Figure 4B**).

Objective response rate

Nineteen studies provide detailed data on ORR, involving 16 different treatment options. Direct comparison was performed using a random-effects model due to the obvious heterogeneity of the data, and the results suggested the effectiveness of CDK4/6 inhibitors, including palbociclib and abemaciclib, for advanced breast cancer in ORR (**Supplementary Figure 1C**). Next, we continuously calculated the results of indirect comparison and cumulative probability of ranking in different treatment strategies. In the indirect comparison, AI combined with everolimus and fulvestrant combined with abemaciclib resulted in higher ORR to some extent (**Figure 6**). **Figure 4C** presents the ranking results, showing that AI plus everolimus had the highest probability to rank first followed by fulvestrant combined with abemaciclib.

Clinical benefit rate

In this network meta-analysis, a total of 19 clinical trials involving 17 different treatments provided detailed data on CBR. The direct comparison results from a random-effects model showed that fulvestrant plus abemaciclib can significantly improve the CBR in advanced breast cancer. The indirect comparison found no significant differences, and fulvestrant plus palbociclib, fulvestrant plus abemaciclib had the highest probability to rank first (71.6% and 73.0%, respectively), followed by AI plus everolimus, as shown in **Figures 4D** and **7**.

Safety

Data on serious adverse events were extracted from articles to evaluate the safety of different treatment options. The results of indirect comparison clearly showed that only fulvestrant combined with capecitabine was significantly different compared with fulvestrant alone, in that the former had a higher incidence of serious adverse events (**Figure 8**). Moreover, fulvestrant plus palbociclib showed the best safety and curative efficacy according to the SUCRA curve (**Figure 4E**).

DISCUSSION

Hormone receptor-positive/HER2-negative is the most common subtype of breast cancer, and ET is the preferred treatment, as it has proven efficacy and tolerable toxicity (except in life-threatening situations) [49]. Recently, many targeted drugs, such as CDK4/6 inhibitors, mTOR inhibitors, PI3KCA inhibitors, etc., have proved to be effective *in vivo* or *in vitro* experiments, particularly against advanced hormone receptor-positive/HER2-negative breast cancer. Although targeted drugs with different mechanisms have exhibited superior curative effects leading to improved prognosis, no study has previously carried out a direct comparison of the efficacy and safety of different treatment options. Therefore, we performed a network meta-analysis to indirectly compare 24 different treatment strategies to determine which options are most appropriate under specific circumstances.

Currently, the crucial role of the PI3K/AKT/mTOR pathway in stimulating tumor cell proliferation, changing metabolism, and inhibiting apoptosis is widely accepted in breast

AI	AI+ABEMA	AI+PALBO	AI+RIBO	AI+TUC	FUL	FUL+ABEMA	FUL+AI	FUL+ALP	FUL+BUP	FUL+CAP	FUL+PALBO	FUL+PIC	FUL+RIBO
2.13 (0.24, 24.15)	0.77 (0.03, 16.36)	1.27 (0.05, 29.72)	2.31 (0.09, 54.13)	0.22 (0.01, 5.68)	2.48 (0.25, 24.34)	0.67 (0.03, 16.29)	1.61 (0.06, 37.78)	0.59 (0.04, 8.84)	14.19 (0.67, 556.90)	0.03 (0.00, 1.08)	1.77 (0.08, 45.74)	3.65 (0.14, 78.27)	
1.63 (0.17, 14.48)	0.97 (0.04, 21.04)	2.85 (0.11, 66.19)	0.48 (0.02, 12.67)	0.56 (0.01, 27.69)	1.67 (0.17, 15.99)	1.07 (0.04, 24.52)	0.93 (0.06, 16.09)	8.21 (0.27, 431.84)	0.44 (0.03, 7.31)	0.06 (0.00, 1.98)	6.32 (0.27, 166.88)		
2.05 (0.20, 19.16)	2.17 (0.07, 48.94)	0.62 (0.02, 14.15)	1.22 (0.02, 66.94)	0.37 (0.01, 18.98)	2.65 (0.26, 25.13)	1.07 (0.04, 24.52)	0.64 (0.04, 9.98)	13.73 (0.41, 782.41)	0.79 (0.05, 14.78)	0.20 (0.00, 6.49)			
4.69 (0.42, 45.73)	0.47 (0.02, 9.76)	1.52 (0.03, 68.91)	0.85 (0.02, 42.08)	0.57 (0.01, 31.16)	1.59 (0.32, 7.39)	0.64 (0.04, 9.98)	1.61 (0.06, 37.78)	0.26 (0.01, 7.01)	0.71 (0.03, 21.22)	0.20 (0.00, 6.49)			
1.00 (0.10, 8.44)	1.19 (0.02, 46.72)	1.05 (0.02, 51.03)	1.27 (0.03, 63.93)	0.36 (0.01, 12.87)	2.212 (1.57, 610.01)	0.70 (0.06, 6.85)	0.75 (0.02, 27.32)	0.40 (0.02, 10.12)	2.62 (0.13, 68.75)	0.20 (0.00, 6.49)			
2.48 (0.09, 60.86)	0.80 (0.01, 34.14)	1.66 (0.03, 77.34)	0.75 (0.02, 27.32)	0.468 (0.08, 446.04)	1.75 (0.07, 48.21)	0.49 (0.02, 13.43)	0.93 (0.06, 16.09)	0.71 (0.03, 21.22)	0.79 (0.05, 14.78)	0.20 (0.00, 6.49)			
1.72 (0.07, 43.52)	1.27 (0.02, 54.01)	0.98 (0.03, 35.53)	1.078 (0.17, 1037.65)	0.16 (0.00, 6.85)	4.36 (0.52, 43.24)	0.96 (0.02, 54.88)	1.59 (0.32, 7.39)	0.28 (0.01, 7.44)	2.62 (0.13, 68.75)	0.20 (0.00, 6.49)			
2.65 (0.10, 56.94)	0.77 (0.02, 22.27)	13.29 (0.21, 1256.25)	0.78 (0.01, 18.21)	0.27 (0.01, 18.42)	0.36 (0.01, 16.38)	2.15 (0.04, 115.14)	1.59 (0.32, 7.39)	0.28 (0.01, 7.44)	2.62 (0.13, 68.75)	0.20 (0.00, 6.49)			
1.56 (0.09, 24.32)	0.77 (0.02, 22.27)	13.29 (0.21, 1256.25)	0.78 (0.01, 18.21)	0.27 (0.01, 18.42)	0.36 (0.01, 16.38)	2.15 (0.04, 115.14)	1.59 (0.32, 7.39)	0.28 (0.01, 7.44)	2.62 (0.13, 68.75)	0.20 (0.00, 6.49)			
22.40 (0.70, 1125.76)	0.42 (0.01, 21.85)	2.36 (0.05, 118.23)	0.34 (0.01, 18.21)	0.16 (0.00, 6.85)	4.36 (0.52, 43.24)	0.96 (0.02, 54.88)	1.59 (0.32, 7.39)	0.28 (0.01, 7.44)	2.62 (0.13, 68.75)	0.20 (0.00, 6.49)			
0.69 (0.03, 18.72)	0.76 (0.02, 42.82)	1.32 (0.02, 63.54)	0.62 (0.01, 37.71)	0.27 (0.01, 18.42)	0.36 (0.01, 16.38)	2.15 (0.04, 115.14)	1.59 (0.32, 7.39)	0.28 (0.01, 7.44)	2.62 (0.13, 68.75)	0.20 (0.00, 6.49)			
1.26 (0.05, 34.64)	2.76 (0.06, 128.20)	0.36 (0.01, 16.38)	2.15 (0.04, 115.14)	0.96 (0.02, 54.88)	4.36 (0.52, 43.24)	0.96 (0.02, 54.88)	1.59 (0.32, 7.39)	0.28 (0.01, 7.44)	2.62 (0.13, 68.75)	0.20 (0.00, 6.49)			
4.44 (0.19, 112.88)	2.76 (0.06, 128.20)	0.36 (0.01, 16.38)	2.15 (0.04, 115.14)	0.96 (0.02, 54.88)	4.36 (0.52, 43.24)	0.96 (0.02, 54.88)	1.59 (0.32, 7.39)	0.28 (0.01, 7.44)	2.62 (0.13, 68.75)	0.20 (0.00, 6.49)			

Figure 8. Indirect comparison of 14 different treatment options in terms of serious adverse events. The values represent HR (95% CI) and bold font indicates that the results had significant difference.

HR = hazard ratio; CI = confidence interval; AI = aromatase inhibitor; AI+ABEMA = aromatase inhibitor plus abemaciclib; AI+PALBO = aromatase inhibitor plus palbociclib; AI+RIBO = aromatase inhibitor plus ribociclib; AI+TUC = aromatase inhibitor plus tucidinostat; FUL = fulvestrant; FUL+ABEMA = fulvestrant plus abemaciclib; FUL+AI = fulvestrant plus aromatase inhibitor; FUL+ALP = fulvestrant plus alpelisib; FUL+BUP = fulvestrant plus buparlisib; FUL+CAP = fulvestrant plus capecitabine; FUL+PALBO = fulvestrant plus palbociclib; FUL+PIC = fulvestrant plus pictilisib; FUL+RIBO = fulvestrant plus ribociclib.

cancer [50]. Some potential interactions between estrogen receptors and the PI3K/AKT/mTOR pathway have been documented [51]. Long-term poverty of estrogen is closely related to increased phosphorylation of AKT and activity of mTOR, which is the mechanism by which tumor cells adapt to low estrogen levels and develop resistance to antiestrogen therapeutics [52]. Everolimus, an oral rapamycin derivative, can suppress PI3K/AKT/mTOR cascades to inhibit downstream signaling events, and exhibits anti-tumor activity when combined with AI in advanced breast cancer, particularly in the presence of endocrine resistance [53]. According to our results, although fulvestrant combined with everolimus also exhibited the capacity to prolong the PFS and improve clinical efficacy, AI plus everolimus showed the highest probability of ranking first terms of its effect on PFS and ORR, while no notable effect was found in terms of OS. Moreover, the results showed that everolimus should certainly not be used alone due to poor efficacy compared with AI or fulvestrant alone. PI3k inhibitors (alpelisib, buparlisib, taselisib, and pictilisib) similarly repressed the PI3K/AKT/mTOR pathway, but tumor regression depended on the presence of PIK3CA mutation, unlike in the case of everolimus [54]. In our results, although PI3K inhibitors (except pictilisib) combined with fulvestrant improved the prognosis compared with endocrine monotherapy, the efficacy was still inferior to that of CDK4/6 inhibitors combined with AI or fulvestrant, regardless of PIK3CA-mutation status.

Another crucial pathway, CDK-Rb-E2F, accelerates G1-S transformation through association with increased CDK4/6 activity [55], and p16^{INK4A}, a suppressor for CDK4/6, is a particularly important protein for inducing cell arrest at G1. Gene mutation of p16^{INK4A} or gene amplification of CDK4/6 both activate the CDK-Rb-E2F pathway, leading to excessively cellular proliferation [7], and could provide a potential therapeutic target for the inhibition of CDK4/6 activity. The highly selective inhibitors of CDK4/6 kinases (palbociclib, ribociclib, and abemaciclib) block Rb phosphorylation, reducing cytotaxis in G1 in luminal breast cancer [56-58]. In this network meta-analysis, although no significant differences were observed in the three CDK4/6 inhibitors combined with AI or fulvestrant, the effects of fulvestrant combined with CDK4/6 inhibitors was slightly better than those of AI plus CDK4/6 inhibitors. The evaluation of OS showed that the efficacy of co-treatment with ribociclib or abemaciclib was superior to that of palbociclib, although no significant difference was found in indirect comparisons. Finally, our safety results indicated that palbociclib, whether combined with AI or fulvestrant, had a more better safety record compared to the other 2 CDK4/6 inhibitors. In addition, our results indicated that the efficacy of palbociclib alone in advanced breast cancer was disappointing. It is worth noting that abemaciclib has been approved to treat breast cancer alone, unlike palbociclib and ribociclib [59].

The results of indirect analysis also included AI plus tyrosine kinase inhibitor, AI plus histone deacetylase (HDAC) inhibitor, AI plus VEGF inhibitor, capecitabine, fulvestrant plus capecitabine, and fulvestrant plus AI. Capecitabine is commonly used as an oral chemotherapy drug for breast cancer patients who have progressed during anti-estrogen therapy, and fulvestrant combined with capecitabine had the most significant effect on prolonging OS, with slightly improved PFS based on our results. The oral HDAC inhibitor (tucidinostat) and dovitinib can promote differentiation and cell cycle arrest, and regulate the tumor microenvironment by inhibiting HDAC1, HDAC2, HDAC3, and HDAC10, and by blocking fibroblast growth factor receptor and the respective downstream signaling pathways of all these proteins [60-62]. Our results demonstrated that AI plus bevacizumab, tucidinostat, or fulvestrant, and fulvestrant combined with dovitinib improved clinical outcomes compared with AI alone, but no significant differences were detected compared

with fulvestrant alone. Because the dosage of fulvestrant was half of the standard usage in those studies which provided detailed data of fulvestrant combined with AI, the efficacy of fulvestrant plus AI may be more remarkable. In conclusion, the above treatment agents are not recommended for first-line treatment. When the patient responds poorly to multiple treatment strategies, including CDK4/6 inhibitors and everolimus, fulvestrant in combination with capecitabine or dovitinib or AI, or AI in combination with bevacizumab or tucidinostat can be used to achieve a longer survival time.

Based on PIK3CA mutations, ET resistance and visceral metastasis, participants were divided into several subgroups to evaluate which ET regimen was more effective in different situations. In patients with PIK3CA mutations, fulvestrant in combination with palbociclib, alpelisib, buparlisib, or taselisib exhibited excellent curative effect on PFS compared to monotherapy, but pictilisib did not. Fulvestrant plus palbociclib was the most likely to rank first, indicating that CDK4/6 inhibitors still had superior efficacy for patients with PIK3CA mutations. Therefore, PI3K inhibitors can be used to treat patients with hormone receptor-positive/HER2-negative advanced breast cancer who are resistant to CDK4/6 inhibitors. For patients without PIK3CA mutations, only palbociclib and buparlisib exhibited an effect of improving the prognosis compared with fulvestrant alone.

Intrinsic or acquired endocrine resistance remains a tough challenge for breast cancer treatment [63]. Many preclinical models have been used to attempt to elucidate the mechanisms of endocrine resistance. At present, researchers have established that disorder of the PI3K/AKT/mTOR axis and CDK/Rb/E2F pathway is the critical mechanism leading to endocrine resistance [64]. In this meta-analysis, fulvestrant combined with palbociclib, abemaciclib, alpelisib, buparlisib, everolimus, or capecitabine showed efficacy in prolonging PFS for patients with ET resistance compared with fulvestrant alone. Although there was no significant difference between the 2 PI3K inhibitors, the efficacy of alpelisib was slightly better than that of buparlisib. Similarly, fulvestrant combined with CDK4/6 inhibitors is the preferred option for ET-resistant patients. Moreover, the results from the visceral metastasis subgroup suggested that fulvestrant combined with CDK4/6 inhibitors or PI3K inhibitors (alpelisib, buparlisib), and AI combined with CDK4/6 inhibitors or tucidinostat can improve the prognosis to different degrees compared with endocrine monotherapy. Although significant differences were not found among the three CDK4/6 inhibitors, the efficacy of palbociclib and abemaciclib was slightly better than that of ribociclib for advanced breast cancer patients with visceral metastasis. Similarly, the curative effect of buparlisib was slightly better to that of alpelisib, although there was no statistical difference.

Our study included a large number of treatment options for hormone receptor-positive/HER2-negative advanced breast cancer, and involved subgroup analysis based on the status of PIK3CA mutation, ET resistance, and visceral metastasis, which no previous study has done [16]. However, there are still many limitations in this network meta-analysis. Firstly, according to the BOLERO-2 study, AI combined with everolimus was extremely effective in patients with visceral metastasis or resistance to ET [22], but everolimus could not be indirectly compared in the ET resistance subgroup and visceral metastasis subgroup due to the lack of complete data. Secondly, we did not distinguish between different dosages of drugs. For example, the standard dosage of fulvestrant was defined as 500 mg fulvestrant on days 1 and 15 of cycle one and then on day one of each subsequent cycle (28 days) [65]. However, the dosage of fulvestrant used in SoFEA and SWOG0226 was 500 mg via intramuscular injection on day 1 followed by 250 mg injections on days 15 and 29, and then

250 mg intramuscular injections were done every 28 days [38,39]. Therefore, the use of non-standard fulvestrant dosages may have influenced the results of indirect comparison. Thirdly, different treatment options were not calculated for premenopausal and perimenopausal patients due to the lack of detailed data. About 85% of the participants in this meta-analysis were post-menopausal, and there may be some differences in the treatment plan for patients with different menstrual conditions.

CONCLUSION

In conclusion, our meta-analysis demonstrated that fulvestrant or AI combined with targeted drugs or other anti-tumor drugs can generally improve the prognosis for hormone receptor-positive/HER2-negative advanced breast cancer with controllable adverse events. Among the options analyzed, the effect of AI combined with everolimus was the most effective for the amelioration of PFS, followed by fulvestrant combined with abemaciclib or palbociclib, and the latter had the smallest incidence of serious adverse events compared to the other 14 treatments. Fulvestrant combined with capecitabine can be used when patients are resistant to everolimus and CDK4/6 inhibitors due to its superior effect on PFS and OS, but application should be scrupulously considered due to its obvious toxic effects. Moreover, fulvestrant combined with CDK4/6 inhibitors was shown to be more effective than AI combined with CDK4/6, and for co-treatment with everolimus, AI combined with everolimus was slightly more effective. PI3K inhibitors are recommended for second-line treatment of patients with PIK3CA mutations.

SUPPLEMENTARY MATERIALS

Supplementary Reference

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Supplementary Table 1

The quality evaluation of included studies

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Supplementary Figure 1

Forest plots of direct comparison. (A) the forest plot of direct comparison for progression-free survival; (B) the forest plot of direct comparison for overall survival; (C) the forest plot of direct comparison for objective response rate; (D) the forest plot of direct comparison for clinical benefit rate.

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Supplementary Figure 2

Indirect comparison of PIK3CA-mutated patients in progression-free survival. The values represent HR (95% CI) and bold font indicates that the results had significant statistical difference.

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Supplementary Figure 3

Indirect comparison of PIK3CA-unmutated patients in progression-free survival. The values represent HR (95% CI) and bold font indicates that the results had significant statistical difference.

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Supplementary Figure 4

Surface under cumulative ranking curve of different endocrine treatment options for progression-free survival. The horizontal ordinate represents the ranking, and the ordinate represents the cumulative possibility. The larger the area under the curve, the better the treatment effect. (A) curve of PI3KCA-mutated patients for progression-free survival; (B) curve of PI3KCA-unmutated patients for progression-free survival; (C) curve of ET resistant patients for progression-free survival; (D) curve of visceral metastasis patients for progression-free survival.

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Supplementary Figure 5

Indirect comparison of ET resistant patients in progression-free survival. The values represent HR (95% CI) and bold font indicates that the results had significant statistical difference.

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Supplementary Figure 6

Indirect comparison of visceral metastases patients in progression-free survival. The values represent HR (95% CI) and bold font indicates that the results had significant statistical difference.

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