# Multiple-Treatments Meta-analysis of Chemotherapy and Targeted Therapies in Advanced Breast Cancer

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- **Background** Many systemic nonhormonal regimens have been evaluated across several hundreds of randomized trials in advanced breast cancer. We aimed to quantify the relative merits of these regimens in prolonging survival.
  - Methods We performed a systematic review of all trials that compared different regimens involving chemotherapy and/or targeted therapy in advanced breast cancer (1973–2007). Regimens were categorized a priori into different treatment types. We performed multiple-treatments meta-analysis and calculated hazard ratios for each treatment category relative to monotherapy with old agents (ie, regimens not including anthracy-clines, anthracenediones, vinorelbine, gemcitabine, capecitabine, taxanes, marimastat, thalidomide, trastuzumab, lapatinib, or bevacizumab).
  - **Results** We identified 370 eligible randomized trials (54189 patients), of which 172 (31552 patients) compared different types of treatment. Survival data from 148 comparisons pertaining to 128 of the 172 trials (26031 patients, 22 different types of treatment) were available for inclusion in the multiple-treatments metaanalysis. Compared with single-agent chemotherapy with old nonanthracycline drugs, anthracycline regimens achieved 22%–33% relative risk reductions in mortality (ie, hazard ratio [HR] for standard-dose anthracycline-based combination: 0.67, 95% credibility interval [Crl] 0.57–0.78). Several newer regimens achieved further benefits (eg, HR [95% Crl] 0.67 [0.55–0.81] for single-drug taxane, 0.64 [0.53–0.78] for combination of anthracyclines with taxane, 0.49 [0.37–0.67] for taxane-based combination with capecitabine or gemcitabine), and similar benefits were seen with several regimens including molecular targeted treatments. Most regimens had very similar efficacy profiles (<5% difference in HR) as first- and subsequentline therapies.
- **Conclusions** Stepwise improvements in efficacy of chemotherapy and targeted treatments cumulatively have achieved major improvements in the survival of patients with advanced breast cancer. Many options that can be used in first and subsequent lines of therapy have comparable efficacy profiles.
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The morbidity band mortality impact of breast cancer is very large worldwide (1-3) and treatment can be challenging. Many pharmacological compounds and administration modalities have been developed for the treatment of advanced breast cancer (4). Key milestones in the systemic treatments for breast cancer were the introduction of hormonal treatment in the 1940s (5), combined chemotherapy in 1969, and anthracyclines in 1972 (6). In the 1990s, gemcitabine, vinorelbine, and capecitabine also entered clinical care, followed by taxanes. Additional expensive targeted treatments were introduced in the last decade.

Although progress has been achieved in the field and patients live longer, the relative merits of the many different chemotherapy and targeted treatment regimens are not well understood. Hundreds of trials have been conducted to compare treatments for advanced breast cancer, but because each has compared only two or a few treatments, it is difficult to integrate information on the relative efficacy of all tested regimens. This integration is important because different regimens vary both in cost and in toxicity. Therefore, we performed a comprehensive systematic review of chemotherapy and targeted treatment regimens in advanced breast cancer and evaluated, through a multiple-treatments meta-analysis (7,8), the relative merits of the many different regimens used to prolong survival in advanced breast cancer patients. Data were analyzed on all eligible trials as well as separately for first- and subsequent-line treatment.

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## Methods

## **Identification of Randomized Studies**

We searched for randomized studies in any language in PubMed and the Cochrane Central Registry of Controlled Trials looking for studies with the key words mammary or breast that also contained the search terms cancer, malign\*, neoplasm\*, or carcinoma\*. The last search update was done on October 9, 2007. PubMed searches used the restriction limit: randomized controlled trial. We searched the reference lists of every primary study and also retrieved previous meta-analyses addressing comparisons of regimens in advanced breast cancer. Considering the recent introduction of targeted treatment agents and the paucity of information available to date on these agents, we ensured that electronic searches would not miss any major reports of eligible studies by hand-searching the volumes published from 2004 to 2007 inclusive of the three journals with the highest number of electronically identified cancer trials (9). Furthermore, because recent trials with novel or targeted treatment agents may still be unpublished, we also reviewed abstract books and presentations of major recent meetings in 2006-2007 of the American Society of Clinical Oncology, the European Society of Clinical Oncology, and the European Cancer Conference to identify any other trial that had presented final (not preliminary) data on drugs that were already studied in the published trials and comparisons that would be eligible for consideration in the multiple-treatments meta-analysis. Earlier meeting abstracts were not included.

## **Eligibility Criteria**

We considered all randomized trials that compared at least two arms of different regimens involving chemotherapy and/or targeted treatments in patients with advanced breast adenocarcinoma in any line of treatment. Advanced-stage disease was defined by the presence of metastatic or recurrent disease that was not amenable to surgical treatment. We excluded trials on earlier stages of the disease (when trials included both advanced and earlier stage, we included only the relevant population) and trials comparing regimens in breast malignancies other than adenocarcinoma (inflammatory breast cancer, sarcoma, etc). We also excluded nonrandomized and pseudo-randomized trials (eg, those with alternate allocation of subjects). If trials included other concomitant interventions such as hormonal treatment, surgery, radiotherapy, or radioisotopic treatment that differed systematically between the investigated arms, they were excluded. Whenever reports pertained to sets of patients that overlapped, only the report with longest follow-up (having the largest number of events) was used in the analysis.

## **Data Extraction**

From each eligible trial, we recorded authors' names, journal and year of publication, country of origin (as noted in their affiliations), years of patient enrolment, sample size (randomized and analyzed) per arm, performance status, regimens used, line of treatment, additional treatments given to both arms, and information pertaining to study design (whether the trial reported the mode of randomization, allocation concealment, description of withdrawals per arm, and blinding). We also recorded the median survival per

# CONTEXT AND CAVEATS

### Prior knowledge

Although many nonhormonal regimens for advanced breast cancer have been evaluated across hundreds of randomized clinical trials, the relative merits of these regimens were not known with precision.

### Study design

Multiple-treatments meta-analysis of survival data based upon direct and indirect comparisons to estimate hazard ratios for different chemotherapy regimens as first- and subsequent-line treatments relative to older nonanthracycline single-agent therapy.

## Contribution

This study quantified the survival gains of 21 different classes of therapy relative to older single-agent therapy.

## Implications

Whether used in first or subsequent lines of therapy, many classes of modern breast cancer therapy, including anthracyclines, taxanes, novel non-taxane agents, and molecular targeted therapies used in the context of single-agent or combination therapy, produce gains in absolute survival over older single agents.

## Limitations

A comprehensive understanding of the effectiveness of newer molecular targeted therapies will require additional trial information. The extent of intertrial variability and its effect on the results of the meta-analysis cannot be known with certainty.

From the Editors

arm and whether there was any statistically significant difference (P < .05) in median survival between any compared arms.

For trials comparing different types of treatment (as defined below), we also extracted or estimated the logarithm of the hazard ratio for death and its variance. We used the hazard ratio and 95% confidence intervals from Cox regressions, as reported or retrieved by contacting the investigators. Unadjusted hazard ratios were preferred over multivariable ones. If only the variance of the hazard ratio was unavailable, we calculated it using the log-rank P value. When neither hazard ratio nor variance was available, we estimated the variance as  $(T_1 + T_2)^2/[(E_1 + E_2)T_1T_2]$ , where  $E_1$  and  $E_2$  are the number of events and  $T_1$  and  $T_2$  the number of randomly assigned patients in each arm, and then estimated the log(HR), such that it would have the P value of the log-rank test (10). When P values were unavailable, the hazard ratio was approximated by the ratio of median survivals.

Data were independently extracted by two investigators (D. Mauri and N. P. Polyzos). Discrepancies were discussed with a third investigator (J. P. A. Ioannidis). J. P. A. Ioannidis and G. Salanti also cross-checked against the original articles all the data entered into calculations.

## **Categories for Analyses**

In our analysis, we grouped the regimens into the following nine categories: anthracyclines, anthracenediones, non-taxane novel chemotherapy agents (vinorelbine, gemcitabine, or capecitabine), taxanes, marimastat, thalidomide, trastuzumab, lapatinib, and bevacizumab. Other compounds, such as cyclophosphamide,

fluorouracil, methotrexate, mitomycin, vincristine, and vinblastine, and all other agents not included in the previous categories were grouped in a miscellaneous category as old agents.

Considering that combined chemotherapy has been demonstrated to produce statistically significant survival advantages over monotherapy (4,11), all categories of drugs analyzed were further split into combination therapies (when two or more different agents were given together) or monotherapies. Regimens using lower dosages of anthracyclines may be inferior to those using higher doses (4); thus, anthracycline-based regimens were further split into low and standard vs high doses. The anthracycline dose equivalence for doxorubicin and epirubicin was derived from isodose studies (12). For monotherapy, we considered as low dose a 21-day dose intensity of less than 60 mg/m<sup>2</sup> doxorubicin or less than 78 mg/m<sup>2</sup> epirubicin. For combination treatment, we designated low dose as a 21-day dose intensity of less than 45 mg/m<sup>2</sup> doxorubicin or less than 58 mg/m<sup>2</sup> epirubicin. For mitoxantrone monotherapy, we designated as low dose a 21-day dose intensity of less than 12 mg/m<sup>2</sup>; a low dose for combination regimens was a 21-day dose intensity of less than 8 mg/m<sup>2</sup>.

#### **Statistical Analyses**

We generated descriptive statistics for trial and study population characteristics across all eligible trials. We described the types of comparisons and how these had evolved over time. We used the probability of interspecific encounter (PIE) and PIE' metrics to evaluate the diversity of the network, and the C metric to measure co-occurrence. Details of these metrics have been described previously (13). In brief, a PIE of less than 0.75 suggests that there is lack of diversity, which could be due to either few regimens or large unevenness of available evidence across the represented regimens. The PIE' index divides PIE by the maximum possible PIE given the number of regimens and thus removes the effect of the number of regimens in the estimation of diversity. The C test, based on a permutation procedure, examines whether there are pairs of regimens in the networks that are systematically preferred or avoided, after accounting for the overall frequency of representation of each regimen in the network.

Quantitative analyses of mortality were limited to trials that compared different types of chemotherapy. In the multipletreatments meta-analyses, we also performed subgroup analyses for first- and subsequent-line treatment.

We conducted a series of "head-to-head" meta-analyses with a random effects model. Between-study heterogeneity was estimated using the  $I^2$  statistic; typically, values greater than 50% are considered large, 25%–50% modest, and less than 25% low heterogeneity. These estimates can have large uncertainty, especially in the presence of few trials, and should be interpreted cautiously (14).

Multiple-treatments meta-analysis synthesizes information from a network of trials (7,10). It combines direct and indirect evidence on the relative effectiveness of two interventions A and B, respecting randomization. Direct evidence comes from trials of A vs B. Indirect evidence, through an "intermediate" comparator C, comes by combining trials of A vs C and of C vs B; many intermediate comparators are possible. Combination of the many sources of evidence increases precision. Combination of direct and indirect evidence for any given treatment comparison can be extended over a complex network of multiple treatments. We assumed a common heterogeneity parameter  $\tau^2$  across all comparisons. The model applied to analyze the data is a Bayesian consistency model as described in reference 7. From each trial (denoted with *i*) with, say, three randomized chemotherapy arms (say A, B, and C), we extract two hazard ratios (in log scale) with their associated variances. We then model the observed log(HR<sub>AB,*i*</sub>), log(HR<sub>AC,*i*</sub>), which compare the effectiveness of A to B and C, respectively, as:

$$\begin{pmatrix} \log(\mathrm{HR}_{\mathrm{AB},i}) \\ \log(\mathrm{HR}_{\mathrm{AC},i}) \end{pmatrix} \sim N \begin{pmatrix} \theta_{\mathrm{AB},i} \\ \theta_{\mathrm{AC},i} \end{pmatrix}, \begin{pmatrix} v_{\mathrm{AB},i} & c \\ c & v_{\mathrm{AC},i} \end{pmatrix} \end{pmatrix},$$

where  $\nu$  denotes the sample variance and *c* is the sample covariance between the log hazard ratios. The random effects  $\theta_{AB,i}$ ,  $\theta_{AC,i}$  are subsequently combined across all studies that address the same comparison as:

$$\begin{pmatrix} \theta_{\mathrm{AB},i} \\ \theta_{\mathrm{AC},i} \end{pmatrix} \sim N\left( \begin{pmatrix} \mu_{\mathrm{AB}} \\ \mu_{\mathrm{AC}} \end{pmatrix}, \begin{pmatrix} 1 & 0.5 \\ 0.5 & 1 \end{pmatrix} \tau^2 \right),$$

where  $\mu$  is the summary effect of a specific comparison and  $\tau^2$  is the between-study variance. In the simpler case, where a study has two arms only, the distributions above reduce to one dimension. The idea described above extends to all reported regimen comparisons. In the Bayesian framework, effect sizes are estimated along with 95% credibility intervals.

Multiple-treatments meta-analysis assumes that the different sources of evidence are coherent (8,15-17). On the top of the estimation of heterogeneity in each head-to-head comparison, we also estimate the incoherence (disagreement between direct and indirect evidence) in each closed loop in the network. Except for one loop with modest incoherence, no incoherence was noted.

Analyses were conducted in WinBUGS 1.4 (MRC Biostatistics Unit, Cambridge, UK; http://www.mrc-bsu.cam.ac.uk/bugs/winbugs/ contents.shtml) and S-PLUS 8 (Insightful Corporation, Seattle, WA; http://www.insightful.com).

## Results

## **Eligible Trials**

The 13651 scrutinized reports yielded 367 eligible trials (Figure 1). Another three eligible trials were identified from recent major meetings, bringing the total to 370 trials with 54189 randomly assigned patients (Table 1). Trials had been published for a period of 35 years (1973–2007), and most included only two arms. Two-thirds addressed first-line treatment, and patients with poor performance status were either absent or a small minority in most trials. Most trials had been performed in the United States or Europe, with approximately a quarter involving several countries. Details on the regimens used over time appear in Supplementary Table 1 (available online,) and detailed study design information on all the 370 trials is available from the authors upon request.

Of the 370 trials, 32 presented formally statistically significant differences between arms in the original report (Supplementary Table 2, available online). Of 198 trials comparing regimens that were categorized as the same type of treatment in our protocol,

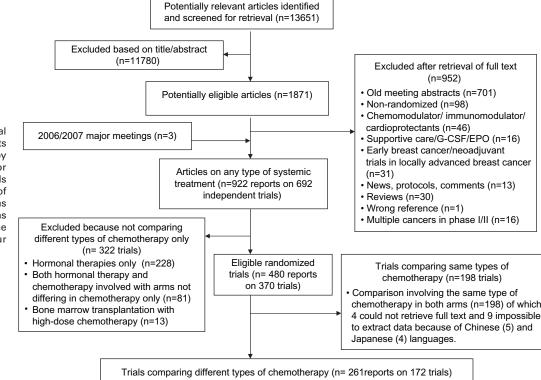


Figure 1. Flow chart for trial selection for multiple-treatments meta-analysis of chemotherapy and targeted therapies for advanced breast cancer. Trials comparing the same types of chemotherapy were defined as those that compared regimens that were considered the same type of treatment in our protocol.

only 8 (4%) presented formally statistically significant results. The other 172 trials with 31552 patients compared different types of treatment (Figure 1; Table 1).

## **Comparisons of Different Types of Treatment**

Of the 172 trials, survival data could be analyzed in 128 trials, representing 26031 randomly assigned patients, of whom 25850 were analyzed in the trials (Supplementary Table 3, available online) (18–144). Data from 121 of these trials could be obtained from published reports, and data from the other seven were obtained by contacting investigators (37,64,75,76,98,114,127). The 44 trials without analyzable survival information included 5521 patients and were mostly small phase 2 trials that did not collect or present survival data. The meta-analysis included 148 comparisons (18 trials had more than two eligible arms).

Twenty-two different types of treatment were represented in the 148 comparisons. The geometry of the network of comparisons was complex (Figure 2). Older regimens generally had somewhat more data than the more recently introduced regimens (Figure 2, A). The PIE index for the network is 0.88, and the PIE' is 0.96. Moreover, although regimens involving older agents had been thoroughly compared against each other, most regimens involving novel agents, taxanes, and/or targeted treatments (the regimens introduced more recently and thus shown with larger circles in Figure 2, B) had been compared against one or few other comparator regimens. The co-occurrence index *C* was 132 (*P* < 0.001), which suggests that there is a clear preference for specific comparisons (or avoidance for others), aside from what would be expected according to the relative representation of each regimen in the network.

## **Direct Comparisons**

For 22 different types of chemotherapy, 231 different comparisons are theoretically possible, but only 45 of these (20%) had been performed (Table 2). Only 10 of these comparisons had been made in more than three trials; thus, the results of these analyses should be interpreted very cautiously. Formally statistically significant differences were seen in 10 analyses, and an estimate consistent with large heterogeneity ( $I^2 > 50\%$ ) was seen in five comparisons; uncertainty in the heterogeneity estimates is unavoidably large with few trials per analysis.

In particular, old agents as monotherapy were formally statistically inferior to novel non-taxane agents, low-dose anthracyclinebased combinations, standard-dose anthracycline, and combinations of the same old agents. Formally statistically significant improvement in survival with the use of combined agents vs single agents was found in six comparisons; conversely, monotherapy with novel non-taxane agents fared better than combinations of old agents. The combination of old agents was statistically significantly inferior to standard-dose anthracycline-based combinations as well but was better than low-dose anthracycline monotherapy. Survival benefits were observed when taxanes were combined with novel non-taxane agents or trastuzumab (Table 2).

#### **Multiple-Treatments Meta-analysis**

Data derived from direct and indirect evidence were analyzed in the multiple-treatment meta-analysis. Results are expressed for convenience as hazard ratio relative to the regimen of monotherapy with an old agent (Table 3). Taxanes in combination with novel agents, trastuzumab, or old agents, had the largest decrease in mortality risk (hazard ratios [HRs] 0.49, 0.50, and 0.53, respectively). Considerable survival benefits, relative to monotherapy

Table 1. Characteristics of the eligible trials
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Characteristics	All trials (n = 370)	Trials eligible for MTM (n = 172)
No. of eligible patients	54 189	31 552
Median sample size (IQR)	104 (53–201)	141 (87–262)
Year of publication		
1971–1980	51	28
1981–1990	111	53
1991–2000	118	51
2001–2007	90	40
Number of eligible armst		
Two	309	136
Three	44	29
Four or more	16	7
Previous chemotherapy, No. (%)	140 (40)‡	55 (32)
Including PS >2, No. (%)	105 (37)‡	62 (46)‡
Had more than 10% of patients	24 (8)‡	9 (7)‡
with PS >2, No. (%)		
Countries involved (investigator		
affiliations)		
Multiple countries	95	52
United States	85	41
Italy	41	16
United Kingdom	28	14
France	22	7
Japan	14	1
China	10	0
Germany	9	4
Denmark	9	6
Other	57	31

 MTM = multiple-treatments meta-analysis; IQR = interquartile range; PS = performance status.

Fifteen trials (of which 12 were eligible for multiple-treatments meta-analysis) also included additional arms that would not be eligible for the systematic review.

Data are not available for all trials, percentages calculated based on trials with available information.

with an old agent, were observed for taxanes in combination with lapatinib, anthracyclines, or both anthracycline + novel non-taxane agents (HRs = 0.57, 0.64, and 0.56, respectively). Combination of trastuzumab with standard-dose anthracyclines offered similar benefits (HR = 0.55). Taxanes as single agents and standard-dose anthracyline-based combinations reduced mortality risk by approximately one-third (HR = 0.67). Old combination regimens without anthracyclines or taxanes achieved a 25% relative risk reduction in mortality.

Therapy line–specific effect estimates were similar to estimates for all available comparisons regardless of line. Effect estimates for the overall analysis and first-line trials only typically differed by less than 5%. Subsequent-line treatment data were more sparse and thus less conclusive, but the overall pattern did not differ, with the exception of a trend for lower efficacy for anthracycline-based combinations when compared with the firstline setting (Table 3).

When plotting hazard ratios as a function of the year of publication of the first trial that used each type of treatment (Figure 3), it appeared that early regimens such as combination of old agents and anthracycline-based regimens achieved a 25%–30% relative risk reduction, with little further progress for two decades; the

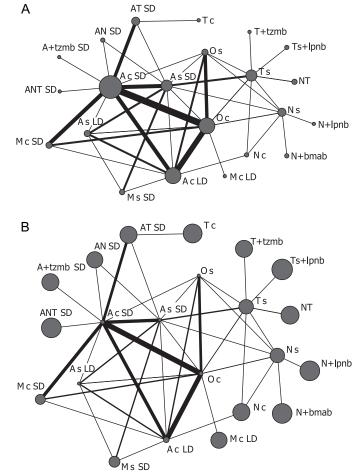


Figure 2. Network of eligible comparisons for the multiple-treatments meta-analysis. Thickness of connecting lines indicates the number of available comparisons. A) The size of each node is proportional to the amount of information (sample size). B) A larger node means more recent year of first publication of a trial for this type of regimen. Ac LD = low-dose anthracycline (combination regimen); Ac SD = standard-dose anthracycline (combination regimen); As LD = low-dose anthracycline (single agent); As SD = standard-dose anthracycline (single agent); A+tzmb SD = standard-dose anthracycline + trastuzumab; AN SD = standard-dose anthracycline + novel non-taxane agents; ANT SD = standard-dose anthracycline + novel non-taxane agents + taxanes; AT SD = standard-dose anthracycline + taxanes; Mc LD = low-dose mitoxantrone (combination regimen); Mc SD = standard-dose mitoxantrone (combination regimen); Ms SD = standard-dose mitoxantrone (single agent); N c = novel non-taxane agents (combination regimen); N s = novel non-taxane agents (single agent); N+bmab = novel non-taxane agents + bevacizumab (single agent); N+lpnb = novel non-taxane agents + lapatinib; NT = novel non-taxane agents + taxanes; Oc = old agents (combination regimen); Os = old agents (single agent); Tc = taxanes (combination regimen); Ts = taxanes (single agent); T+tzmb = taxanes + trastuzumab; Ts+lpnb = taxanes + lapatinib.

introduction of additional agents attained larger (30%–50%) relative risk reductions.

## Discussion

Our meta-analysis quantifies the progress achieved in the treatment of advanced breast cancer with nonhormonal systemic treatment in the last 35 years. Several regimens have shown effectiveness, and for some of them, the treatment effects are practically indistinguishable in magnitude. Given that subsequent lines of treatment

Arm 1	Arm 2	No. of studies	No. of patients	HR† (95% CI)	I² (%)
Low-dose anthracycline (combination regimen)	Standard-dose anthracycline (combination regimen)	16	3347	1.05 (0.98 to 1.13)	0
Low-dose anthracycline (combination regimen)	Low-dose anthracycline (single agent)	4	725	0.93 (0.83 to 1.05)	0
Low-dose anthracycline (combination regimen)	Standard-dose anthracycline (single agent)	1	115	0.90 (0.62 to 1.32)	NA
Low-dose anthracycline (combination regimen)	Standard-dose mitoxantrone (combination regimen)	2	440	0.79 (0.58 to 1.07)	51
Low-dose anthracycline (combination regimen)	Standard-dose mitoxantrone (single agent)	1	260	0.89 (0.67 to 1.18)	NA
Low-dose anthracycline (combination regimen)	Novel non-taxane agents (combination regimen)	1	84	1.08 (0.67 to 1.75)	NA
Low-dose anthracycline (combination regimen)	Old agents (combination regimen)	19	2257	0.96 (0.86 to 1.08)	23
Low-dose anthracycline (combination regimen)	Old agents (single agent)	3	458	0.82‡ (0.68 to 1.00)	0
Standard-dose anthracycline (combination regimen)	Low-dose anthracycline (single agent)	2	565	0.87 (0.73 to 1.04)	0
Standard-dose anthracycline (combination regimen)	Standard-dose anthracycline (single agent)	8	1030	0.84‡ (0.74 to 0.96)	0
Standard-dose anthracycline (combination regimen)	Standard-dose anthracycline + trastuzumab	1	281	1.22 (0.91 to 1.63)	NA
Standard-dose anthracycline (combination regimen)	Standard-dose anthracycline + novel non-taxane agents	1	170	1.03 (0.73 to 1.45)	NA
Standard-dose anthracycline (combination regimen)	Standard-dose anthracycline + novel non-taxane agents + taxanes	1	243	1.18 (0.80 to 1.77)	NA
Standard-dose anthracycline (combination regimen)	Standard-dose mitoxantrone (combination regimen)	9	1322	0.95 (0.84 to 1.08)	9
Standard-dose anthracycline (combination regimen)	Old agents (combination regimen)	14	2529	0.89‡ (0.80 to 0.98)	11
Standard-dose anthracycline (combination regimen)	Old agents (single agent)	1	60	1.14 (0.60 to 2.18)	NA
Standard-dose anthracycline (combination regimen)	Standard-dose anthracycline + taxanes	9	2201	1.06 (0.89 to 1.27)	62
Low-dose anthracycline (single agent)	Standard-dose anthracycline (single agent)	5	202	1.03(0.70 to 1.54)	63
Low-dose anthracycline (single agent)	Standard-dose mitoxantrone (single agent)	2	99	0.77(0.50 to 1.17)	0
Low-dose anthracycline (single agent)	Old agents (combination regimen)	2	88	1.54‡(1.01 to 2.32)	0
Standard-dose anthracycline (single agent)	Standard-dose anthracycline + novel non-taxane agents	2	687	1.09 (0.93 to 1.29)	0.
Standard-dose anthracycline (single agent)	Standard-dose anthracycline + taxanes	1	454	0.98 (0.79 to 1.22)	NA
Standard-dose anthracycline (single agent)	Standard-dose mitoxantrone (single agent)	3	621	0.87 (0.65 to 1.18)	61
Standard-dose anthracycline (single agent)	Novel non-taxane agents (single agent)	2	698	0.75 (0.46 to 1.22)	84
Standard-dose anthracycline (single agent)	Old agents (combination regimen)	1	64	1.26 (0.71 to 2.25)	NA
Standard-dose anthracycline (single agent)	Old agents (single agent)	2	155	0.52‡ (0.35 to 0.77)	10
Standard-dose anthracycline (single agent)	Taxanes (single agent)	3	1110	1.02 (0.89 to 1.17)	0
Standard-dose anthracycline + taxanes	Taxanes (combination regimen)	1	327	1.20 (0.88 to 1.64)	NA
Standard-dose anthracycline + taxanes	Taxanes (single agent)	1	459	1.07 (0.88 to 1.30)	NA
Low-dose mitoxantrone (combination regimen)	Old agents (combination regimen)	1	312	0.92 (0.61 to 1.41)	NA
Standard-dose mitoxantrone (combination regimen)	Novel non-taxane agents (single agent)	1	65	1.06 (0.65 to 1.72)	NA
Standard-dose mitoxantrone (combination regimen)	Old agents (combination regimen)	3	116	0.88 0.68 to 1.14)	0
Novel non-taxane agents (combination regimen)	Novel non-taxane agents (single agent)	1	251	1.04 (0.78 to 1.38)	NA
Novel non-taxane agents (combination regimen)	Taxanes (single agent)	1	176	1.07 (0.71 to 1.59)	NA
Novel non-taxane agents (single agent)	Novel non-taxane agents + bevacizumab	1	462	1.04 (0.77 to 1.40)	NA
Novel non-taxane agents (single agent)	Novel non-taxane agents + lapatinib	1	324	1.09 (0.69 to 1.72)	NA
Novel non-taxane agents (single agent)	Old agents (combination regimen)	3	482	0.78‡ (0.62 to 0.97)	0
Novel non-taxane agents (single agent)	Old agents (single agent)	1	179	0.67‡ (0.47 to 0.97)	NA
Novel non-taxane agents (single agent)	Taxanes (single agent)	1	41	1.56 (0.69 to 3.53)	NA
Novel non-taxane agents + taxanes	Taxanes (single agent)	3	1140	0.76‡ (0.66 to 0.88)	0
Old agents (combination regimen)	Old agents (single agent)	9	950	0.72‡ (0.55 to 0.95)	72
Old agents (combination regimen)	Taxanes (single agent)	4	1076	1.10 (0.82 to 1.47)	76
Old agents (single agent)	Taxanes (single agent)	1	81	1.72 (0.82 to 3.61)	NA
Taxanes (single agent)	Taxanes + trastuzumab	2	374	1.32‡ (1.06 to 1.65)	0
Taxanes (single agent)	Taxanes + lapatinib	1	579	1.16 (0.93 to 1.46)	NĂ

Forest plots are available in Supplementary Figure 1 (available online). NA = not applicable (one trial); HR = hazard ratio; CI = confidence interval

HR estimates less than 1.00 suggest that arm 1 has better survival. t

‡ *P* < .05.

can confer similar relative benefits as the first-line setting, one can exploit the survival benefits conferred by several effective regimens used in sequential fashion. For example, if three regimens are used, each one prolonging survival by 1 year for a patient who would have had 1 year of survival without effective treatment, life expectancy may be quadrupled. The availability of multiple effective options suggests that it is essential to identify appropriate indications for timely changes of regimens in patients in whom treatment is no longer effective.

Progress in chemotherapeutic treatment of advanced breast cancer has been stepwise. After the achievement of a 25% reduction in the risk of death in 1975 with the use of anthracyclinebased combination regimens (145), little additional progress was made for almost two decades. Subsequently, several new agents (vinorelbine, gemcitabine, capecitabine, taxanes, and molecular targeted treatments) were introduced that proved more effective (59,85,146-148). These agents have not yet been tested in all possible combinations. Moreover, less data have been collected on combinations involving these newer agents than are available for the standard earlier treatments. Comparisons for which there is at present little or no data should be the focus of future investigations so that more direct evidence about the relative merits of these regimens can be obtained.

Interventions should be tailored to the individual patient, and benefits should be weighed against adverse events for each treatment. Toxicity should not be underestimated. For example,

Table 3. Multiple-treatments r	meta-analysis and subgro	oup analyses by line of treatment
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Treatment	All available comparisons in any line (n = 148), HR (95% Crl)	Survival gain over Os* (mo)	First-line comparisons (n = 107), median HR (95% Crl)	Subsequent-line comparisons (n = 41), median HR (95% Crl)
Old agents (single agent)	1.00 (referent)		1.00 (referent)	1.00 (referent)
Novel non-taxane agents + taxanes	0.49 (0.37 to 0.67)	12.5	0.52 (0.32 to 0.86)	0.47 (0.30 to 0.71)
Taxanes + trastuzumab	0.51 (0.35 to 0.72)	11.5	0.51 (0.33 to 0.79)	0.56 (0.35 to 0.90)
Taxanes (combination regimen)	0.53 (0.34 to 0.85)	10.6	0.54 (0.32 to 0.90)	_
Standard-dose anthracycline + trastuzumab	0.55 (0.36 to 0.84)	9.8	0.54 (0.34 to 0.88)	_
Standard-dose anthracycline + novel non-taxane agents + taxanes	0.56 (0.34 to 0.94)	9.4	0.56 (0.32 to 0.98)	—
Taxanes + lapatinib	0.57 (0.38 to 0.86)	9.1		_
Standard-dose anthracycline + taxanes	0.64 (0.53 to 0.78)	6.8	0.64 (0.50 to 0.83)	_
Standard-dose anthracycline + novel non-taxane agents	0.65 (0.49 to 0.85)	6.5	0.67 (0.46 to 0.97)	0.64 (0.41 to 1.03)
Taxanes (single agent)	0.67 (0.55 to 0.81)	5.9	0.67 (0.50 to 0.91)	0.65 (0.49 to 0.87)
Standard-dose anthracycline (combination regimen	0.67 (0.57 to 0.78)	5.9	0.67 (0.54 to 0.81)	0.74 (0.55 to 0.99)
Novel non-taxane agents + lapatinib	0.68 (0.38 to 1.23)	5.6	0.69 (0.39 to 1.22)	0.62 (0.33 to 1.12)
Low-dose mitoxantrone (combination regimen)	0.69 (0.41 to 1.18)	5.4	_	_
Low-dose anthracycline (combination regimen)	0.70 (0.61 to 0.81)	5.1	0.70 (0.57 to 0.85)	0.76 (0.57 to 1.01)
Standard-dose anthracycline (single agent)	0.71 (0.60 to 0.84)	4.9	0.74 (0.58 to 0.94)	0.72 (0.55 to 0.94)
Novel non-taxane agents + bevacizumab (single agent)	0.71 (0.45 to 1.12)	4.9	_	0.64 (0.39 to 1.06)
Novel non-taxane agents (combination regimen)	0.72 (0.53 to 0.99)	4.7	0.65 (0.35 to 1.18)	0.69 (0.46 to 1.04)
Novel non-taxane agents (single agent)	0.74 (0.60 to 0.91)	4.2	1.08 (0.69 to 1.68)	0.66 (0.50 to 0.88)
Old agents (combination regimen)	0.75 (0.65 to 0.85)	4.0	0.75 (0.63 to 0.90)	0.72 (0.54 to 0.95)
Standard-dose mitoxantrone (combination regimen)	0.75 (0.62 to 0.90)	4.0	0.74 (0.59 to 0.94)	0.71 (0.41 to 1.26)
Low-dose anthracycline (single agent)	0.78 (0.64 to 0.94)	3.4	0.80 (0.62 to 1.03)	0.79 (0.54 to 1.18)
Standard-dose mitoxantrone (single agent)	0.82 (0.64 to 1.06)	2.6	0.81 (0.56 to 1.18)	0.85 (0.58 to 1.26)

\* Absolute prolongation of survival with various regimens for a patient with an anticipated survival on an old agent (single agent) of 1 y; it is calculated as [(12/HR) - 12] mo. Crl = credibility interval; HR = hazard ratio; Os = old agents (single agent).

taxane-based combinations that include capecitabine and gemcitabine cause serious hematologic toxicity, diarrhea, mucositis, hand-foot syndrome, neurosensory disorders, and fatigue (39,125,128). Administration of anthracycline with trastuzumab was associated with major cardiotoxicity (84), and preliminary data (149) suggest that the combination of gefitinib with docetaxel is very toxic. Very few patients with poor performance status, who may be especially susceptible to adverse effects, have been enrolled in recent trials.

Our meta-analysis includes data on results of treatment with trastuzumab, bevacizumab, and lapatinib as of October 2007. The available data on lapatinib and bevacizumab in particular are quite limited. After data freeze and analysis, a trial on 722 patients (150) was published, showing no statistically significant survival benefit with bevacizumab + paclitaxel vs paclitaxel, but clear benefit was seen for disease progression. It is possible that many patients switched treatment upon disease progression and that this may have eroded any survival difference. With many available treatment options and early switches to other effective agents, differences in survival by intention to treat may become small. Cautious optimism as to the effectiveness of molecular targeted treatments is needed until more data accumulate, and it is important to correctly identify the patients who would benefit from them. Targeted therapies are not appropriate for all breast cancer patients. Trastuzumab and lapatinib are effective in treating HER2-positive cancers, but these account only for approximately 20% of breast cancers (129). Although molecular targeted therapies have an increasing range of applications for different cancers, trials with less impressive results

(not stopped early) may still be ongoing, and the complete picture of effectiveness and toxicity has not yet emerged.

Overall, it would be useful to design trials that fill in important gaps for key treatment comparisons that have minimal or no information. For some treatment comparisons in the examined network of treatments, no direct evidence was available, and thus, evaluation of incoherence (ie, the extent of disagreement between direct and indirect evidence) was also impossible.

Furthermore, it has to be considered that over time, anthracycline- and taxane-based chemotherapy has been increasingly used for the treatment of early-stage disease. Thus, estimates of the relative merits of these agents in the advanced setting could be in part biased by when the study was conducted.

Also in the last three decades, there has been much change in supportive care of cancer patients. Thus, there is uncertainty as to the validity of the transitivity assumption (ie, that the treatment effect of a regimen does not change over time) for early introduced regimens. Would the relative risk reduction be the same as several decades ago, if early trials were to be repeated? Obviously, repetition of the trials is not feasible for ethical reasons. Changes in supportive care and other adjunct management would certainly change the absolute survival for both compared arms, but there is no strong reason to believe that these changes would influence the relative performance of the compared regimens, and it is the relative performance of one regimen vs the other that enters in the multiple-treatments meta-analysis calculations.

A possible limitation of the current analysis is that it is based on published group data, rather than individual patient information.

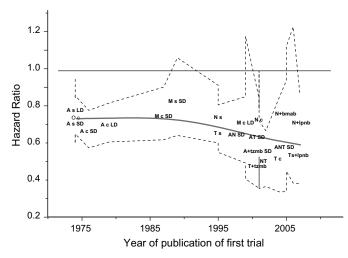


Figure 3. Hazard ratios for mortality using monotherapy, with old agent as the reference comparator as a function of the first year of publication of a trial for each type of treatment. The abbreviated name of each regimen is placed at the point representing the first year of publication, and the hazard ratio estimated from the multipletreatments meta-analysis. Also shown is a spline fitted line. In dashed lines are the piecewise 95% credibility intervals. Ac LD = low-dose anthracycline (combination regimen); Ac SD = standard-dose anthracycline (combination regimen); As LD = Low-dose anthracycline (single agent); As SD = standard-dose anthracycline (single agent); A+tzmbSD = standard-dose anthracycline + trastuzumab; AN SD = standard-dose anthracycline + novel non-taxane agents; ANT SD = standard-dose anthracycline + novel non-taxane agents + taxanes; AT SD = standard-dose anthracycline + taxanes; Mc LD = low-dose mitoxantrone (combination regimen); Mc SD = standard-dose mitoxantrone (combination regimen); Ms SD = standard-dose mitoxantrone (single agent); Nc = novel non-taxane agents (combination regimen); Ns = novel non-taxane agents (single agent); N+bmab = novel non-taxane agents + bevacizumab (single agent); N+lpnb = novel non-taxane agents + lapatinib; NT = novel non-taxane agents + taxanes; Oc = old agents (combination regimen); Tc = taxanes (combination regimen); Ts = taxanes (single agent); T+tzmb = taxanes + trastuzumab; Ts+lpnb = taxanes + lapatinib.

However, we have included patients across a rather narrow range of prognoses, namely, metastatic breast cancer (stage IV). Individual patient data might allow a more detailed appraisal of outcomes at different risk groups (151). Still, the power to detect effect modifications might still be limited even with individual patient information in such a complex network of multiple treatments. The available data for each regimen are relatively limited, and thus, strong inferences about the specific superiority of one particular regimen should be avoided. Finally, given the retrospective nature of the meta-analysis, publication bias and selective reporting biases cannot be excluded. Although such biases may affect sporadic comparisons, they are unlikely to refute the overall picture of progress. Prospective registration of trials has been adopted in the last few years. Registration, if complete and transparent, would allow updating this and similar analyses with more comprehensive information and keeping track of the improvements in the treatment of advanced breast cancer.

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