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Epidemiology, biology, and treatment of triple-negative breast cancer in women of African ancestry

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Abstract

Breast cancer incidence is increasing worldwide, and breast cancer-related mortality is highest in women of African ancestry, who are more likely to have basal-like or triple-negative breast cancer (TNBC) than are women of European ancestry. Identification of cultural, epidemiological, and genetic risk factors that predispose women of African ancestry to TNBC is an active area of research. Despite the aggressive behaviour of TNBC, achievement of a pathological complete response with chemotherapy is associated with good long-term survival outcomes, and sensitivity to chemotherapy does not seem to differ according to ethnic origin. Discovery of the molecular signalling molecules that define TNBC heterogeneity has led to the development of targeted agents such as inhibitors of poly (ADP-ribose) polymerase-1 and mTOR and immunomodulatory drugs that are in the early stages of clinical testing. First, we summarise the existing published work on the differences reported on the epidemiology, biology, and response to systemic treatment of TNBC between women of African ancestry and white women, and identify some gaps in knowledge. Second, we review the opportunities for development of new therapeutic agents in view of the potential high clinical relevance for patients with TNBC irrespective of race or ethnic origin.

Introduction

Breast cancer is the most common cancer in women worldwide, with around 1 676 633 new cases diagnosed in 2012 (figure 1).¹ Breast cancer has become the most prevalent cancer in women in sub-Saharan Africa, and in most of the 22 countries of the Americas, including the Caribbean.^{2,3}

The annual incidence of breast cancer varies in women of African ancestry and accurate reporting of the data is affected by the inadequacy of cancer registries. The age-adjusted

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Declaration of interests

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incidence ranges from 30 cases per 100 000 women in eastern Africa¹ to 78·1 cases per 100 000 in a Caribbean population⁴ and 120·5 cases per 100 000 in African-American women.⁵ Since the publication of GLOBOCAN 2008 cancer statistics, breast cancer incidence has increased by more than 20% worldwide.¹ In Africa, breast cancer incidence increased annually in Hare, Zimbabwe, by around 4·9% and in Kampala, Uganda, by about 4·5% between the early 1990s and mid-2000s.^{6,7} In addition to population growth and ageing, increases in breast cancer incidence in Africa, particularly in wealthy women, might be attributable to women having fewer children and giving birth for the first time later in their lives than in previous decades⁸ partly as a result of an increase in use of oral contraception, and to increases in prevalence of lifestyle risk factors, such as obesity and low levels of physical activity.^{9,10}

Breast cancer is the main cause of cancer-related mortality in women worldwide; 324 000 deaths were reported in low-income countries in 2012,¹ with 68 100 occurring in Africa.³ Breast cancer-related mortality in African-American women (30.8 deaths per 100 000)¹¹ is quite similar by comparison with the age-standardised cancer-related mortality of women living in the non-Latin Caribbean who are predominantly of African ancestry (25.0 deaths per 100 000), despite decreased health-care capacity.²

Racial disparities in breast cancer-related mortality have also been noted in high-income countries with mixed racial populations. In the USA, African-American women have a 41% higher breast cancer-related death rate than do white women.¹¹ In the UK, 5-year distant breast cancer relapse-free survival is 62.8% for young black women, compared with 77% for young white women with equal access to health care (p=0.0053).¹² Racial disparities in breast cancer survival between and within countries are linked to the availability of early detection, access to diagnosis and treatment, cultural differences in lifestyle behaviours, socioeconomic factors, and differences in the biological characteristics of breast cancer.¹³

Subtypes of breast cancer

Breast cancer has traditionally been subtyped into oestrogen receptor (ER)-positive (60– 70% of breast cancer cases) and ER-negative breast cancer (30–40% of cases).¹⁴ ERpositive breast cancer typically expresses the progesterone receptor (PR), another molecular marker of breast cancer. ER-positive or PR-positive breast cancer typically responds to ERtargeted therapy, such as selective ER modulators (eg, tamoxifen) or aromatase inhibitors, which lower the serum concentrations of oestrogen. ER-negative breast cancer is aggressive and is most common in women of African ancestry and women who carry *BRCA1* mutations.^{15,16}

15–20% of breast cancers amplify or overexpress the oncogene *HER2/neu* oncogene.¹⁴ When discovered, HER2-positive breast cancer—which can be either ER-positive or ER-negative—was associated with a poor prognosis.¹⁴ However, the advent of HER2-targeted treatments (eg, trastuzumab) have greatly improved the outcomes of women with HER2-positive breast cancer.¹⁴

These three molecular markers distinguish breast cancer into subgroups of ER-positive, PR-positive, or HER2- positive breast cancer. Breast cancer that does not express any of the

three markers is commonly termed triple-negative breast cancer (TNBC). The recommended definition of ER-negative and PR-negative breast cancer are based on tests that distinguish immunoreactive cells; ER-negative and PR-negative breast cancers are those that express less than 1% immunoreactive cells, based on studies that have shown assocations of tumour response and clinical outcomes to endocrine therapy.¹⁷ However, many clinical trials and epidemiological studies often use a threshold of more than 10% immunoreactive cells to define receptor negativity, which was the previously defined threshold assumed to correspond with the cutoff point of cytosol protein 10 fmol per mg for the ligand-binding assay, the first test used to define receptor presence or absence based on the odds of response to endocrine treatment. TNBC accounts for about 10–15% of all breast cancers but it is much more common in women of African ancestry than any other ethnic origns, particularly in women who develop breast cancer before the age of 50 years.¹⁵

More recently, gene-expression profiling has also been used to subtype breast cancer, especially TNBC. Initial investigations done by Perou and colleagues¹⁸ and Sorlie and colleagues¹⁹ showed that breast cancer could be separated into five groups using RNA expression profiling: luminal A (ER-positive), luminal B (ER-positive), HER2 (HER2-positive), basal-like (mostly ER-negative), and normal-like breast cancer. Subsequently, Prat and colleagues²⁰ showed that basal-like breast cancer included another subtype, termed claudin-low breast cancer. Basal-like and claudin-low breast cancer are both predominantly (but not exclusively) ER-negative and triple-negative.

TNBC subgroups, incidence, and prevelence

TNBC and basal-like breast cancer are more common in women of African ancestry than women of other ethnic origins.^{15,21} TNBC is often aggressive, and is associated with a higher mortality than the other subtypes of breast cancer (p<0.01).¹⁹ TNBC is characterised by high cell proliferation, poor cellular differentiation, many recurrent copy number imbalances, and, in most cases, mutations in the *TP53* tumour suppressor gene.^{18,22,23}

TNBC has been further subdivided by use of meta-analysis of gene-expression profiling data. Lehmann and colleagues²⁴ analysed RNA expression profiling data from 14 breast cancer gene expression datasets that used breast tumours from women in the USA, Europe, and China as the training set to develop gene signatures for TNBC subgroups. They then validated these gene signatures by use of RNA profiling data from seven additional datasets of tumours from the USA and Europe. No breakdown of race or ethnic origin was provided in this analysis; however, due to the different ethnic population distributions of the countries, women of African ancestry were likely to be included in the US datasets but not in those from Europe or China. Their analysis shows that TNBC can be subdivided into six subgroups: basal-like 1, basal-like 2, immunomodulatory, mesenchymal-like, mesenchymal stem-like, and luminal androgen-receptor (AR) expressing. Lehmann and colleagues²⁴ also showed that the breast cancer subtypes have varying sensitivity to therapeutic agents. Basallike 1 and basal-like 2 cells were most sensitive to cisplatin, luminal AR cells to bicalutamide (an anti-androgen) and alvespimycin (an Hsp90 inhibitor), and the mesenchymal subgroups to the SRC inhibitor dasatinib and the phosphoinositol-3 kinase (PI3K)/mTOR inhibitor NVP-BEZ235. These results suggest that different forms of TNBC

Population-based incidence for TNBC in women of African ancestry have mainly been reported from the USA's Surveillance, Epidemiology, and End Results (SEER) California Cancer Registry, which has collected data on the status of ER and PR subtypes since 1990 and on HER2/neu status since 1999.25 By use of the SEER California database, Clarke and colleagues²⁵ showed that incidence of TNBC is higher in African-American women than other racial or ethnic origin groups at all ages (p<0.05).²⁵ African-American women were twice as likely to be diagnosed with ER-positive, PR-positive, and HER2-negative breast cancer than TNBC; however, the ratio was substantially lower than reported for white (ratio 6.9:1) or Asian women (ratio 6.1:1). An analysis of the entire US SEER data of women diagnosed with breast cancer in 2010, added evidence to support this result, showing that African-American (odds ratio [OR] 1.4, 95% CI 1.2–1.6) and Hispanic women (1.3, 1.2– 1.5) were more likely to be diagnosed with TNBC than were white women.²⁶ Since mammography screening is more likely to detect ER-positive breast cancer,²⁷ differences in screening patterns by race might account for the under-representation of ER-positive breast cancer in African- American women than in women of other ethnic origins. Data for population-based rates of mammography screening by race and ethnic origin are conflicting. Data from the 2010 National Health Interview Survey of self-reported mammography screening showed that black women were as likely as white women to report having had a mammogram in the past 2 years (73.2% and 72.8%, respectively), and 71.9% of all women aged 65-74 years reported mammography screening.²⁸ Silber and colleagues²⁹ used information from Medicare and the SEER database to show that in women diagnosed with breast cancer after the age of 66 years, black women were significantly less likely than white women (23.5% vs 35.7%, p<0.001) to have received any breast cancer screening in the 6–18 months before diagnosis. However, in women younger than 40 years in the USA, who are not eligible for population-based mammography screening, African-American women had 000) or Hispanic women $(2.1 \text{ per } 100\ 000 \text{ women})$. This result suggests that differential mammography screening rates and oversampling of ER-positive breast cancer might not fully explain racial and ethnic differences in the incidence of TNBC.³⁰

Although the frequency of TNBC varies across regional populations of women of African ancestry, it is consistently higher than that reported in other racial or ethnic origin groups. In the population-based North Carolina Breast Cancer cohort of 878 African-American women with breast cancer, premenopausal African-American women had higher rates of basal breast cancer (39%) than did white women of a similar age (16%) or postmenopausal African-American women (14%).²¹ Similarly, Bowen and colleagues²⁹ showed in a retrospective cohort study of patients with breast cancer in the UK that 22% of black women had TNBC compared with 15% of white women.³¹

Huo and colleagues³² investigated the distribution of molecular subtypes of invasive tumours in women (mean age 44.8 years) in different geographical regions in Nigeria and Senegal (507 women) and reported that TNBC, including basal-like TNBC, was the predominant type of cancer (27%).³² Of consecutive cases of breast cancer reported in a

Bamako University hospital in Mali, the mean age of patients was 46 years (range 25–85 years) and 46% of tumours were triple-negative.³³ In a large case study³⁴ of 1216 women with breast cancer in Soweto, South Africa, 90% were black, and 20% of breast cancers were triple-negative, which is consistent with the frequency reported in African-American women. The frequency of TNBC was highest in women aged 50–59 years (29·2%), and was higher in women of African ancestry than in women of other ethnic origins (OR 2·2, 95% CI 1·1–3·8).³⁴ Smaller, hospital-based studies have reported a higher prevalence of TNBC in a sample of Ghanaian women (79%) compared with African-American (32%) and white American women (10%).³⁵

Overall, epidemiological data support the conclusion that although TNBC is not restricted to a specific age or ethnic group, this cancer is of a higher frequency and is a contributor to the survival disadvantage of women of African ancestry with breast cancer. Factors that might account for variations in the incidence and prevalence of TNBC in women of African ancestry include differences in methods of case ascertainment, population age structure, genetic and lifestyle risk factor distribution, and access to mammography screening.^{34,35} Overestimation of ER and PR negativity is of particular concern because poor or unreliable laboratory standards for tissue handling, type of fixation used, and the initiation and duration of fixation can affect the optimum performance of immunohistochemical testing and thus the results.¹⁷

Epidemiological risk factors

Few epidemiological studies have included large numbers of women of African ancestry to examine reproductive risk factors associated with TNBC or ER-negative breast cancer by ethnic origin. Most evidence is from studies done in the USA in which at least 10% of patients with breast cancer were of African ancestry, and suggests a positive association between higher parity or having more than one child and risk of development of TNBC, and a negative association between duration of breastfeeding and risk of development of TNBC (table).

The population-based North Carolina Breast Cancer study,³⁶ which assessed 878 African-American women and 187 white women with breast cancer, showed that having both multiple children and not breastfeeding those children were risk factors for basal-like TNBC across ethnic groups. A population-based case study⁴¹ of black women from the USA and Caribbean showed that, although having more children was associated with an increased risk of TNBC (p_{trend}=0·03), the association with breastfeeding was not significant. Another large population-based, case-control study³⁹ of 873 African- American and 1072 white women with breast cancer living in California reported a significant trend between longer durations of breastfeeding and reduced risk of development of TNBC (p_{trend}=0·02), and oral contraceptive use was associated with a 2·9 times increased risk of women aged 45–64 years developing TNBC.³⁹ Development of TNBC in the absence or short duration of breastfeeding might be caused by progenitor cells in breast tissue not undergoing terminal differentiation and apoptosis that usually occurs with prolonged lactation, resulting in a persistent pool of progenitor cells that are at risk for carcinogenesis.⁴⁰

Data for the relation between obesity and TNBC are conflicting.^{37,38} A meta-analysis⁴² of case-case and case-control studies, some of which included women of African ancestry, showed that obesity was associated with an increased risk of development of TNBC for all women (OR 1·20, 95% CI 1·03–1·40). An increased hip-to-waist ratio has also been associated with an increased risk of development of TNBC and ER/PR-negative breast cancer^{36,43} in African-American women. Additionally, a suggested role for diet in the risk of development of TNBC is derived from epidemiological studies that showed a slightly significant inverse association between high total vegetable intake (relative risk 0·82, 95% CI 0·74–0·90),⁴⁴ or a diet high in fruits and vegetables (0·85, 0·76–0·95),⁴⁵ and risk of development of ER-negative breast cancer.

Few epidemiological studies have investigated risk factors for breast cancer in African women, because routine assessment of ER, PR, and Her-2/*neu* on breast tumours is not done and thus few data are available about specific risk factors for subtypes. Similar causal associations are assumed to exist for breast cancer development in African-American women to women of African ancestry living elsewhere in the world.⁴⁶ In view of the differences in the distribution of reproductive risk factors between African women and women of African ancestry in high-income countries, unknown risk factors could be associated with risk of developing TNBC in Africa. Galukande and colleagues⁴⁷ reported a small case study of 113 women with breast cancer diagnosed in Uganda, with information available about ER and PR status. The prevalence of ER/PR-negative tumours was much the same between women younger than 50 years and women older than 50 years. No significant association between parity, breastfeeding, or body-mass index and ER/PR-positive versus ER/PR-negative disease was recorded; however, the sample size of this study was potentially too small to provide adequate power to detect associations.

The role that socioeconomic status has in the cause of TNBC is of interest because low socioeconomic status is associated with many of the shared characteristics of breast tumours that occur in women of African ancestry, including high grade, high clinical stage, and ER-negative status.^{15,21} Data from the California Cancer Registry showed that irrespective of race or ethnic origin, women living in areas of low socioeconomic status were more likely than women living in areas of high socioeconomic status to be diagnosed with TNBC than any other type of breast cancer.¹⁵ Socioeconomic status is intrinsically linked with race and lifestyle behaviours, such as physical activity, obesity, diet, reproductive experiences such as having more children, and screening behaviours, which vary in prevalence across different populations of women.⁴⁸ Therefore, the challenge faced in assessment of the causes of TNBC in women of African ancestry is to distinguish the contributions of women.

Genetic susceptibility

In view of the higher incidence of TNBC in women of African ancestry than of European ancestry, African ancestry might be associated with inherited genetic variants that predispose carriers to TNBC. By use of genome-wide admixture mapping in 1484 African-American women with invasive breast cancer, assessed for around 2400 ancestry informative markers (AIMs) that distinguish European and African ancestral origins,

Fejerman and colleagues⁴⁹ reported no association between African or European ancestry, for subtype-specific breast cancer risk at any specific loci. However, global ancestry results showed that women with a higher percentage of European ancestry, estimated from AIMs, had an increased risk of having ER-positive or PR-positive breast cancer rather than ER-positive or PR-negative breast cancer (OR 2·84, 95% CI 1·13–7·14 for a 0–100% change in European ancestry), and were more likely to be diagnosed at an earlier stage of disease.⁴⁹ In another study of African-American women, Reding and colleagues⁵⁰ estimated the percentage of African ancestry with 128 AIMs. They reported no variation within ethnic groups for risks of ER or PR subtypes, stage, or grade by percentage ancestry (95% African ancestry versus <80% African ancestry). Because of the large amounts of genetic variation in self-identified African-Americans, random samples of African-Americans recruited from different regions will probably have very different allele frequencies that might lead to inconsistent results in studies assessing genetic associations with TNBC.⁵¹

Women with a *BRCA1* mutation have more than a 50% chance of development of TNBC, and unique and founder *BRCA1* mutations have been identified in patients of African ancestry with breast cancer.⁵² The estimated prevalence of *BRCA1* mutations in African-American women not selected for family history is low, at about 1.4%.^{53,54} In a cohort of 434 unselected Nigerian patients with breast cancer, 31 women (7.0%) had a *BRCA1* mutation,⁵⁵ which might suggest an increased contribution of deleterious mutations in *BRCA1* causing TNBC in Nigerian women. A study⁵⁶ of 214 Bahamian women with breast cancer who were not selected for age or family history reported that 52 (24%) carried *BRCA1* mutation (of which, most mutations were African founder mutations), and is the highest prevalence reported so far in any population. Knowledge of the prevalence of TNBC in women of African ancestry, is restricted by the low rates of genetic testing in all women of this ethinic origin.

Low-penetrance genetic variants might contribute to genetic predisposition to TNBC. In a genome-wide association study⁵⁷ of breast cancer in women of African ancestry (3153 cases and 11 330 controls), two novel risk loci for breast cancer were identified (rs432260 at chromosome 14q31, OR 1.18, $p=4.3 \times 10^{-6}$; and rs10510333 at chromosome 3p26, 1.15, $p=1.5 \times 10^{-5}$). However, in this study, the statistical power was too low to assess a potential association between single-nucleotide polymorphisms (SNPs) and ER-negative breast cancer risk, and so the associations did not meet the standard level of genome-wide significance.⁵⁷ A meta-analysis⁵⁸ of genome-wide association studies of ER-negative breast cancer using data from the breast cancer genome-wide association study in women of African ancestry and a European ancestry genome-wide association study identified a novel TERT-rs10069690 SNP in 5p15 for ER/PR-negative and TNBC risk. The risk allele frequency of rs10069690 was greater in women of African than European ancestry. These results were replicated in the Black Women's Health Study (which examined only African-American women; 1199 patients and 1948 controls), and showed that the rs10069690 SNP in 5p13·33 was significantly associated with ER/PR-negative breast cancer (OR 1·29, 95% CI 1.04-1.59) and TNBC (1.42, 1.02-1.99). In this independent sample of African-American women, the mean percentage of African ancestry estimated from AIMs was also

significantly higher in women with TNBC (83.0%) than in those with ER/PR-positive breast cancer (79.9%; p=0.008).⁵⁹ Although the functional significance of these findings are unknown, these results support a genetic cause of TNBC that might partly contribute to racial or ethnic origin differences in the incidence of the disease.

Response to systemic treatments

Chemotherapy is the standard treatment for TNBC. Despite the aggressive course of TNBC disease, it can be very sensitive to cytotoxic drugs: the proportion of patients achieving a pathological complete response after treatment with modern adjuvant or neoadjuvant chemotherapy range from 30–45%.^{60,61} Preliminary reports suggest that the proportion of patients achieving a pathological complete response might be increased by adding platinum salts to standard anthracyline and taxane-based chemotherapy. In a large meta-analysis⁶² including 12 neoadjuvant studies, 389 (33·6%, 95% CI 30·9–36·4) of 1157 patients with TNBC achieved a pathological complete response.

Irrespective of breast cancer subtype, patients who achieve a pathological complete response after neo-adjuvant chemotherapy have improved long-term outcomes compared with patients with residual disease identified at the time of surgery.^{63,64} Despite the poor prognosis associated with TNBC, patients who achieve a pathological complete response have excellent long-term outcomes, with overall survival of about 94%. Furthermore, if a pathological complete response is achieved, patients with TNBC and other subtypes of breast cancer have similar overall survival (p=0.24). However, patients with TNBC and residual disease have worse outcomes than do patients with residual disease and other subtypes of breast cancer (3-year overall survival 68% vs 88%, p<0.0001; figure 2).⁶³

TNBC represents a heterogeneous group of tumours. Masuda and colleagues⁶¹ assessed differential chemotherapeutic sensitivity according to Lehmann's TNBC subtype classification. Of 130 patients, there was a strong correlation between TNBC subtype and proportion of patients achieving a pathological complete response. The proportion of all 130 assessable patients with TNBC who achieved a pathological complete response was 28%; basal-like 1 subtype had the highest proportion of achievement of a pathological complete response (52%), followed by mesenchymal-like (31%), immunomodulatory (30%), and mesenchymal stem subtypes (23%). 10% of patients with the luminal AR subtype achieved a pathological complete response, and no patients with basal-like 2 subtype achieved a pathological complete response. Despite the low proportion of patients with luminal AR TNBC achieving a pathological complete response, this subtype was associated with the best overall survival.⁶¹

Few data exist for the chemotherapeutic sensitivity of TNBC in women of African ancestry. In a retrospective study that aimed to exclusively assess patients with TNBC, Dawood and colleagues⁶⁴ reported that 17% of African-American women achieved a pathological complete response, compared with a proportion of 25% for women of other ethnicities (p=0.09). After controlling for patient and tumour characteristics, ethnicity was not associated with relapse-free survival (hazard ratio [HR] 1.08, 95% CI 0.69–1.68), or overall

survival (1.08, 0.69-1.68) in patients who were white or of another race compared with African-American patients.⁶⁴

Similarly, we assessed the association between race or ethnic origin, and pathological complete response in 2074 patients treated with anthracycline-based and taxane-based chemotherapy.⁶⁵ Of 490 patients with TNBC, 19·4% of patients attained a pathological complete response, and we noted no significant differences according to race or ethnic origin. The proportions of patients achieving a pathological complete response in patients who identified their race were 16·8% in African- American patients, 20·8% in Hispanic patients, 20·2% in white patients, and 16% in patients of other racial or ethnic backgrounds (p=0·843). Although these results show the excellent outcome for patients with TNBC and chemosensitive tumours, and similar proportions of patients achieving pathological complete response in African-American patients who have treatment-resistant tumours. In a retrospective study⁶⁶ to assess predictors of tumour progression during neoadjuvant systemic chemotherapy, pretreatment clinical T3 status (OR 6·31, [95% CI $1\cdot81-21\cdot97$]), ER negativity (ER-positive status with 0·24 [0·13–0·44]), and African ancestry (2·07 [1·12–3·84]) were the most important predictors of tumour progression.

Treatment options for patients with TNBC

No data suggest that patients with TNBC of African ancestry should be treated differently from patients of European or Asian ancestry; therefore, present therapy guidelines apply to all patients irrespective of race or ethnic origin. Data from clinical trials stratifying outcomes according to race or ethnic origin are sparse, and the number of participants of African ancestry is very small, usually ranging between 5% and 10% of participants, or up to 15% in a few trials. The challenge for future clinical trials in TNBC is to improve the racial and ethnic diversity of participant enrolment to enhance the applicability of the results to other populations. Development of evidence-based and locally appropriate strategies to treat TNBC in women of African ancestry living in Africa will need a commitment to capacity building in cancer research and continuous efforts towards attaining this important goal.⁶⁷ For patients with early-stage or locally advanced TNBC, treatment with anthracycline and taxane-based regimens is preferred because it is associated with high rates of pathological complete response, which is a good surrogate marker for improved survival.^{63,64} In patients with metastatic disease, sequential single agents are frequently used, although combination chemotherapy is also part of the standard of care. No specific strategy is associated with improved survival; however, combination chemotherapy is associated with higher proportions of patients achieving an overall response, longer time to progression, and increased toxicity than are sequential single agents.

Ixabepilone and eribulin are some of the newest agents used in the treatment of metastatic TNBC. In the CA163- 046 trial 752 patients were randomly assigned to receive ixabepilone in combination with capecitabine, or capecitabine alone; subgroup analysis of the 187 patients with TNBC significantly favoured the combination group.⁶⁸ In a similar study with 1221 patients with metastatic breast cancer, Sparano and colleagues⁶⁹ reported that in the 256 patients with TNBC, the combination treatment was significantly associated with

improved durations of progression-free survival (HR 0.64, 95% CI 0.48–0.98). A subgroup analysis of 40 African-American patients did not show any benefit from the combination treatment against capecitabine alone, but the conclusions were restricted by the small number of patients in this analysis.⁶⁹

Tumours of *BRCA* mutation carriers have a deficiency in the repair of DNA double stranded breaks by homologous recombination and this phenotype has been termed BRCAness (or BRCA-like).⁷⁰ 30–50% of patients with TNBC are estimated to have BRCAness with loss of BRCA1 function. Identification of these BRCA-like tumours might have important therapeutic implications because of their frequent sensitivity to alkylating and platinum drugs. In patients with *BRCA* mutations, the reported proportions of patients achieving pathological complete responses with single-agent cisplatin were 83–100%.^{71,72} In a meta-analysis,⁷³ investigators assessed the effect of addition of platinum salts to standard neoadjuvant chemotherapy, which doubled the increase in proportions of pathological complete response (19·6% *vs* 48·4%, p<0·0001) in patients with TNBC.⁷³ The proportion of patients with *BRCA* mutations participating in these trials is unclear. The benefits of the addition of carboplatin or cisplatin could be driven by the very high proportions of pathological complete response in patients with *BRCA* mutations.

In the metastatic setting, the activity of single-agent platinum salts against TNBC is low, but the combination of platinum salts with other agents, such as gemcitabine, is synergistic. In a small phase 2 study⁷⁴ to assess the combination of gemcitabine and carboplatin to treat metastatic breast cancer, the median time to progression was $5\cdot3$ months (95% CI $2\cdot4-6\cdot7$ months) and the proportion of patients achieving an overall response was 31%.⁷⁴ In patients with TNBC, the combination was associated with a similar proportion of patients achieving an overall response (27% for patients who had not previously been treated with taxane and 32% for those who were pretreated with taxane).⁷⁵ These results are consistent with the proportion of patients achieving a clinical benefit of 35%, and the median duration of progression-free survival of $3\cdot6$ months recorded in the control group of a phase 2 study in which investigators assessed the use of gemcitabine and carboplatin with or without iniparib in patients with TNBC.⁷⁶

New targeted therapies for TNBC

Different subtypes of breast cancer are now being treated with different molecularly targeted therapies (figure 3). The gene profiling and DNA mutation data of breast cancer subgroups have identified many molecular signalling molecules that could be targeted for the treatment of breast cancer, including those in women of African ancestry. A major aim of present research is to discover effective therapies for TNBC.

Signalling inhibitors have activity in preclinical or early clinical tests. Poly (ADP-ribose) polymerase-1 (PARP) plays a key part in the repair of DNA single-strand breaks through the base excision repair pathway. It binds directly to the site of DNA damage where the polymerase recruits additional DNA repair enzymes.⁷⁷ PARP inhibitors cause specific DNA lesions that need homologous repair and are therefore interesting agents to target in cells that are deficient in repairing double-stranded DNA breaks. Data support the postulation that

BRCA1-mutated tumours are sensitive to PARP inhibitors via synthetic lethality and drugs, including olaparib, have proven to have significant single-agent activity in patients with *BRCA1* and *BRCA2* mutations.⁷⁸ Because of the important similarities between sporadic TNBC and *BRCA*-mutated breast cancers, the efficacy of these agents has been assessed in the treatment of TNBC. A randomised phase 2 study for patients with metastatic TNBC of iniparib in combination with gemcitabine and carboplatin seemed more effective than chemotherapy alone (progression-free-survival is 5.9 months [95% CI 4.5–7.2] for combination therapy *vs* 3.6 months [2.6–5.2] for chemotherapy alone; p=0.01 and overall survival 12.3 months [9.8–21.5] *vs* 7.7 months [6.5–13.3]; p=0.01);⁷⁹ however, these results could not be reproduced in a phase 3 trial of identical design. PARP inhibitors are likely to be active in *BRCA*-mutant cancers, but its efficacy against sporadic TNBC has not been shown.

Additional promising molecular targets include the androgen receptor (for luminal androgen-receptor TNBC), src (for mesenchymal TNBC), and insulin-like growth factor receptor, PI3K, protein kinase B (AKT), and mTOR (all molecules that transduce growth signals from growth factor receptors).²⁴ PI3K is especially promising because it is frequently mutated in human breast cancers.²³ Another promising target is the cyclin-dependent kinases.⁸⁰ These molecules phosphorylate crucial cell cycle regulators (cyclins) that regulate cancer cell growth. Specific small-molecule inhibitors of all these targets are being tested in early clinical trials for the treatment of breast cancer.

Immunomodulatory drugs are another important class of drugs that target the T-cell surface molecules CTLA4 (ipilimumab) or PD-1 (nivolumab). These drugs have shown great potential for treatment of melanoma and other cancers and are being considered for testing in breast cancer.⁸¹ The *TP53* tumour suppressor gene is another important target. This critical cell-cycle regulatory gene is the most commonly mutated gene in ER-negative breast cancers (around 80% of cases of TNBC show *TP53* gene mutations).²³ The P53 protein has been very difficult to target because of its dynamic and complex protein– protein interactions and substantial number of DNA promotor binding.⁸² No P53-specific therapies are clinically available. However, development of P53-specific therapies is a major effort in pharmaceutical companies and research laboratories.

Conclusion

Breast cancer is an increasing health-care problem throughout the world with large populations of women of African ancestry, whose disease is diagnosed at an advanced stage and is associated with high mortality. Women of African ancestry have a disproportionately high frequency of the aggressive TNBC subtype (20–79%) and present with disease at a younger age than do women of European ancestry. In African-American women, premenopausal status, increased parity, and shorter duration of breastfeeding are positively associated with an increased risk of TNBC, but whether these represent shared risk factors across other populations of African women is unknown. Genetic markers of African ancestry that predispose women to TNBC independent of known environmental risk factors have not been identified, but this is an area of active investigation. TNBC or basal-like breast cancer is a heterogeneous disease with variable responsiveness to chemotherapy

regardless of race or ethnic origin. High-priority clinical trials that combine novel targeted agents with standard chemotherapeutic regimens are being undertaken. An urgent need exists for collaborative multinational research that focuses on the causes, prevention, and treatment of TNBC in women of African ancestry.

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Search strategy and selection criteria

We searched PubMed for articles published in English between Jan 1, 2000, to July 1, 2014. Relevant articles were identified with the MeSH search terms "triple negative", "basal", "breast neoplasms", "African continental ancestry group", "African American", "tumour markers/biology", "epidemiology", "genetics", and "drug therapy". We obtained additional statistics on breast cancer incidence and mortality from population-based registries available on the websites of the International Agency for Research on Cancer, Pan American Health Organization, and Surveillance Epidemiology, and End Results Program.

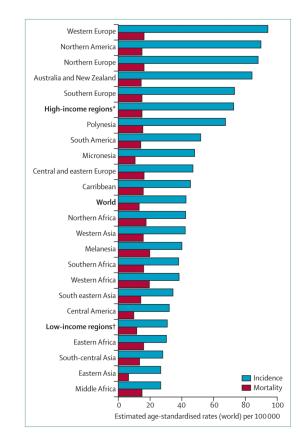


Figure 1. Global breast cancer incidence and mortality in women in 2012

*Includes all regions of Europe, Northern America, Australia, New Zealand, and Japan. †Includes all regions of Africa, Asia (excluding Japan), Latin America, the Caribbean, Melanesia, Federated States of Micronesia, and Polynesia. Region definitions from IARC WHO. Data from the International Agency for Research on Cancer, GLOBOCAN 2012.¹

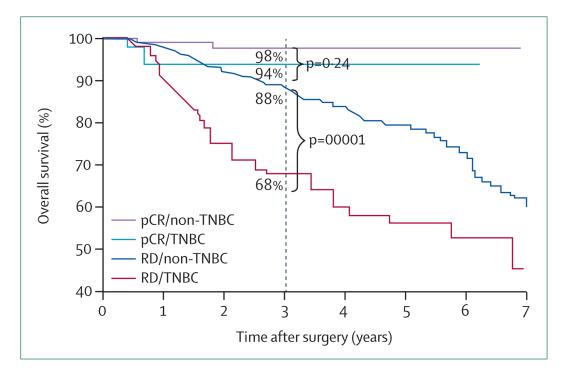


Figure 2. Overall survival after breast cancer surgery

Lines show survival as a function of response to chemotherapy (pathological complete response [pCR] *vs* residual disease [RD]) and triple-negative status (triple-negative breast cancer [TNBC] *vs* non-TNBC). Reproduced with permission from Liedtke and colleagues.⁶³

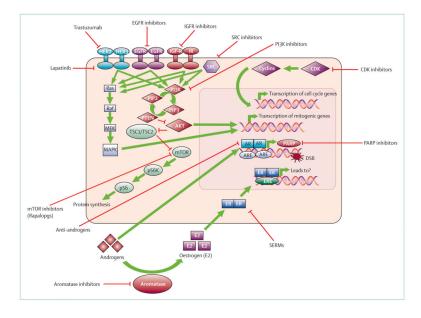


Figure 3. Oncogenic pathways associated with breast cancer

Selected genes and pathways differentially associated with specific breast cancer subtypes. Green arrows show activating signals. Red lines show inhibitory effects. Akt=akt proto-oncogene. AMPK=AMP-activated protein kinase. AR=androgen receptor. ARE=androgen response element. CDK=cyclin-dependent kinase. DSB=double-strand break. ER=oestrogen receptor. ERE=oestrogen response element. E2=oestrogen. IGFR=insulin-like growth factor receptor. IR=insulin receptor. MAPK=mitogen-activated protein kinase. MAPKK/MEK=mitogen-activated protein kinase kinase. P13K=phosphatidylinositol 3-kinase. PARP=poly ADP-ribose polymerase. pS6=ribosomal protein S6. pS6K=pS6 kinase. PTEN=phosphate and tensin homologue. Raf=raf-1 protooncogene. SRC=SRC proto-oncogene. TSC1=tuberous sclerosis 1. TSC2=tuberous sclerosis 2.

Table

Epidemiological risk factors for triple-negative breast cancer in women of African ancestry

	Number of cases of African ancestry (% of total cases)	Study design	Factors associated with triple-negative breast cancer
Bauer et al (2007) ¹⁵	636 (10%)	Cohort	Aged younger than 40 years (OR 1.53, 95% CI 1.37–1.70); low socioeconomic status (1.12, 1.01–1.24)
Millikan et al (2008) ³⁶	787 (43%)	Case-control	Parity ($p_{trend}=0.04$); short duration of breastfeeding ($p_{trend}=0.03$); large waist-to-hip ratio ($p_{trend}=0.002$)
Stead et al (2009) ³⁷	178 (43%)	Case-control	Overweight or obese (both non-significant)
Trivers et al (2009)38	116 (24%)	Case-control	Obesity (OR 1.89, 95% CI 1.22–2.92)
Ma et al (2010) ³⁹	849 (44%)	Case-control	Short duration of breastfeeding (p _{trend} =0.03); aged 45–65 years with oral contraceptive use before age 18 years (OR 2.87, 95% CI 1.44–5.74)
Shinde et al (2010) ⁴⁰	254 (10%)	Case-case	Parity (OR 1·12, 95% CI 1·06–1·20); short duration of breastfeeding (0·93, 0·90–0·97)
Ambrosone et al (2014) ⁴¹	786 (100%)	Case-control	Parity (p_{trend} =0.03); shorter duration of breastfeeding (non-significant)

OR=odds ratio. All studies were done in the USA.