









Genetic Counseling, Testing, and Family Communication Into Survivorship After Diagnosis of Breast Cancer

Steven J. Katz, MD, MPH^{1,2,3} ; Paul Abrahamse, MA⁴ ; Allison Furgal, PhD⁴; Rachel Hodan, MS⁵ ; Rachel S. Tocco, MA¹ ; Kevin C. Ward, PhD⁶ ; Ann S. Hamilton, PhD⁷ ; Lauren P. Wallner, PhD, MPH^{1,8} ; and Allison W. Kurian, MD, MSc^{9,10} 

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ABSTRACT

PURPOSE To examine receipt of genetic testing and communication with relatives about results into survivorship after diagnosis of breast cancer.

METHODS Women age 20–79 years diagnosed with early-stage breast cancer in 2014–2015 and reported to the Georgia and Los Angeles County SEER registries were surveyed approximately 7 months and 6 years after diagnosis (n = 1,412). We asked about genetic counseling, testing, and communication with relatives about results. We categorized women into indications for testing on the basis of clinical guidelines at the time of diagnosis and at the time of the follow-up survey (FUPs).

RESULTS A total of 47.4% had indications for genetic testing at any time: 28.0% at baseline and an additional 19.4% at the time of the FUPs (only); 71.9% (95% CI, 67.4 to 76.4) of those with a baseline indication reported genetic testing versus 53.3% (95% CI, 47.3 to 59.2) with an indication at FUPs only and 35.0% (95% CI, 31.6 to 38.4) with no indication ($P < .001$). There were no significant racial or ethnic differences in receipt of testing, controlling for age and clinical indications ($P = .239$); results for genetic counseling were similar. Only 3.4% of survivors had direct-to-consumer genetic testing (DTCT) for cancer. Testers who reported a pathogenic variant (n = 62) were much more likely to have talked to most or all their first-degree adult relatives about genetic testing than those with a variant of unknown significance (n = 49) or a negative finding (n = 419): 62.7% versus 38.8% and 38.0%, respectively ($P < .001$).

CONCLUSION Many women with indications for genetic counseling and testing into survivorship do not receive it. But those tested reach out to family members on the basis of the clinical relevance of their results. Very few patients obtained DTCT, which suggests that these tests do not substitute for clinical testing in breast cancer survivors.

ACCOMPANYING CONTENT

 Appendix

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INTRODUCTION

Support for universal germline genetic testing after a breast cancer diagnosis is growing because of concerns that targeted guidelines^{1–3} fail to identify many patients who could benefit from genetic counseling and testing. Studies show that about one third of patients with breast cancer meet clinical practice guidelines for genetic counseling and testing at the time of diagnosis, but many do not receive it.^{4–8} Access to genetic counseling and testing for the 280,000 people diagnosed with breast cancer in the United States each year is increasingly important because results influence locoregional and systemic treatment decisions and inform the risk of second primary cancers over time.^{9,10} In addition, genetic testing results in patients have important implications for

cancer risk stratification and prevention for relatives. Thus, genetic counseling and testing is important for the nearly four million survivors of breast cancer living in the United States and their family members. However, virtually nothing is known about its uptake in the years after a diagnosis of breast cancer. Untested patients with clinical indications for genetic counseling and testing at the time of diagnosis continue to benefit from counseling and testing in the survivorship period, which we consider as the period after completion of first-course therapy with surgery, radiation, and/or chemotherapy. In addition, patients who did not have an indication at the time of initial treatment management may meet criteria later because of a diagnosis of a new primary cancer or metastatic recurrence, a change in pertinent family history, or a change in testing guidelines.

CONTEXT

Key Objective

What is the uptake of genetic testing and counseling in a cancer-registry based cohort of survivors of breast cancer up to 6 years after diagnosis?

Knowledge Generated

Clinical indications for genetic testing increased into survivorship, but many eligible women did not receive it. Tested patients reach out to family members on the basis of the clinical relevance of their results. Few patients reported interest in direct-to-consumer testing and fewer obtained it, which suggests that these test options are not substituted for clinical testing in breast cancer survivors.

Relevance (S.B. Wheeler)

Germline genetic testing is increasingly important after a diagnosis of cancer, for treatment management and for cancer risk reduction in families with hereditary cancer syndromes, yet many survivors and their family members do not receive it, representing an important area for future inquiry.

*Relevance section written by JCO Associate Editor Stephanie B. Wheeler, PhD, MPH.

We examined patient report of genetic counseling, clinical genetic testing, and use of direct-to-consumer genetic testing (DTCT) from diagnosis through the first 6 years of survivorship in a diverse, population-based cohort of women diagnosed with breast cancer in 2014–2015 as reported to the SEER registries of Georgia and Los Angeles County. We hypothesized that a substantial proportion of survivors who did not receive genetic counseling and testing at the time of diagnosis did so during the survivorship years. We further hypothesized that a considerable number of survivors who did not receive clinical testing might have used DTCT instead.

METHODS

The iCanCare Study is a population-based, longitudinal survey study of women with early-stage breast cancer and their clinicians. As detailed previously,⁷ women age 20–79 years who were newly diagnosed with early-stage breast cancer (stages 0–II) in 2014–2015 as reported to the SEER registries of Georgia and Los Angeles County were surveyed. African American, Asian, and Latina women were oversampled. Women were ineligible if they had stage III or IV disease, had tumors larger than 5 cm, or could not complete a questionnaire in English or Spanish (n = 258). A total of 2,502 women completed surveys, resulting in a 68.0% baseline response rate. The median time from diagnosis to completion of the initial baseline survey was 7.8 months (25%–75% range, 5.6–10.1 months) and 83.6 months (25%–75% range, 53.1–86.8 months) for the follow-up survey (FUPs).

We sent respondents a paper FUPs approximately 6 years after their initial diagnosis in 2021–2022, with an option to complete the survey online. As in previous work, in this

study, we used a modified Dillman approach to patient recruitment, including reminders to nonrespondents and a \$20 in US dollars (USD) up-front cash incentive. Patients were deemed ineligible for the follow-up study if they were deceased (n = 108) or were too ill (n = 33) to participate. The FUPs was completed by 1,412 of the 2,361 eligible women (FUPs response rate of 59.8%; Appendix Fig A1 [online only]). Responses to the surveys were merged with SEER clinical data, and a deidentified analytic data set was created. The study was approved by the University of Michigan Institutional Review Board (IRB) and the state and institutional IRBs of the SEER registries. We obtained informed consent from each participant.

We asked women in the FUPs about the occurrence of new primary cancers by cancer type (including breast), recurrence of breast cancer (including anatomic location), their family history of cancer, receipt of genetic counseling, receipt of clinical germline genetic testing and results, communication with relatives about cancer genetic testing, and use of DTCT after diagnosis. We categorized women into indications (yes/no) on the basis of the National Comprehensive Cancer Network (NCCN) Clinical Practice Guidelines in Oncology (NCCN Guidelines) at the time of diagnosis (genetic counseling and testing indication baseline) and at the time of the FUPs (genetic counseling and testing indication at FUPs only).

Outcome Measures

All outcome measures were derived from the FUPs:

1. Patients were first asked “How long has it been since you last had a counseling session with a genetic counseling expert—that is, an appointment where the whole discussion is about genetic cancer risk?”

2. The next question was “How long has it been since you last had a blood or saliva genetic test for future cancer risk that was ordered by a doctor or genetic counselor?” Response categories were: Never, within the past 2 years, 2–5 years ago, and more than 5 years ago.
3. Patients who reported receipt of testing were then asked: What were the results of the most recent genetic test that was ordered by a doctor or genetic counselor? Please mark all that applies (negative, positive for a gene mutation, uncertain [variant of unknown significance, VUS]). Patients who reported a result of positive for a gene mutation were coded as positive, regardless of other responses checked. Patients who reported a VUS result and who did not report a positive result were coded as VUS. Patients who reported negative results and no positive or VUS results were coded as negative.
4. Patient report of DTCT followed the questions above. We framed these questions on tests sold by companies that offer genetic tests for cancer risk on the internet, without the need to involve your doctor. Anyone can buy these tests online, get a testing kit in the mail, collect their spit in a special cup or tube, and mail the test kit back to the company for analysis. Examples of companies offering this direct-to-consumer testing include 23andMe, AncestryDNA, and Color.
5. We first asked “How much have you researched these types of tests online?” Response categories used a five-point Likert scale from not at all to a lot. We then asked, “Have you ever taken a direct-to-consumer genetic test for cancer risk that you ordered on the internet?” Examples included 23andMe, AncestryDNA, Color, and Other, and for each example, response categories were yes and no.
6. Patient report of communication with family members about their test results: “Have you talked with your immediate adult blood relatives (parents, brothers and sisters, children) about getting clinical genetic testing to learn more about their own future cancer risk?” Response categories were as follows: Yes, I have talked to most or all of my adult family members; Yes, I have talked to some of my adult family members (but not all); and No, I haven’t talked to any adult family members.

Independent Variables

Appendix [Table A1](#) shows the criteria for genetic counseling and testing indications on the basis of NCCN guidelines at the time of baseline¹¹ and FUPs.¹² Indications at baseline were derived from the baseline survey and the SEER data (presence of triple-negative breast cancer [TNBC] subtype). Indications at the time of the FUPs were derived from that survey and the SEER data (presence of TNBC subtype). Other covariates included age at diagnosis, race and ethnicity, education level, and annual household income all derived from the baseline survey; clinical stage and grade at the time of diagnosis and geographic site were derived from the SEER data.

Analytic Plan

We first described patient characteristics for the analytic sample of 1,412 respondents who completed the baseline and FUPs after diagnosis ([Table 1](#)). We then showed patient report of genetic risk evaluation by indication for genetic counseling and testing and by timing of testing and counseling during the study period. We then examined patient report of testing by race and ethnic groups, controlling for age, education, and clinical stage. Next, we described engagement and communication with relatives about genetic test results by self-reported results outcomes (pathogenic variant [PV], VUS, negative). Finally, we described patient report of use of DTCT during the study period. To account for the effects of differential response rates, we repeated all analyses using analytic weights on the basis of covariates with significantly different response rates and examined the results for differences from the unweighted analyses.

RESULTS

Population Characteristics

[Table 1](#) shows characteristics of the study population. The median age was 55.3 years; about half were of minoritized race or ethnicity (18.2% Black, 18.0% Latina, and 9.5% Asian); 27.1% had a high school education or less; and 31.8% reported annual income of less than \$40,000 (USD). The distribution of clinical stage and grade at the time of diagnosis reflected the selection of patients, with more favorable disease in the inception cohort. Patient report of a second primary breast cancer (3.5%) or metastatic recurrence (1.5%) was uncommon. The patient sample was nearly equally distributed between the two state registries (51.6% from Georgia v 48.4% from Los Angeles County). Less than half of the respondents (47.4%) had indications for genetic risk evaluation over the study period: 28.0% at baseline and an additional 19.4% at the time of the FUPs (only).

Genetic Testing and Counseling

[Figure 1](#) shows patient-reported genetic testing and genetic counseling at FUPs by indication category (baseline, FUPs only, no indication). Nearly three quarters (71.9%; 95% CI, 67.4 to 76.4) of those with a baseline indication reported genetic testing over the observation period, versus 53.3% (95% CI, 47.3 to 59.2) with an indication at FUPs only and 35.0% (95% CI, 31.6 to 38.4) of those with no indication ($P < .001$). A substantial proportion of those who reported testing received the test during the survivorship period: 13.0% of testers with baseline indications tested in the past 2 years of the survey versus 19.5% of those with indications in FUP only ($P < .001$). Assessment of confounding showed no substantial effects of age, education, or clinical stage, and thus, these descriptive results are not adjusted.

TABLE 1. Characteristics of the Study Population

Characteristic	No. (%)
Age, years (missing = 1)	
<40	28 (2.0)
40-49	168 (11.9)
50-59	364 (25.8)
60-69	522 (37.0)
≥70	329 (23.3)
Race (missing = 29)	
White	750 (54.2)
Black	252 (18.2)
Latina	249 (18.0)
Asian	132 (9.5)
Education (missing = 33)	
High school or less	373 (27.1)
Some college or technical school	401 (29.1)
College graduate or higher	605 (43.9)
Income (missing = 232)	
<\$20,000 USD	174 (14.7)
\$20,000-~\$40,000 USD	201 (17.0)
\$40,000-~\$60,000 USD	202 (17.1)
\$60,000-~\$90,000 USD	212 (18.0)
≥\$90,000 USD	391 (33.1)
Stage (missing = 32)	
0	267 (19.4)
I	773 (56.0)
II	340 (24.6)
Grade (missing = 63)	
1	381 (28.2)
2	615 (45.6)
3	353 (26.2)
New breast cancer since diagnosis (missing = 31)	
No	1,332 (96.5)
Yes	49 (3.5)
Breast cancer distant recurrence since diagnosis (missing = 29)	
No	1,362 (98.5)
Yes	21 (1.5)
Geographic site	
Georgia	729 (51.6)
Los Angeles	683 (48.4)
Indication for genetic risk evaluation after diagnosis	
Baseline	395 (28.0)
Follow-up only	274 (19.4)
No indication	743 (52.6)

Abbreviation: USD, US dollars.

Figure 2 shows that there were no significant racial or ethnic differences in receipt of testing during survivorship, controlling for age and clinical indications ($P = .239$). Results for report of genetic counseling were very similar (Appendix Figs A2 and A3).

Communication of Test Results With Family Members

Testers who reported a PV result ($n = 62$) were much more likely to have talked to most or all their first-degree adult relatives about genetic testing than those with a VUS ($n = 49$) or a negative finding ($n = 419$): 62.7% versus 38.8% and 38.0%, respectively ($P < .001$).

Interest and Receipt of DTCT

Overall, there was very little interest in DTCT for cancer risk: 5.1% researched DTCT online somewhat to a lot and only 3.4% had DTCT.

Follow-up response rates differed by income, employment, education, race, receipt of hormonal therapy, and having subsequent nonbreast cancer. To account for possible bias related to differential response, we generated weights and reran all analyses using the weights. There were no meaningful differences in our results.

DISCUSSION

We performed a large, SEER-based longitudinal survey study of patients diagnosed with early-stage breast cancer in 2014–2015 at two time points: 7 months after diagnosis and then 6 years into survivorship. We found that a substantial proportion of women met NCCN guidelines for genetic counseling and testing that were published during the initial diagnosis and treatment periods. In addition, many of those who were not candidates for genetic counseling and testing at the time of diagnosis became eligible over the course of the survivorship period because of pertinent new cancers or additional family history and the somewhat broader indications promulgated by NCCN guidelines over the course of the study period. Yet, many women eligible for genetic counseling and testing did not receive it. We observed this gap uniformly across race and ethnic groups, with no significant differences across subgroups.

In response to growing evidence for the clinical utility of testing and studies suggesting undertesting, several professional organizations have expanded the criteria for genetic counseling and testing^{13,14} and there is growing advocacy for near-universal germline testing after diagnosis of breast cancer.^{15–17} It has become even more important to increase testing after diagnosis as evidence continues to grow about the need for germline test results for both locoregional and systemic management.¹⁰ In addition, germline genetic testing after diagnosis of breast cancer is an essential strategy to close the unacceptable gap in cascade testing of families with hereditary cancer risk.^{18–21}

The survivorship period that immediately follows an often-arduous initial course of therapy is an essential time of recovery for patients with breast cancer. However, there are important clinical issues during survivorship that warrant close engagement and continuity with medical oncology,

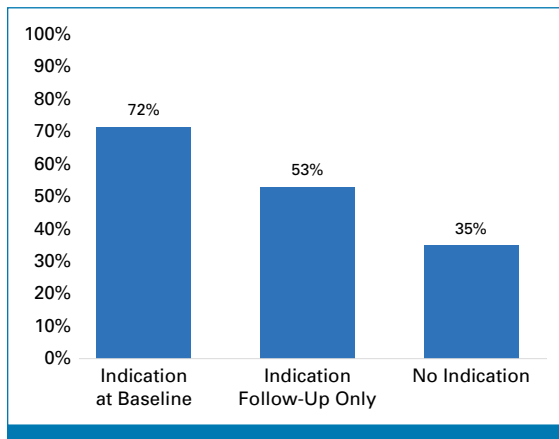


FIG 1. Percent of respondents who reported germline genetic testing in the FUPs by clinical guidelines at the time of the baseline survey (at the time of diagnosis) and at the time of the FUPs (approximately 6 years after diagnosis). FUPs, follow-up survey.

including treatment related side effects, medication management for patients on longer-term therapies, and assessment and management of future cancer risk in patients and their families through genetic counseling and testing. Oversight by medical oncologists and other clinicians including primary care during the survivorship period may result in missed opportunities to optimize patient and family outcomes. A particular challenge is the need to record an accurate and up-to-date family history of cancer, which may be underascertained in follow-up encounters with patients.

Our results suggest optimism in engaging more patients in genetic counseling, testing, and family communication during survivorship. During the study period, rates of

testing and counseling after a diagnosis of breast cancer increased in the geographic regions of our study²² and our findings demonstrate high rates of testing and counseling over time, especially in patients who had indications at the time of cancer diagnosis. We did not observe significant race and ethnic disparities in counseling or testing. In addition, our results suggest that patients reach out to family members on the basis of the clinical relevance of their results. Finally, a reassuring finding from our study is that very few patients reported any interest in DTCT and fewer obtained it across the long survivorship period. There has been concern that a substantial number of patients may seek DTCT and fail to differentiate DTCT from clinical grade testing.²³⁻²⁷ Our findings suggest that DTCT is not substituted for clinical testing by breast cancer survivors.

Although the response rate for the two surveys in this longitudinal study was high, our results could have been biased by differential response rates. We accounted for this possibility by performing weighted analyses on the basis of measured covariates, but there might have been differential response rates by unmeasured ones. We might have misclassified indications for genetic counseling and testing because some pertinent patient personal or family cancer history was not ascertained. Outcomes were derived from patient reports, which may be prone to recall bias. Reassuringly, in our previous work with this cohort, we found good concordance between patient self-report of genetic testing in the baseline survey (29%)⁶ and linkage to genetic testing data obtained directly from laboratories (26%) near the time of cancer diagnosis.²⁸ In addition, survey face and construct validity were high. Finally, the lower rate of testing observed in patients with indications at follow-up only versus baseline is partly related to the shorter observation period between the two groups.

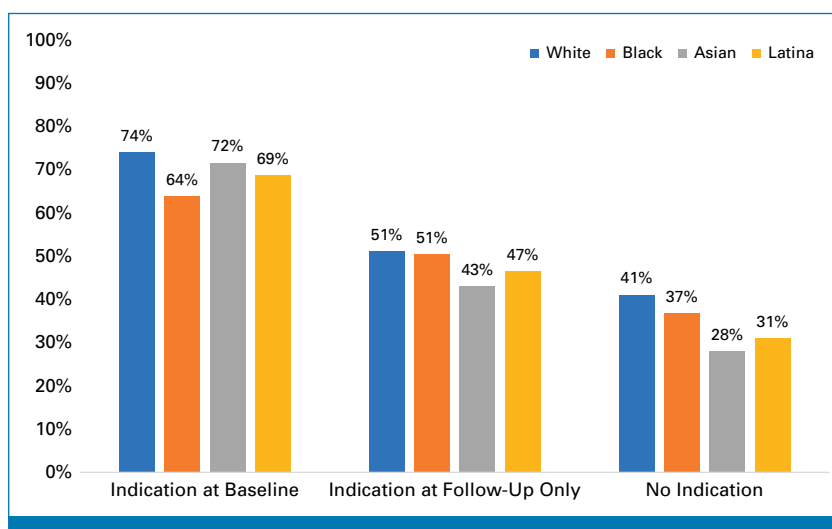


FIG 2. Percent of respondents who reported genetic testing in the follow-up survey by guideline indication and race and ethnic group identity. Results are adjusted for age, education, and clinical stage at diagnosis.

In conclusion, germline genetic testing is increasingly important after a diagnosis of cancer, for treatment management and for cancer risk reduction in families with hereditary cancer syndromes. While it is ideal to obtain genetic testing results during treatment planning, the survivorship period remains a major missed opportunity to engage patients and family members who may be at risk for hereditary cancer susceptibility and may be candidates for effective risk reduction and treatment strategies. Proponents of universal testing of all patients with breast cancer argue that it would increase the detection of clinically meaningful results, reduce disparities in receipt of testing, and facilitate a clear focus on cascade testing and cancer risk reduction for survivors and family members.¹⁴ However, a potential adverse outcome is more detection of meaningless results, particularly VUS, which are more frequently detected in racially and ethnically minoritized

groups. Indeed, broadening clinical guidelines in conjunction with the advent of larger multigene test panels has already markedly increased the rate of clinically less meaningful results, particularly VUS.^{22,29} Proponents of broadening guidelines argue that clinically noncontributory findings such as VUS can be managed successfully by clinicians and that the failure to detect meaningful PVs is a much bigger problem. More research is needed on the potential adverse consequences of less clinically meaningful test results on the management of breast cancer and engagement with families regarding cascade genetic risk evaluation. In addition, more testing of patients diagnosed with cancer yearly, and of the growing number of cancer survivors, should motivate more research to evaluate and implement multipronged strategies to facilitate genetic counseling, testing, and outreach to family members in clinical practice.

AFFILIATIONS

¹Department of Medicine, University of Michigan, Ann Arbor, MI

²Department of Health Management and Policy, School of Public Health, University of Michigan, Ann Arbor, MI

³Department of Internal Medicine, University of Michigan, Ann Arbor, MI

⁴Department of Biostatistics, School of Public Health, University of Michigan, Ann Arbor, MI

⁵Cancer Genetics, Stanford Health Care, Stanford, CA

⁶Department of Epidemiology, Rollins School of Public Health, Emory University, Atlanta, GE

⁷Department of Population and Public Health Sciences, Keck School of Medicine, University of Southern California, Los Angeles, CA

⁸Department of Epidemiology, School of Public Health, University of Michigan, Ann Arbor, MI

⁹Department of Medicine, Stanford University, Stanford, CA

¹⁰Department of Epidemiology and Population Health, Stanford University, Stanford, CA

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AUTHORS' DISCLOSURES OF POTENTIAL CONFLICTS OF INTEREST

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AUTHOR CONTRIBUTIONS

Conception and design: Steven J. Katz, Paul Abrahamse, Rachel S. Tocco, Lauren P. Wallner, Allison W. Kurian

Financial support: Steven J. Katz, Lauren P. Wallner

Administrative support: Steven J. Katz, Lauren P. Wallner

Provision of study materials or patients: Steven J. Katz, Kevin C. Ward, Ann S. Hamilton, Lauren P. Wallner

Collection and assembly of data: Steven J. Katz, Paul Abrahamse, Allison Furgal, Rachel S. Tocco, Kevin C. Ward, Ann S. Hamilton, Lauren P. Wallner

Data analysis and interpretation: Steven J. Katz, Paul Abrahamse, Allison Furgal, Rachel Hodan, Rachel S. Tocco, Allison W. Kurian

Manuscript writing: All authors

Final approval of manuscript: All authors

Accountable for all aspects of the work: All authors

CORRESPONDING AUTHOR

Steven J. Katz, MD, MPH; e-mail: skatz@umich.edu.

DISCLAIMER

The ideas and opinions expressed herein are those of the authors. The State of California, Department of Public Health, the NCI, and the CDC and their contractors and subcontractors had no role in design and conduct of the study; collection, management, analysis, and interpretation of the data; and preparation, review, or approval of the manuscript.

EQUAL CONTRIBUTION

L.P.W. and A.W.K. contributed equally to this work.

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Lauren P. Wallner

Honoraria: Kaiser Permanente

Consulting or Advisory Role: Gilead Sciences

Travel, Accommodations, Expenses: Gilead Sciences

Allison W. Kurian

Other Relationship: Ambry Genetics, Color Genomics, GeneDx/
BioReference, InVita, Genentech, Myriad Genetics, Adela, Merck, Gilead
Sciences

No other potential conflicts of interest were reported.

APPENDIX

TABLE A1. Indications for Genetic Risk Evaluation During the Study Period

Indications at Baseline Survey	Indications at Follow-Up Survey
TNBC age ≤ 60 years	TNBC at any age
Patient age ≤ 45 years at diagnosis	Patient age ≤ 45 years at diagnosis
Breast cancer in any male relative	Breast cancer in any male relative
First-degree relative diagnosed with breast cancer at age < 50 years	Close relative diagnosed with breast cancer at age < 50 years
First-degree relative diagnosed with sarcoma	First-degree relative diagnosed with sarcoma
First-degree relative diagnosed with ovarian cancer	Close ^a relative diagnosed with ovarian cancer
Two or more first-degree relatives diagnosed with breast cancer	Two or more close relatives diagnosed with breast cancer
	Patient with breast cancer between age 46 and 50 years and close relative with breast cancer at any age
	Patient with breast cancer between age 46 and 50 years and a second primary breast cancer diagnosed at any age
	Two primary breast cancers in patient and family history of breast cancer in close relative at any age
	Patient with pancreatic cancer
	Patient ^b with metastatic breast cancer

NOTE. Cancer diagnosis including biologic subtype reported through SEER; all other information is from patient surveys. Indications per NCCN guidelines that could not be included because the information was not ascertained: (1) lobular breast cancer and family history of diffuse gastric cancer; (2) family history of pancreatic cancer; (3) family history of high-grade prostate cancer; (4) has a mutation probability model score of $> 5\%$; and (5) personal history of breast cancer < 50 with unknown or limited family history. Citations for indications for genetic risk evaluation: At baseline survey: Daly et al.¹¹ At follow-up survey: NCCN.¹²

Abbreviations: NCCN, National Comprehensive Cancer Network; TNBC, triple-negative breast cancer.

^aClose relative includes first-degree, second-degree, and third-degree relatives.

^bPatient reported recurrence distant from breast.

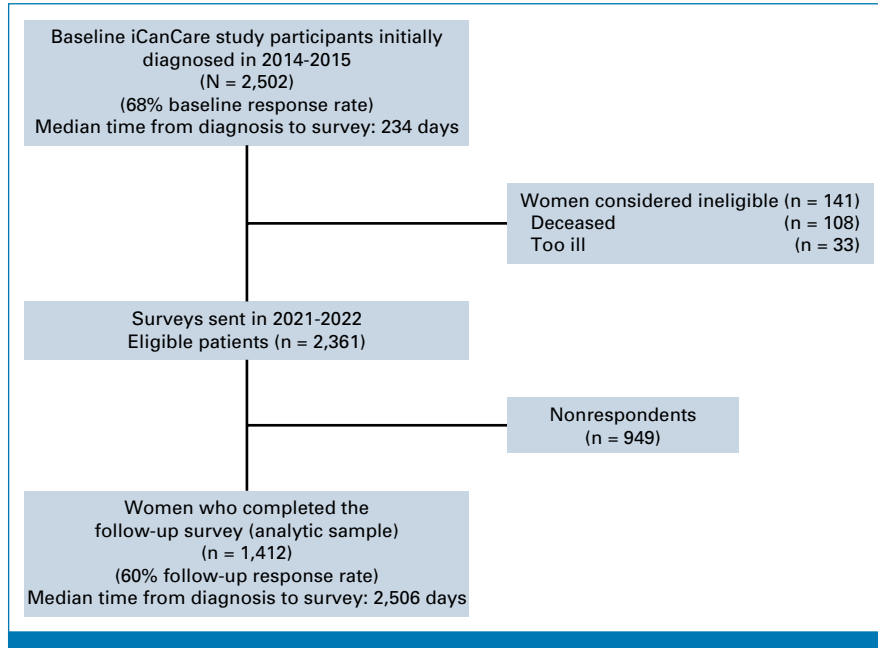


FIG A1. iCanCare follow-up study flow.

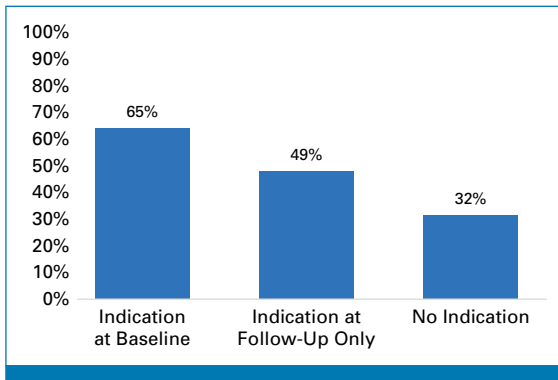


FIG A2. Rates of genetic counseling by indication.

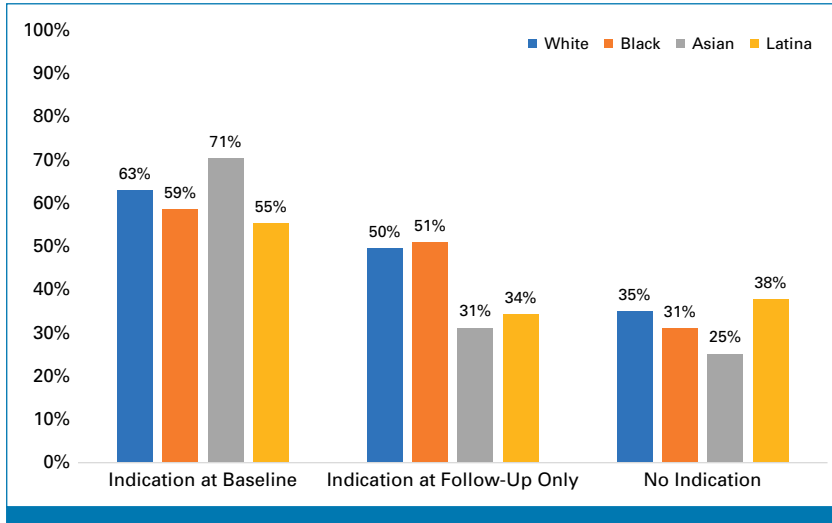


FIG A3. Adjusted rate of genetic counseling by race and indication. Adjusted for age, education, and stage.