

Breast Cancer in Latinas: Gene Expression, Differential Response to Treatments, and Differential Toxicities in Latinas Compared with Other Population Groups

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ABSTRACT

Disparities in clinical outcomes of breast cancer have been described among different racial and ethnic groups in the U.S. Convincing data exist showing that Latina women have a lower incidence of breast cancer but a higher breast cancer-related mortality rate compared with white women. Noticeable differences in breast cancer incidence are present even within different Latina subsets with a higher incidence in secondand third-generation women compared with foreign born. An increasing amount of data exists pointing to significant differences in the genetics and biology of breast cancer in Latinas as a significant contributor to the higher mortality, including a higher incidence of triple-negative breast cancers (which do not overexpress HER-2 protein and are negative for estrogen receptors and progesterone receptors). Other social and environmental factors are likely to play a significant role as well, including a lower rate of screening mammography, variable access to medical care, among others. Recent data are inconclusive regarding differences among racial/ethnic groups in the response to chemotherapy. Data on racial/ethnic variations in the pharmacogenomics of chemotherapy, endocrine treatments, and toxicity are more limited, with some data suggesting differences in frequencies of polymorphisms of genes involved in the metabolism of some of these agents. Further studies are needed on this subject. *The Oncologist* 2010;15: 466–475

INTRODUCTION

Several studies have shown that Hispanics have a lower incidence of breast cancer but a higher mortality rate compared with whites [1, 2]. Ethnic-specific studies have the potential to elucidate genetic causes of breast cancer, as illustrated by recent reports of the relationship between gene mutations and differential outcomes of high-risk childhood acute lymphoblastic leukemia in various racial and ethnic groups. These leukemia studies led to the discovery of specific genetic alterations that

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are more common in Latino children; these mutations help explain their poorer prognosis and helped introduce targeted therapies that could potentially improve outcomes [3, 4].

Latino/Hispanic populations are composed of a very heterogeneous group that includes many different racial and national backgrounds; thus, caution about generalizations is warranted [5]. In the U.S. in 2008, Latinos/Hispanics represented 46.9 million individuals (15% of the overall population) [6]. The majority of these individuals were of Mexican origin (64%) followed by individuals of Central American or South American origin (14.4%), Puerto Rican origin (9%), and Cuban origin (3.5%) [6]. Marked differences in breast cancer incidence rates exist among the Latina groups. For example, among Latina women in the San Francisco Bay area, the incidence of breast cancer is much higher among second- and third-generation women than among foreign-born women, suggesting a considerable influence from environmental factors [7].

In this article, we present a thorough review of the published literature on breast cancer in Latinas, emphasizing the data on differences in outcomes, differences in breast cancer gene expression, as well as the response to and toxicity of breast cancer treatment in the Latina population. The terms Latina and Hispanic will be used interchangeably.

DIFFERENTIAL OVERALL OUTCOME OF BREAST CANCER AMONG LATINA WOMEN

Breast cancer has a lower incidence rate among Latina women in the U.S. (90.1 cases per 100,000) compared with white women (130.6 per 100,000), although the incidence in Latinas has been increasing over the past 10 years [1, 2]. Although the reasons for this increase are unclear, obesity may be a contributing factor [8]. Nevertheless, breast cancer, despite its lower incidence rate, is the most common cancer and the leading cause of cancer death for Latina women. An estimated 14,300 Latina women were diagnosed with breast cancer in 2006, and an estimated 1,740 died of breast cancer in the U.S. in the same year [9]. Studies have attributed this difference in breast cancer outcome between Latinas and white women to several factors: 1) variable access to medical care, resulting in more advanced stage at diagnosis; 2) differing breast cancer risk factors; and 3) differences in genetics and biology. Some newer data are available that address these possible explanations.

Access to Medical Care

Low socioeconomic status is associated with worse outcome in breast cancer, regardless of ethnic origin. Latina women are more likely to be uninsured, have lower educa-

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tional achievements, and have higher poverty rates than white women in the U.S.; all of these factors potentially contribute to mortality rate differences [10]. In addition, Latina women are reported to be less likely to be compliant with treatment, probably because of fear associated with the diagnosis and treatment [11]. Language, cultural, and financial barriers may also significantly contribute to a higher noncompliance rate among Latinas [11].

Latina women are more likely to present with advancedstage breast cancer, presumably because of lower use of mammography screening services [10, 12-17]. Elledge et al. [18] reported that a higher percentage of white women present with node-negative breast cancer (55%) compared with Hispanics (43%) and blacks (42%), although no statistically significant differences were observed in overall survival or disease-free survival for patients with nodenegative disease among the different ethnic and racial groups. Fewer white women (39%) presented with nodepositive disease or locally advanced breast cancer compared with Hispanics (50%) and blacks (48%) (p < .001), and significant differences in overall survival (p = .008)and disease-free survival (p = .001) were observed among the racial/ethnic groups [18]. In addition, the proportion of women who presented with distant metastatic disease varied significantly by race (white, 5.4%; Hispanic, 6.9%; black, 9.5%; p < .001).

Multiple studies have evaluated breast conservation rates and postoperative radiotherapy in blacks, but similar studies in Hispanics are not available at this time. Studies evaluating practice patterns for patients with breast carcinoma who underwent breast conservation surgery showed that only 78.7% of black and Hispanic women received recommended systemic adjuvant therapy compared with 85.3% of white women [19, 20]. However, Elledge et al. [18] reported that for women with node-negative and nodepositive cancers, a greater proportion of Hispanic women received adjuvant chemotherapy compared with black and white women (10%–20% more). More white women received endocrine therapy than chemotherapy, but this likely was because of their higher rate of hormone receptor positivity [18].

Differing Breast Cancer Risk Factors

Although some studies have suggested that socioeconomic factors explain the increased frequency of more advanced stage of breast cancer in black and Hispanic women, most studies to date suggest that black and Latina women have a significantly higher incidence of late-stage breast cancer at presentation even after adjustment for a higher incidence of low socioeconomic status among minorities [21–23]. This suggests that biological and environmental factors specific

to the minority population are significant contributors to these observed disparities in outcome. The prevalence of established breast cancer risk factors such as reproductive risk factors (e.g., age at first birth, parity, and lactation), use of hormone-replacement therapy, and obesity varies across racial and ethnic subpopulations and likely contributes to the overall increased incidence of breast cancer in white women and to the more advanced-stage cancer at presentation in other ethnic groups. The increase in risk of breast cancer due to reproductive risk factors appears to vary in magnitude as well as pattern in different ethnic/racial groups. For example, nulliparous Hispanic women were found to be at a higher risk of developing breast cancer compared with white women, but increased parity has less of a protective effect on Hispanic women compared with white women [24]. In addition, breastfeeding, which is generally considered relatively protective against subsequent breast cancer, is more so in black and white women compared with Hispanic women [24]. These findings suggest that there may be significant differences in the effects of hormone and other reproductive factors on the causation of breast cancer in Hispanics, which point to the possibility of differences in polymorphisms in genes involved in estrogen metabolism in Hispanics. Further studies in this area are indicated.

A review of factors influencing differences in incidence and outcome of breast cancer in 156,570 postmenopausal women participating in the Women's Health Initiative study showed that black and Latina women had a lower incidence of breast cancer that was explainable by a lower prevalence of known risk factors (age; number of first-degree relatives with breast cancer; age at menarche, first birth, and menopause; and prior breast biopsy for benign breast disease) for breast cancer in minority population [10]. A population-based, case-control study conducted in New Mexico of 694 Hispanic and 813 non-Hispanic patients with breast cancer evaluated weight change and body mass index from age 18 to usual adult weight. Weight change and obesity were shown to be risk factors for breast cancer in Hispanic and non-Hispanic white women; however, the risk for Hispanic women was independent of menopausal status, whereas the risk for non-Hispanic women was seen only in postmenopausal women [8].

Differences in Genetics and Biology

Differences in the genetics and biology of breast cancer are likely to be significant contributors to the disparities in breast cancer outcomes among black and Latina women compared with white women [18]. Elledge et al. [18] evaluated 4,885 white patients, 1,016 black patients, and 777

Hispanic patients with breast cancer and reported significant differences in 5-year survival rates. Five-year survival was $75\% \pm 1\%$ for white patients (median survival, 166 months), $70\% \pm 2\%$ for Hispanic patients (median, 156 months), and $65\% \pm 2\%$ for black patients (median, 117 months). This survival pattern was noted even when evaluating patients with untreated, nonmetastatic disease (Table 1). In a recent study comparing a group of Hispanic women who had the same access to care as white women, more Hispanic women presented with larger, triple-negative breast cancers (these cancers do not overexpress HER-2 protein and are negative for estrogen receptors [ERs] and progesterone receptors [PRs]) at a younger age, suggesting that genetic and biologic differences may have a major role in explaining these differences in presentation and outcome [25]. In addition, studies have shown that Latina women with a family history of breast cancer have a markedly higher incidence of breast cancer [26].

Similar to black women. Latina women from Colorado who had the same access to care and same screening mammography rates as non-Hispanic white women were reported to present with breast cancer at a younger age, have a higher incidence of tumors larger than 5 cm (Table 2), and have a higher incidence of triple-negative tumors (Table 3) [18, 25, 27-30]. Using Arizona Cancer Registry data, investigators from the University of Arizona recently published similar findings of a higher incidence of triplenegative breast cancer among Latinas [31]. These findings suggest the presence of influential biologic and possibly genetic differences in breast cancer among Latina women compared with white women. Data on Hispanic women's breast cancer gene microarray analysis including genetic polymorphism analysis, estrogen and progesterone receptor expression, HER-2 receptor gene, BRCA1 and BRCA2 gene expression, and population genetics studies will be reviewed.

Breast Cancer Gene Microarray and Genetic Polymorphism Analysis

Several studies have shown differences in breast cancer gene expression among Latina women compared with white women, but more work is definitely needed. Most of the studies reported to date have attempted to identify genetic polymorphisms that affect inflammation, ER pathways, and lipid catabolism pathways and to determine their potential role in determining the risk of breast cancer. Studies evaluating the incidence of *BRCA1* and *BRCA2* mutations in the Hispanic population have also been reported [32]. Gene expression analyses and correlation with breast cancer treatment outcomes in Latinas are lacking in the medical literature.



Study	Hispanic		Black		White	
	No. of patients	Mortality rate	No. of patients	Mortality rate	No. of patients	Mortality rate
Jemal et al. [2]	NR	15.8/100,000	NR	33.5/100,000	NR	24.4/100,000
Curtis et al. [34]	1,172	HR, 1.24	2,479	HR, 1.63 ^a	35,878	HR, 1
Boyer-Chammard et al. [27]	875	HR, 2.3	185	HR, 1.2	10,937	HR, 1
Li et al. [35]	4,471	HR, 1.3 ^a	6,915	HR, 2 ^a	75,978	HR, 1
O'Malley et al. [36]	1,100	HR, 1.39	940	HR, 1.81	10,414	HR, 1

Baumbach et al. [33] recently presented genetic microarray analysis of 28 paraffin-embedded, triple-negative breast cancer samples from Hispanic (10 samples), white (8 samples), and black (10 samples) women. For each sample, normal tissue from the same patient was used as control to determine differential gene expression. A total of 9,399 transcripts were expressed in normal tissues from all 3 ethnic groups, and 10,296 transcripts were noted in the tumor tissues. Ethnic-specific expression patterns were observed in both tumor and normal tissue specimens. Significant differential expression of DNA repair pathway genes was observed in tumor samples from all 3 ethnic groups. These studies add further evidence for major differences in gene expression among ethnic groups, which may contribute to the observed disparities in outcome.

On the other hand, a recent single institution study of gene expression in breast cancer did not report any significant differences with regard to clustering of gene expression by ethnicity or genes differentially expressed after adjusting for multiple confounders [37]. This new study evaluated gene expression profiles of 98 triple-negative breast cancer biopsy specimens. These included specimens from 19 black, 23 Hispanic, and 56 white women.

The γ -aminobutyric acid A receptor is a multisubunit chloride channel that mediates the fastest inhibitory synaptic transmission in the central nervous system. γ -Aminobutyric acid A receptor, π (GABRP) is a type A receptor subunit that is not expressed in normal adult neuronal tissue but is expressed in reproductive and endocrine tissues. Breast tissue undergoes considerable physiologic change during pregnancy and the postpartum period. M.D. Anderson researchers [38] hypothesized that during pregnancy, progenitor cells that coexpress GABRP proliferate within the breast lobules and then are progressively lost during breastfeeding. Therefore, a full-term pregnancy followed by a short duration of lactation may allow survival of progenitor cells, with a higher risk of malignant transformation. Their evaluation of 203 newly diagnosed invasive breast cancers showed that higher *GABRP* gene expression indeed was more common in younger women with a limited history of lactation after pregnancy. Interestingly, the same study showed that *GABRP* gene expression was higher in Hispanic women compared with age-matched white controls. The authors suggested that *GABRP* gene expression may be associated with high-grade breast cancer in Hispanic women; this requires further study.

Slattery et al. [39] reported an analysis of 5 single-nucleotide polymorphisms of the *IL-6* gene and reported that among postmenopausal women not exposed to exogenous estrogen, the AG/GG genotype of rs1800797 (–596A>G) and the GC/CC genotype of rs1800795 (–174G>C) were associated with a significantly lower risk of breast cancer among Hispanic and non-Hispanic white women. Aspirin use among non–estrogen-using Hispanic women was associated with a lower incidence of breast cancer, particularly in carriers of the rs1800796 *IL-6* polymorphism.

Vitamin D receptors have a considerable role in modulating transcription of genes involved in calcium and phosphorous homeostasis, as well as those involved in cellular differentiation and proliferation. Polymorphisms in the 5' and 3' ends of the vitamin D receptor gene appear to confer phenotypic consequences for calcium and vitamin D metabolism, bone density, and prostate cancer risk [40]. Available data suggest that normal human breast epithelial cells and most breast cancers express the vitamin D receptor. Ingles et al. [40] genotyped polymorphisms in the vitamin D receptor gene for 143 Latina women with breast cancer and 300 controls and reported that polymorphic variation in or near the 3' end of the vitamin D receptor gene influences breast cancer risk in Latina women.

The genome-wide association studies evaluating genetic polymorphisms and breast cancer risk have not included Hispanic populations [41].

Study/Variable	Hispanics	Blacks	Whites	<i>p</i> -Value
Chlebowski et al. [10]				
No. of patients	103	242	3,455	
Tumor characteristics				
Mean size, cm	1.89	1.60	1.56	.12
Size 2–5 cm, %	15	18	13	.58
Size >5 cm, %	1	2	2	.58
Poorly differentiated, %	17.7	31.8	10.0	<.001
Gapstur et al. [42]				
No. of patients	410	1,114	11,715	
Tumor characteristics, %				
Size 2–5 cm	35	38	33	<.001
Size >5 cm	9	8	5	<.001
Positive LN, %	41	42	34	<.001
Elledge et al. [18]				
No. of patients	777	1,016	4,885	
Tumor characteristics, %				
Size 2–5 cm	51.3	42.6	44.6	<.001 (all)
Size >5 cm	16.6	27.7	10.9	<.001 (all)
Positive LN, %				
1–3	24.1	24.6	21.1	<.001 (H vs. W); <.001 (B vs. W); .31 (H vs. B)
>4	30	26.6	21	
Li et al. [35]				
No. of patients	7,219	10,560	97,999	
Tumor characteristics, %				
Size 2–4.9 cm	42.8	40.7	33.5	NR
Size >5 cm	13.9	4.6	3.3	NR
Positive LN, %				
2–4	12.2	11.7	9.8	NR
>5	12.6	11.6	8.7	NR
Tumor grade 3, %	37.8	40.8	29.4	NR
Hausauer et al. [43]				
No. of patients	15,355	19,105	193,513	
Tumor size >2 cm, %	43.5	45.6	34.4	NR
Shavers et al. [44]				
No. of patients	616	724	2,638	
Median tumor size, cm	3	2.8	2.5	<.001
Positive LN, %				
1–3	23.4	22.4	21.1	NR
4-10+	11.5	12.4	9.3	.004
Tumor grade 3, %	51.5	54.8	43.3	<.001
Abbreviation: B, black; H, His	ania IN Irma	h noda. ND	mot nomonto.	A W white

ER and PR Status

Few studies have evaluated the association between race/ ethnicity and tumor expression of ER and PR after adjusting for other tumor biologic characteristics. In general, Hispanic patients with breast cancer tend to have ER-negative tumors more frequently than non-Hispanic white women,



Study	Hispanic		Black		White	
	No. of patients	ER negative, %	No. of patients	ER negative, %	No. of patients	ER negative, %
Elledge et al. [18]	777	29.9	1,016	37.9	4,885	22.1
Chlebowski et al. [10]	103	17	242	29	3,455	13
Dunnwald et al. [45]	5,585	29.8	5,724	39.5	78,805	21.6
Chu et al. [46]	9,860	21.1	7,896	26.4	96,272	17.3
Li et al. [35]	4,471	24.1	6,915	33.7	75,978	18.8
Hausauer et al. [43]	15,355	17.3	19,105	24.2	193,513	14.2
Gapstur et al. [42]	410	30	1,114	42	11,715	26

although the difference was not as great as that seen between black and non-Hispanic white women [47]. Elledge et al. [18] also showed that ER-negative tumors were more common in Hispanic patients with breast cancer compared with non-Hispanic patients (29.9% vs. 22.1%; p < .0001) but less frequent than in black women (29.9% vs. 37.9%; p < .0001). Another study used data from 13 Surveillance, Epidemiology, and End Results cancer registries over a study period from 1992 through 2004 [43]. Researchers analyzed age-adjusted incidence rates and trends from women older than 50 years, stratified by race or ethnicity and tumor subtype, and showed an increase in incidence rates of ERand PR-negative tumors among Hispanic women compared with white women (14.2% in white women compared with 17.3% in Hispanic women). They reported a relative increase in rates (cases per 100,000 person-years) of triplenegative breast cancer in Hispanic women (from 2001 through 2004) of 26.8%, with an absolute increase from 34.6 (95% confidence interval [CI], 29.9-39.9) to 43.9 (95% CI, 39.0-49.9) [43].

HER-2 Receptor Status

In general, 10% to 15% of breast cancers are HER-2 positive, as determined by protein overexpression or gene amplification [48]. Bauer et al. [49] used the population-based California Cancer Registry to analyze the triple-negative phenotype. They showed that the most powerful risk factors for the triple-negative subtype were age younger than 40 years, non-Hispanic black race, and, to a lesser extent, Hispanic race (hazard ratio [HR] for non-Hispanic blacks, 1.77 [95% CI, 1.59–1.97]; for Hispanics, 1.23 [95% CI, 1.14– 1.34]). Elledge et al. [18] did not observe any significant differences in the frequency of HER-2-positive breast cancers in patients of different ethnic/race groups.

Increased activation of the AKT1 gene (also termed

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AKT) has been associated with HER-2/neu overexpression and with worse outcomes. In a recent analysis by Wu et al. [50], more than 70% of black and Latina women with HER-2/neu–positive tumors had overexpression of *AKT1*. In contrast, only 43% to 44% of Japanese and Swedish women with HER-2/neu–positive tumors had increased *AKT1* expression.

BRCA1 and BRCA2 Mutations

Among the nonwhite minorities, Hispanics with breast cancer have the highest incidence of pathogenic *BRCA1* mutations [32]. In a report of data collected from the Breast Cancer Family Registry of women younger than 65 years with breast cancer enrolled at the Northern California site, prevalence of the *BRCA1* mutation was 3.5% for Hispanics, 1.3% for blacks, 0.5% for Asian Americans, 8.3% for Ashkenazi Jews, and 2.2% for other non-Hispanic whites. The authors postulated that the higher prevalence of the *BRCA1* mutations among Hispanics may reflect a higher incidence of Jewish ancestry in this subset [32]. Given the higher incidence of triple-negative breast cancer among carriers of pathogenic *BRCA1* mutations, this particular genetic factor may heavily contribute to the described disparities in breast cancer outcome.

Large genetic rearrangements account for 8% to 15% of *BRCA* mutations. Recently, a deletion involving exons 9 through 12 of *BRCA1* was identified in 5 unrelated Mexican families that did not have an identifiable *BRCA* mutation [51]. The authors suggest that an assay for this mutation should be considered for sequence-negative, high-risk Hispanic patients. A large multiplex panel that tests for 18 mutations in *BRCA1* and *BRCA2*, which would replace complete sequencing of the *BRCA* genes, is currently being developed.

Population Genetics

Most individuals of Hispanic ancestry have genetic backgrounds that combine 2 of 3 different populations: European, African, and Native American [52]. Among Latinas in the San Francisco Bay area, women with a greater European background had a higher incidence of breast cancer [52]. In addition, marked differences between Mexicanborn women and U.S.-born Mexican American women have been described recently [53]. In that study, 652 women were recruited (309 from the U.S. and 343 from Mexico). The women differed with respect to the age at diagnosis of breast cancer (average age 48.7 years in the U.S. vs. 53.8 years in Mexico), number of live births (3.2 in the U.S. vs. 3.7 in Mexico), and the prevalence of family history of breast cancer (18.1% in the U.S. vs. 6.2% in Mexico).

The 4-Corners Breast Study is a case-control, population-based study of women in the southwestern U.S. The study enrolled 3,074 non-Hispanic white women and 1,674 Hispanic women. Findings from this study have suggested that genetics may significantly influence the disparities observed in breast cancer outcome in the Hispanic population. This study examined those with a family history of breast cancer and showed that Hispanic women had a higher incidence of triple-negative breast cancer, whereas non-Hispanic white women had a higher incidence of postmenopausal hormone receptor-positive breast cancer [54]. Among women younger than 50 years, a family history of breast cancer was associated with a higher risk of breast cancer among non-Hispanic whites compared with Hispanics, suggesting a stronger genetic predisposition among white women [55]. This difference in risk was not observed among older women in the study; for those women, a family history of breast cancer was associated with a similar risk of breast cancer for Hispanic and non-Hispanic white women [55].

DIFFERENTIAL RESPONSE TO TREATMENT

Chemotherapy

A number of studies have examined the differential responses and survival outcomes among women of different racial groups (especially black women) in response to chemotherapy and have shown conflicting results. After controlling for numerous factors, including socioeconomic status, most trials show that chemotherapy treatments have comparable efficacy among patients with breast cancer, at least when comparing blacks with whites [56]. This was further shown in a recent M.D. Anderson Cancer Center study, which demonstrated similar responses to chemotherapy in the neoadjuvant setting in black and white women [57]. An older study with a small number of patients (n =37) reported that different ethnic groups of poor urban women with locally advanced breast cancer did not significantly differ in their response to adjuvant chemotherapy with cyclophosphamide, methotrexate, and 5-fluorouracil or with 5-fluorouracil, doxorubicin, and cyclophosphamide plus radiotherapy [58]. Another study evaluated neoadjuvant chemotherapy and pathologic complete response to anthracycline- and taxane-based neoadjuvant chemotherapy in patients with stage II and III breast cancer [59]. Of 2,074 total patients, 14.6% were black, 15.2% Hispanic, 64.3% white, and 5.9% other. There was no differences in pathologic complete response rates among racial/ethnic group (12.5% in blacks, 14.24% in Hispanics, 12.3% in whites, and 11.5% others, p = 0.788). Although authors concluded that race/ethnicity was not a predictor of survival outcome, there appeared to be a trend toward improvement in relapse-free survival (HR 0.69 [95% CI 0.49-0.97]) and overall survival (HR 0.63 [95% CI 0.41-0.97]) in Hispanic women [59].

Contrary to the above studies, when racial disparities in survival were evaluated in 19,457 participants of Southwest Oncology Group trials, Albain et al. found that black patients had worse survival than white patients for early-stage premenopausal and postmenopausal breast cancer, advanced staged ovarian cancer, and advanced stage prostate cancer [60]. No statistically significant association between race and survival was seen for lung cancer, colon cancer, lymphoma, leukemia, or myeloma [60]. Another study, reported by Balmanoukian et al. [61], included 38 patients and showed a lower pathologic complete response in black patients compared with women in the white/other group.

Most studies have not been sufficiently powered to determine whether specific treatments have different efficacy and whether survival in minority populations differs from the overall survival of the study group.

Antiestrogen Therapy

Tamoxifen is extensively metabolized to several primary and secondary metabolites, including 4-hydroxytamoxifen and endoxifen, by various cytochrome P450 genes such as *CYP2D6* and *CYP2C9*. A recent study evaluated clinical data and blood samples from 301 patients (299 females, 2 males) who received at least 8 weeks of adjuvant tamoxifen therapy [62]. Most patients were white (68%) or Hispanic (26%). Hispanics had significantly higher serum levels of tamoxifen (p = .02) and 4-hydroxytamoxifen (p = .007) than whites.

Polymorphisms have been identified in the genes encoding a number of cytochrome P450 enzymes that lead to variable drug clearance rates. Considerable ethnic differences in the *CYP2D6* allele frequencies have been reported [63]. *CYP2D6* polymorphisms can be classified according to one of four metabolizer activity levels—poor, intermediate, extensive, and ultrarapid. Strong association was seen between *CYP2D6* activity and plasma levels of endoxifen. The extensive metabolizer phenotype is expressed by most people and is considered the normal phenotype. The prevalence of the poor metabolizer phenotype in the Hispanic population is reported to be 2.2% to 6.6%; in whites, it is 8.9% to 10%; and in Asians, it is 0% to 1.2%.

Human sulfotransferase 1A1 (*SULT1A1*) is an enzyme involved in the metabolism of 4-hydroxytamoxifen. Homozygosity for *SULT1A1*2/*2* leads to lower catalytic activity [64]. Nowell et al. [65] have shown that patients with breast cancer who were treated with tamoxifen and were homozygous for the *SULT1A1*2* allele had increased risk of death compared with those who were homozygous or heterozygous for *SULT1A1*1* [65]. Grabinski et al. [64] showed that Hispanics have a higher genotypic frequency of the *2/*2 variant genotype compared with whites, although the difference was not statistically significant.

DIFFERENTIAL TOXICITY RELATED TO TREATMENT

Chemotherapy

Unfortunately, the data are very limited with regard to differences in toxicity from treatment of breast cancer in different ethnic and racial groups. Han et al. [66] recently reviewed toxicity data among different ethnic groups after neoadjuvant chemotherapy with 5-fluorouracil, epirubicin, and cyclophosphamide. Significantly higher hematologic toxicity was observed in Asian women from Japan and Hong Kong compared with white and black women. Their analysis did not include Hispanic women. Some data report lower white blood cell counts in black women, which may contribute to delays in chemotherapy or dose reductions [20].

Antiestrogen Therapy

Differences in the incidence of polymorphisms of the aromatase gene among different ethnic groups have been reported [67]. These differences may lead to differential toxicity among ethnic groups. The MA.17 trial [68] studied postmenopausal women with hormone receptor–positive breast cancer who had completed 5 years of tamoxifen therapy. Study investigators compared data from 5 years of adjuvant aromatase inhibitor therapy versus placebo and showed that women of minority backgrounds were less likely to have toxicity from aromatase inhibitors. However, the beneficial effect of aromatase inhibitor therapy also was not observed in the subset of minority women. A significantly greater number of minority women reported not taking the letrozole daily. Latina women represented 1.5% of the participants in that study.

CONCLUSION

Differences in the incidence and outcome of breast cancer among Latina women compared with white women are well documented in the medical literature and likely explained by issues of access to medical care, different incidence of environmental factors including known breast cancer risk factors, and differences in genetics and biology of breast cancer. Data on the pharmacogenomics of chemotherapy and endocrine therapy response and toxicity are more limited, but some data do suggest differences in frequencies of polymorphisms involved in the metabolism of some of these agents. Further studies elucidating the pathogenesis of these differences are very important because they may lead to genetic discoveries with potentially significant therapeutic implications.

AUTHOR CONTRIBUTIONS

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