

Protocol

Genetic Testing for Breast Cancer Gene Expression Prognosis Assay

(20436, 20476)

(Content formerly found in: Assays of Genetic Expression in Tumor Tissue as a Technique to Determine Prognosis in Patients With Breast Cancer and Quantitative Assay for Measurement of HER2 Total Protein Expression and HER2 Dimers)

Medical Benefit		Effective Date: 07/01/17	Next Review Date: 09/17
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Preauthorization is required.

The following protocol contains medical necessity criteria that apply for this service. The criteria are also applicable to services provided in the local Medicare Advantage operating area for those members, unless separate Medicare Advantage criteria are indicated. If the criteria are not met, reimbursement will be denied and the patient cannot be billed. Please note that payment for covered services is subject to eligibility and the limitations noted in the patient's contract at the time the services are rendered.

Populations	Interventions	Comparators	Outcomes
Individuals: • With early-stage node-negative invasive breast cancer considering adjuvant chemotherapy	Interventions of interest are: • Gene expression profiling testing	Comparators of interest are: • Clinical risk prediction algorithms	Relevant outcomes include: • Disease-specific survival • Change in disease status
Individuals: • With early-stage node-positive invasive breast cancer considering adjuvant chemotherapy	Interventions of interest are: • Gene expression profiling testing	Comparators of interest are: • Clinical risk prediction algorithms	Relevant outcomes include: • Disease-specific survival • Change in disease status
Individuals: • With ductal carcinoma in situ considering radiotherapy	Interventions of interest are: • Gene expression profiling testing	Comparators of interest are: • Clinical risk prediction algorithms	Relevant outcomes include: • Change in disease status
Individuals: • With early-stage node-negative invasive breast cancer free of distant recurrence at 5 years	Interventions of interest are: • Gene expression profiling testing	Comparators of interest are: • Clinical risk prediction algorithms	Relevant outcomes include: • Disease-specific survival • Change in disease status
Individuals: • With breast cancer who are undergoing assessment of HER2 status	Interventions of interest are: • Assessment of HER2 status using quantitative total HER2 protein expression and HER2 homodimer measurement	Comparators of interest are: • Assessment of HER2 status using immunohistochemistry or fluorescence in situ hybridization	Relevant outcomes include: • Overall survival • Disease-specific survival • Test accuracy • Test validity

Description

Laboratory tests have been developed that detect the expression, via messenger RNA, of many different genes in breast tumor tissue and combine the results into a prediction of distant recurrence risk for women with early-stage breast cancer. Test results may help providers and patients decide whether to include adjuvant chemotherapy in postsurgical management of breast cancer or to alter treatment in patients with ductal carcinoma in situ (DCIS).

Novel assays that quantitatively measure total human epidermal growth factor receptor 2 (HER2) protein expression and homodimers have been developed to improve the accuracy and consistency of HER2 testing.

Summary of Evidence

For all tests and all indications, relevant outcomes include disease-specific survival and changes in disease status.

Early-Stage Node-Negative Invasive Breast Cancer

Only studies presenting ten year distant recurrence rates in node-negative women not receiving adjuvant chemotherapy were included in this review. In addition to negative nodes, the type of patient considered for this indication have positive hormone receptors and are human epidermal growth factor receptor 2 (HER2) negative.

21-Gene Assay (Oncotype DX)

For individuals who have early-stage node-negative invasive breast cancer considering adjuvant chemotherapy who receive gene expression profiling with the 21-gene assay (Oncotype DX), the evidence includes multiple prospective clinical trials and prospective-retrospective studies. Patients classified as low risk with Oncotype DX have a low risk of recurrence in which avoidance of adjuvant chemotherapy is reasonable (average risk at 10 years, 7%-9%; upper bound of the 95% confidence intervals, 11% to 15%). These results have been demonstrated with stronger study designs for evaluating biomarkers. The evidence is sufficient to determine that the technology results in a meaningful improvement in the net health outcome.

70-Gene Signature (MammaPrint)

For individuals who have early-stage node-negative invasive breast cancer considering adjuvant chemotherapy who receive gene expression profiling with the 70-gene signature (MammaPrint), the evidence includes one study with outcomes in node-negative patients. Although the study showed a low risk of 10-year distant recurrence, it did not derive from high-quality data sources. A recently reported study of clinical utility only reported five year results and may not identify a group with sufficiently low risk. The evidence is insufficient to determine the effects of the technology on health outcomes.

Prosigna

For individuals who have early-stage node-negative invasive breast cancer considering adjuvant chemotherapy who receive gene expression profiling with Prosigna, the evidence includes two prospective-retrospective studies evaluating the prognostic ability of Prosigna. Both studies showed a low absolute risk of distant recurrence in patients with low risk scores. The evidence is sufficient to determine that the technology results in a meaningful improvement in the net health outcome.

*Early-Stage Node-Positive Invasive Breast Cancer*70-Gene Signature (MammaPrint)

For individuals who have early-stage node-positive invasive breast cancer considering adjuvant chemotherapy

who receive gene expression profiling with the 70-gene signature (MammaPrint), the evidence includes prospective-retrospective studies. Existing studies have not reported 10-year distant recurrence outcomes in the patients of interest. The studies are confounded by various factors (e.g., receipt of treatment) or do not report the outcome of interest. The evidence is insufficient to determine the effects of the technology on health outcomes.

Continuation of Tamoxifen Therapy Beyond Five Years

EndoPredict

For individuals who have early-stage node-negative invasive breast cancer free of distant recurrence at five years considering extending tamoxifen treatment who receive gene expression profiling with EndoPredict, the evidence includes one study of archived tissue samples from a previously conducted clinical trial. The study showed low distant recurrence rates in patients classified at low risk with the test. Further studies are necessary to reinforce this finding. It is unclear what an appropriate risk threshold is to consider avoidance of extending tamoxifen treatment. The evidence is insufficient to determine the effect of the technology on health outcomes.

Breast Cancer Index

For individuals who have early-stage node-negative invasive breast cancer free of distant recurrence at five years considering extending tamoxifen treatment who receive gene expression profiling with BCI, the evidence includes two studies of archived tissue samples from previously conducted clinical trials. The study showed low distant recurrence rates in patients classified at low risk with the test. Further studies are necessary to reinforce this finding. It is unclear what an appropriate risk threshold is to consider avoidance of extending tamoxifen treatment. The evidence is insufficient to determine the effect of the technology on health outcomes.

HER2

The evidence for assessment of HER2 status using quantitative total HER2 protein expression and HER2 homodimer measurement in patients who have breast cancer and are undergoing assessment of HER2 status includes validation studies and retrospective analysis of association between levels and survival outcomes. Relevant outcomes are overall survival, disease-specific survival, test accuracy, and test validity. Retrospective analysis using HERmark[®] have shown that the assay may predict a worse response to trastuzumab in certain populations. However, findings have been inconsistent, and no clear association with clinical outcomes has been shown. Additionally, cut points for defining patient groups varied across studies. Clinical utility of the HERmark[®] assay has not been demonstrated, and clinical trials are needed to determine the impact on clinical outcomes of patients stratified by the HERmark[®] assay. The evidence is insufficient to determine the effects of the technology on health outcomes.

Policy

The use of gene expression assays for deciding the value of adjuvant therapy in individuals with intermediate risk breast cancer may be considered **medically necessary** for women with breast cancer meeting all of the following characteristics:

- Unilateral tumor; AND
- hormone receptor-positive (that is estrogen-receptor [ER]-positive or progesterone receptor [PR]-positive); AND
- human epidermal growth factor receptor 2 (HER2) negative; AND
- tumor size 0.6 to one cm with moderate/poor differentiation or unfavorable features OR tumor size larger than one cm; AND

- node negative or lymph nodes with micrometastases less than two mm in size; AND
- who will be treated with adjuvant endocrine therapy, e.g., tamoxifen or aromatase inhibitors; AND
- when the test result will aid the patient in making the decision regarding chemotherapy (i.e., when chemotherapy is a therapeutic option); AND
- when ordered within six months after diagnosis.

The use of the Oncotype DX or Prosigna RT-PCR assays may be considered **medically necessary** if the individual meets the criteria above.

Gene expression assays for deciding the value of adjuvant therapy in individuals with intermediate risk breast cancer and with ER+, Her2-cases with one to three ipsilateral lymph nodes may be considered **medically necessary**.

The 21-gene RT-PCR assay Oncotype DX™ should only be ordered on a tissue specimen obtained during surgical removal of the tumor and after subsequent pathology examination of the tumor has been completed and determined to meet the above criteria (i.e., the test should not be ordered on a preliminary core biopsy). The test should be ordered in the context of a physician-patient discussion regarding risk preferences and when the test result will aid the patient in making decisions regarding chemotherapy.

For patients who otherwise meet the above characteristics but who have multiple ipsilateral primary tumors, a specimen from the tumor with the most aggressive histological characteristics should be submitted for testing. It is not necessary to conduct testing on each tumor; treatment is based on the most aggressive lesion.

All other indications for multigene breast cancer panel assays, including determination of recurrence risk in invasive breast cancer patients with positive lymph nodes or patients with bilateral disease, are considered **investigational**.

The use of a subset of genes from the Oncotype DX or Prosigna RT-PCR assay for predicting recurrence risk in individuals with noninvasive ductal carcinoma in situ to inform treatment planning following excisional surgery is considered **investigational**.

The use of other gene expression assays, including but not limited to MammaPrint® 70-gene signature, Mammostrat® Breast Cancer Test, the Breast Cancer IndexSM, BreastOncPx™, NexCourse® Breast IHC4, BreastPRS™, and EndoPredict™ for any indication is considered **investigational**.

The use of gene expression assays in men with breast cancer is considered **investigational**.

The use of gene expression assays to molecularly subclassify breast cancer is considered **investigational**.

The use of gene expression assays for quantitative assessment of ER, PR, and HER2 overexpression is considered **investigational**.

The use of gene expression assays in tumors that are hormone receptor positive, Her2 Negative, or less than or equal to .5 cm is **investigational**.

Medicare Advantage

The above medical necessity criteria applies for Medicare Advantage members for all gene expression assays in breast tumor tissue tests when no separate Medicare Advantage criteria exists.

The Prosigna breast cancer gene signature assay may be considered **medically necessary** for members that meet the following criteria consistent with the FDA indications for use:

- Post-menopausal female **either**

- ER+, lymph node-negative, stage I or II breast cancer; or
- ER+, lymph node-positive (1-3 positive nodes), stage II breast cancer.

PROSIGNA® Breast Cancer Prognostic Gene Signature Assay is considered **medically necessary** in patients who have undergone surgery in conjunction with locoregional treatment consistent with standard of care, either as:

- A prognostic indicator for distant recurrence-free survival at 10 years in post-menopausal women with Hormone Receptor-Positive (HR+), lymph node-negative, Stage I or II breast cancer to be treated with adjuvant endocrine therapy alone, when used in conjunction with other clinicopathological factors.
- A prognostic indicator for distant recurrence-free survival at 10 years in post-menopausal women with Hormone Receptor-Positive (HR+), lymph node-positive (one to three positive nodes), Stage II breast cancer to be treated with adjuvant endocrine therapy alone, when used in conjunction with other clinicopathological factors. The device is not intended for patients with four or more positive nodes.

The Breast Cancer Index (aka BCI) (bioTheranostics) may be considered **medically necessary** for members that meet the following criteria:

- Post-menopausal female with non-relapsed, ER+ BREAST CANCER, and
- Was lymph node negative, and
- Is completing five (5) years of tamoxifen therapy, and
- Patient must be eligible for consideration of extended endocrine therapy based on published clinical trial data or practice guidelines, and
- Physician or patient is concerned about continuing anti-hormonal therapy because of documented meaningful toxicity or possible significant patient-specific side effects, and
- The test results will be discussed with the patient (including the limitations of the testing method, the risks and benefits of either continuing or stopping the therapy based on the test, and current cancer management guidelines).

Oncology (breast), mRNA, gene expression profiling by real-time RT-PCR of 21 genes, utilizing formalin-fixed paraffin embedded tissue, algorithm reported as recurrence score is considered **medically necessary** to guide therapeutic decision-making in patients with the following findings:

- estrogen-receptor positive, node-negative carcinoma of the breast
- estrogen-receptor positive micrometastases of carcinoma of the breast, and
- estrogen-receptor positive breast carcinoma with one to three positive nodes.

The Oncotype DX DCIS assay is **medically necessary** only when the following clinical conditions are met:

- Pathology (excisional or core biopsy) reveals ductal carcinoma in situ of the breast (no pathological evidence of invasive disease), and
- FFPE specimen with at least 0.5 mm of DCIS length, and
- Patient is a candidate for and is considering breast conserving surgery alone as well as breast conserving surgery combined with adjuvant radiation therapy, and
- Test result will be used to determine treatment choice between surgery alone vs. surgery with radiation therapy, and
- Patient has not received and is not planning on receiving a mastectomy.

For Medicare Advantage the HERmark®, Oncotype DX® Breast and the MammaPrint™ tests may be considered **medically necessary**.

Background

An important part of treatment planning for women with breast cancer involves determining which patients could benefit from adjuvant treatments. For example, for women with early-stage, invasive breast cancer (i.e., cancer extending beyond the basement membrane of the mammary ducts into adjacent tissue), adjuvant cytotoxic chemotherapy consistently provides approximately a 30% relative risk reduction in 10-year breast cancer mortality regardless of prognosis. However, the absolute benefit of chemotherapy depends on the baseline risk of recurrence. Women with the best prognosis have small tumors, are estrogen receptor–positive, and are lymph node–negative. These women have an approximately 15% baseline risk of recurrence; approximately 85% of these patients would be disease-free at 10 years with tamoxifen treatment alone and could avoid the toxicity of chemotherapy, if they could be accurately identified. Conventional risk classifiers (e.g., Adjuvant! Online) estimate recurrence risk by considering criteria such as tumor size, type, grade, and histologic characteristics; hormone receptor status; and lymph node status. Consensus guidelines for defining receptor status exist.¹ However, no single classifier is considered a criterion standard, and several common criteria have qualitative or subjective components that add variability to risk estimates. As a result, more patients are treated with chemotherapy than can benefit. Better predictors of baseline risk could help women’s decision making, some who may prefer to avoid chemotherapy if assured that their risk is low.

In other clinical scenarios involving breast cancer, accurate assessment of prognosis may affect the decision to offer certain treatments. Recently, several groups have identified panels of gene expression markers (“signatures”) that appear to predict the baseline risk of invasive breast cancer recurrence after surgery, radiotherapy, and endocrine therapy (for hormone receptor–positive tumors). Several gene expression tests commercially available in the United States are listed in Table 1. If these panels are more accurate risk predictors than current conventional classifiers, they could be used to aid decision making on adjuvant treatments without greatly affecting disease-free survival and overall survival (OS). This review focuses on gene expression profiling (GEP) panels that have prognostic or predictive ability in individuals with early-stage, invasive breast cancer with known estrogen receptor and progesterone receptor and human epidermal growth factor receptor (HER2) status. The proposed clinical utility of these tests varies depending on the clinical context; these specific indications are discussed in this review:

1. Prognosis and/or prediction of treatment response in patients with node-negative, early-stage, HER2-negative invasive breast cancer who will receive adjuvant hormonal therapy for the purpose of determining whether patients can avoid adjuvant cytotoxic chemotherapy.
2. Prognosis and/or prediction of treatment response in patients with node-positive (one to three nodes), early-stage, HER2-negative invasive breast cancer who will receive adjuvant hormonal therapy for the purpose of determining whether patients can avoid adjuvant cytotoxic chemotherapy.
3. Prognosis and/or prediction of treatment response in patients with ductal carcinoma in situ (DCIS) for the purpose of determining whether patients can avoid radiation therapy.
4. Prognosis and/or prediction of treatment response in patients with node-negative, early-stage, HER2-negative invasive breast cancer, receiving adjuvant hormonal therapy, who have survived without progression to five years postdiagnosis, for the purpose of determining whether patients should continue adjuvant hormonal therapy.

For each of these clinical indications, clinical trials have shown that there is some clinical benefit to receiving the additional therapy under consideration. However, each of the additional treatments has potential adverse effects. If a patient subgroup can be defined that has an extremely low risk of distant recurrence, or a subgroup

can be defined that does not respond to the treatment, then the additional treatment can be forgone with little effect on cancer outcome due to the low risk of poor outcome or lack of response to treatment.

Table 2. Gene Expression Tests Reporting Recurrence Risk for Breast Cancer Considered Herein

Test	Manufacturer	Description
Oncotype DX	Genomic Health (Redwood City, CA)	21-gene RT-PCR
EndoPredict	Sividon Diagnostics (acquired by Myriad [Salt Lake City, UT] in 2016)	12-gene real-time RT-PCR
Breast Cancer Index	bioTheragnostics (San Diego, CA)	Combines MGI and the HOXB13:IL17BR Index using RT-PCR
MammaPrint	Agendia (Amsterdam, The Netherlands)	70-gene DNA microarray
Prosigna	NanoString Technologies (Seattle, WA)	nCounter® digital analysis system based on PAM50 breast cancer intrinsic subtype classifier

MGI: Molecular Grade Index; PAM50: prediction analysis of microarray 50 gene set; RT-PCR: reverse transcriptase polymerase chain reaction.

Additional commercially available tests may provide some prognostic or predictive information for breast cancer. Tests intended to assess estrogen receptor, progesterone receptor, and HER2 status, such as TargetPrint® (Agendia; via quantitative microarray), are outside the scope of this review. In addition, tests that do not provide a specific recurrence risk are outside the scope of this review.

Other commercially available biomarkers are designed to provide information about tumors' molecular subtypes (i.e., luminal A, luminal B, HER2 type, and basal type). Prosigna was initially offered as a molecular subtype test. The Blueprint 80-gene molecular subtyping assay (Agendia) is offered in combination with MammaPrint to augment predictive data about response to chemotherapy.

Many studies have investigated individual biomarkers or combinations of biomarkers that are associated with breast cancer outcomes. Determining which studies constitute sufficient evidence that the test or biomarker is likely to be clinically useful depends on attributes of the test such as its performance and the quality of the study generating the results. Simon et al has described a framework to evaluate prognostic biomarker evidence.² Study designs such as prospective clinical trials or previously conducted clinical trials with archived tumor samples constitute stronger evidence than studies with less planned and systematic patient recruitment and data collection. Randomized trials allow determination of treatment-biomarker interactions that may be clinically important. In some clinical scenarios, demonstration of a treatment-biomarker interaction is not critical, because the decision to withhold chemotherapy in a low-risk group (to avoid chemotherapy-related morbidity) does not require the presence of a biomarker-treatment interaction. The study must generate an absolute estimate of outcome in the patient group of interest that would result in a change in management (e.g., withholding of chemotherapy), and the study must have sufficient precision (narrow confidence intervals). Results of the same test across studies should show consistency of results and more than one study demonstrating the desired result should be available. Simon has proposed that at least two Simon category B studies showing results consistent with clinical utility are necessary to demonstrate adequate evidence of a biomarker.²

The main outcome of interest to this review is 10-year distant recurrence-free survival. Distant recurrence is a hallmark of advanced breast cancer and thus more informative of OS than disease-free survival. Disease-free survival also includes local recurrence, which has a much better treatment prognosis than distant disease. For one of the indications in this review, the main outcome of interest is 10-year distant recurrence-free survival conditional on recurrence-free survival for five years. There is no definitive threshold for an acceptable trade-off of distant recurrence risk for avoidance of treatment toxicity and inconvenience that is derived from empirical evidence on patient preferences. While some studies have shown that patients are willing to accept intensive chemotherapy for even a small chance of benefit, individual patients will vary in their preferences and tolerance for adverse effects.

HER2

The human epidermal growth factor receptor (HER) family of receptor tyrosine kinases (EGFR/HER1, ErbB2/HER2, ErbB3/HER3, ErbB4/HER4) plays a major role in the pathogenesis of many solid tumors. In approximately 25% to 30% of breast cancers, overexpression of HER2 has been linked to shorter disease-free (DFS) and overall survival (OS), lack of responsiveness to tamoxifen antiestrogen therapy, and altered responsiveness to a variety of cytotoxic chemotherapy regimens.

Trastuzumab, a monoclonal antibody directed at the extracellular domain of HER2 has offered significant DFS and OS advantages in the metastatic and adjuvant settings in HER2-overexpressing patients, although not all patients respond. Fewer than 50% of patients with metastatic HER2-positive breast cancer show initial benefit from trastuzumab treatment, and many of those eventually develop resistance.¹

Current methodologies for the selection of HER2-positive patients include immunohistochemistry (IHC) to detect HER2 protein overexpression, and fluorescence in situ hybridization (FISH) to detect HER2 gene amplification. However, controversy still exists regarding the accuracy, reliability, and interobserver variability of these assay methods. IHC provides a semiquantitative measure of protein levels (scored as 0, 1+, 2+, 3+) and the interpretation may be subjective. FISH is a quantitative measurement of gene amplification, in which the HER2 gene copy number is counted. However, FISH, which is considered to be more quantitative analytically, is not always representative of protein expression, and multiple studies have failed to demonstrate a relationship between HER2 gene copy number and response to trastuzumab. Whereas patients who overexpress HER2 protein (IHC) or show evidence of HER2 gene amplification (FISH) have been shown to experience better outcomes on trastuzumab than those scored negative by those assays, differences in the degree of expression or amplification by these methods have generally not been shown to discriminate between groups with different outcomes. IHC and FISH testing may be affected by interlaboratory variability, and neither test provides quantitative data that reflect the activation state of signaling pathways in tumors, which may limit their utility in patient selection.² Most laboratories in North America and Europe use IHC to determine HER2 protein status, with equivocal category results (2+) confirmed by FISH (or more recently by chromogenic in situ hybridization [CISH]).

Normally, HER2 activates signaling pathways by dimerizing with ligand-bound EGFR-family members such as HER1 and HER3. A HER2 ligand has not been identified, but overexpressed HER2 is constitutively active. When HER2 is pathologically overexpressed, the receptor may homodimerize and activate signaling cascades in the absence of the normal regulatory control imposed by the requirement for ligand binding of its heterodimerization partners.

A novel assay (HERmark[®] Breast Cancer Assay; Monogram Biosciences, South San Francisco, CA) was developed to quantify total HER2 protein expression (H2T) and HER2 homodimers (H2D) in formalin-fixed, paraffin-embedded tissue samples.

Regulatory Status

Clinical laboratories may develop and validate tests in-house and market them as a laboratory service; laboratory-developed tests (LDTs) must meet the general regulatory standards of the Clinical Laboratory Improvement Amendments (CLIA). Oncotype DX[®] and other tests listed herein are available under the auspices of CLIA. Laboratories that offer LDTs must be licensed by CLIA for high-complexity testing. To date, the U.S. Food and Drug Administration (FDA) has chosen not to require any regulatory review of this testing.

In February 2007, MammaPrint[®] (Agendia) was cleared for marketing by FDA through the 510(k) process. In January 2015, MammaPrint[®] was cleared for marketing by FDA through the 510(k) process for use in fresh-frozen, paraffin-embedded breast cancer tissue.

In September 2013, Prosigna[®] was cleared for marketing by FDA through the 510(k) process. FDA determined that Prosigna[®] was substantially equivalent to MammaPrint[®].

Product Code: NYI.

Services that are the subject of a clinical trial do not meet our Technology Assessment Protocol criteria and are considered investigational. *For explanation of experimental and investigational, please refer to the Technology Assessment Protocol.*

It is expected that only appropriate and medically necessary services will be rendered. We reserve the right to conduct prepayment and postpayment reviews to assess the medical appropriateness of the above-referenced procedures. **Some of this protocol may not pertain to the patients you provide care to, as it may relate to products that are not available in your geographic area.**

References

We are not responsible for the continuing viability of web site addresses that may be listed in any references below.

1. Wolff AC, Hammond ME, Hicks DG, et al. Recommendations for human epidermal growth factor receptor 2 testing in breast cancer: american society of clinical oncology/college of american pathologists clinical practice guideline update. *J Clin Oncol.* Nov 1 2013; 31(31):3997-4013. PMID 24101045
2. Simon RM, Paik S, Hayes DF. Use of archived specimens in evaluation of prognostic and predictive biomarkers. *J Natl Cancer Inst.* Nov 4 2009; 101(21):1446-1452. PMID 19815849
3. Slevin ML, Stubbs L, Plant HJ, et al. Attitudes to chemotherapy: comparing views of patients with cancer with those of doctors, nurses, and general public. *BMJ.* Jun 2 1990; 300(6737):1458-1460. PMID 2379006
4. Ravdin PM, Siminoff IA, Harvey JA. Survey of breast cancer patients concerning their knowledge and expectations of adjuvant therapy. *J Clin Oncol.* Feb 1998; 16(2):515-521. PMID 9469335
5. Duric VM, Stockler MR, Heritier S, et al. Patients' preferences for adjuvant chemotherapy in early breast cancer: what makes AC and CMF worthwhile now? *Ann Oncol.* Nov 2005; 16(11):1786-1794. PMID 16126738
6. Stiggelbout AM, de Haes JC, van de Velde CJ. Adjuvant chemotherapy in node negative breast cancer: patterns of use and oncologists' preferences. *Ann Oncol.* May 2000; 11(5):631-633. PMID 10907961
7. Buus R, Sestak I, Kronenwett R, et al. Comparison of EndoPredict and EPclin with Oncotype DX Recurrence Score for prediction of risk of distant recurrence after endocrine therapy. *J Natl Cancer Inst.* Nov 2016; 108(11). PMID 27400969
8. Paik S, Shak S, Tang G, et al. A multigene assay to predict recurrence of tamoxifen-treated, node-negative breast cancer. *N Engl J Med.* Dec 30 2004; 351(27):2817-2826. PMID 15591335
9. Paik S, Tang G, Shak S, et al. Gene expression and benefit of chemotherapy in women with node-negative, estrogen receptor-positive breast cancer. *J Clin Oncol.* Aug 10 2006; 24(23):3726-3734. PMID 16720680
10. Tang G, Shak S, Paik S, et al. Comparison of the prognostic and predictive utilities of the 21-gene Recurrence Score assay and Adjuvant! for women with node-negative, ER-positive breast cancer: results from NSABP B-14 and NSABP B-20. *Breast Cancer Res Treat.* May 2011; 127(1):133-142. PMID 21221771
11. Sparano JA, Gray RJ, Makower DF, et al. Prospective validation of a 21-gene expression assay in breast cancer. *N Engl J Med.* Nov 19 2015; 373(21):2005-2014. PMID 26412349
12. Filipits M, Rudaš M, Jakesz R, et al. A new molecular predictor of distant recurrence in ER-positive, HER2-negative breast cancer adds independent information to conventional clinical risk factors. *Clin Cancer Res.* Sep 15 2011; 17(18):6012-6020. PMID 21807638

13. Sgroi DC, Sestak I, Cuzick J, et al. Prediction of late distant recurrence in patients with oestrogen-receptor-positive breast cancer: a prospective comparison of the breast-cancer index (BCI) assay, 21-gene recurrence score, and IHC4 in the TransATAC study population. *Lancet Oncol.* Oct 2013; 14(11):1067-1076. PMID 24035531
14. Zhang Y, Schnabel CA, Schroeder BE, et al. Breast cancer index identifies early-stage estrogen receptor-positive breast cancer patients at risk for early- and late-distant recurrence. *Clin Cancer Res.* Aug 1 2013; 19(15):4196-4205. PMID 23757354
15. Bueno-de-Mesquita JM, Sonke GS, van de Vijver MJ, et al. Additional value and potential use of the 70-gene prognosis signature in node-negative breast cancer in daily clinical practice. *Ann Oncol.* Sep 2011; 22(9):2021-2030. PMID 19955335
16. Cardoso F, van't Veer LJ, Bogaerts J, et al. 70-gene signature as an aid to treatment decisions in early-stage breast cancer. *N Engl J Med.* Aug 25 2016; 375(8):717-729. PMID 27557300
17. Dowsett M, Sestak I, Lopez-Knowles E, et al. Comparison of PAM50 risk of recurrence score with Oncotype DX and IHC4 for predicting risk of distant recurrence after endocrine therapy. *J Clin Oncol.* Aug 1 2013; 31(22):2783-2790. PMID 23816962
18. Gnant M, Filipits M, Greil R, et al. Predicting distant recurrence in receptor-positive breast cancer patients with limited clinicopathological risk: using the PAM50 Risk of Recurrence score in 1478 postmenopausal patients of the ABCSG-8 trial treated with adjuvant endocrine therapy alone. *Ann Oncol.* Feb 2014; 25(2):339-345. PMID 24347518
19. Albain KS, Barlow WE, Shak S, et al. Prognostic and predictive value of the 21-gene recurrence score assay in postmenopausal women with node-positive, oestrogen-receptor-positive breast cancer on chemotherapy: a retrospective analysis of a randomised trial. *Lancet Oncol.* Jan 2010; 11(1):55-65. PMID 20005174
20. Dowsett M, Cuzick J, Wale C, et al. Prediction of risk of distant recurrence using the 21-gene recurrence score in node-negative and node-positive postmenopausal patients with breast cancer treated with anastrozole or tamoxifen: a TransATAC study. *J Clin Oncol.* Apr 10 2010; 28(11):1829-1834. PMID 20212256
21. Goldstein LJ, Gray R, Badve S, et al. Prognostic utility of the 21-gene assay in hormone receptor-positive operable breast cancer compared with classical clinicopathologic features. *J Clin Oncol.* Sep 1 2008; 26(25):4063-4071. PMID 18678838
22. Gluz O, Nitz UA, Christgen M, et al. West German Study Group Phase III PlanB Trial: first prospective outcome data for the 21-gene recurrence score assay and concordance of prognostic markers by central and local pathology assessment. *J Clin Oncol.* Jul 10 2016; 34(20):2341-2349. PMID 26926676
23. Chang JC, Makris A, Gutierrez MC, et al. Gene expression patterns in formalin-fixed, paraffin-embedded core biopsies predict docetaxel chemosensitivity in breast cancer patients. *Breast Cancer Res Treat.* Mar 2008; 108(2):233-240. PMID 17468949
24. Gianni L, Zambetti M, Clark K, et al. Gene expression profiles in paraffin-embedded core biopsy tissue predict response to chemotherapy in women with locally advanced breast cancer. *J Clin Oncol.* Oct 10 2005; 23(29):7265-7277. PMID 16145055
25. Mook S, Schmidt MK, Viale G, et al. The 70-gene prognosis-signature predicts disease outcome in breast cancer patients with 1-3 positive lymph nodes in an independent validation study. *Breast Cancer Res Treat.* Jul 2009; 116(2):295-302. PMID 18661261
26. van de Vijver MJ, He YD, van't Veer LJ, et al. A gene-expression signature as a predictor of survival in breast cancer. *N Engl J Med.* Dec 19 2002; 347(25):1999-2009. PMID 12490681
27. Esserman LJ, Berry DA, Cheang MC, et al. Chemotherapy response and recurrence-free survival in neoadjuvant breast cancer depends on biomarker profiles: results from the I-SPY 1 TRIAL (CALGB 150007/150012; ACRIN 6657). *Breast Cancer Res Treat.* Apr 2012; 132(3):1049-1062. PMID 22198468
28. Saghathian M, Mook S, Pruneri G, et al. Additional prognostic value of the 70-gene signature (MammaPrint((R))) among breast cancer patients with 4-9 positive lymph nodes. *Breast.* Oct 2013; 22(5):682-690. PMID 23347730

29. Solin LJ, Gray R, Baehner FL, et al. A multigene expression assay to predict local recurrence risk for ductal carcinoma in situ of the breast. *J Natl Cancer Inst.* May 15, 2013; 105(10):701-710. PMID 23641039
30. Rakovitch E, Nofech-Mozes S, Hanna W, et al. A population-based validation study of the DCIS Score predicting recurrence risk in individuals treated by breast-conserving surgery alone. *Breast Cancer Res Treat.* Jul 2015; 152(2):389-398. PMID 26119102
31. Dubsy P, Brase JC, Jakesz R, et al. The EndoPredict score provides prognostic information on late distant metastases in ER+/HER2-breast cancer patients. *Br J Cancer.* Dec 10 2013; 109(12):2959-2964. PMID 24157828
32. Filipits M, Nielsen TO, Rudas M, et al. The PAM50 risk-of-recurrence score predicts risk for late distant recurrence after endocrine therapy in postmenopausal women with endocrine-responsive early breast cancer. *Clin Cancer Res.* Mar 1 2014; 20(5):1298-1305. PMID 24520097
33. Sestak I, Dowsett M, Zabaglo L, et al. Factors predicting late recurrence for estrogen receptor-positive breast cancer. *J Natl Cancer Inst.* Oct 2 2013; 105(19):1504-1511. PMID 24029245
34. Hornberger J, Alvarado MD, Rebecca C, et al. Clinical validity/utility, change in practice patterns, and economic implications of risk stratifiers to predict outcomes for early-stage breast cancer: a systematic review. *J Natl Cancer Inst.* Jul 18 2012; 104(14):1068-1079. PMID 22767204
35. Fan C, Oh DS, Wessels L, et al. Concordance among gene-expression-based predictors for breast cancer. *N Engl J Med.* Aug 10 2006; 355(6):560-569. PMID 16899776
36. Espinosa E, Vara JA, Redondo A, et al. Breast cancer prognosis determined by gene expression profiling: a quantitative reverse transcriptase polymerase chain reaction study. *J Clin Oncol.* Oct 10 2005; 23(29):7278-7285. PMID 16129846
37. Kelly CM, Bernard PS, Krishnamurthy S, et al. Agreement in risk prediction between the 21-gene recurrence score assay (Oncotype DX(R)) and the PAM50 Breast Cancer Intrinsic Classifier in early-stage estrogen receptor-positive breast cancer. *Oncologist.* 2012; 17(4):492-498. PMID 22418568
38. Prat A, Parker JS, Fan C, et al. Concordance among gene expression-based predictors for ER-positive breast cancer treated with adjuvant tamoxifen. *Ann Oncol.* Nov 2012; 23(11):2866-2873. PMID 22532584
39. Badve SS, Baehner FL, Gray RP, et al. Estrogen- and progesterone-receptor status in ECOG 2197: comparison of immunohistochemistry by local and central laboratories and quantitative reverse transcription polymerase chain reaction by central laboratory. *J Clin Oncol.* May 20, 2008; 26(15):2473-2481. PMID 18487567
40. Khoury T, Yan L, Liu S, et al. Oncotype DX RT-qPCR assay for ER and PR correlation with IHC: a study of 3 different clones. *Appl Immunohistochem Mol Morphol.* Mar 2015; 23(3):178-187. PMID 24992175
41. Drukker CA, Elias SG, Nijenhuis MV, et al. Gene expression profiling to predict the risk of locoregional recurrence in breast cancer: a pooled analysis. *Breast Cancer Res Treat.* Dec 2014; 148(3):599-613. PMID 25414025
42. Fitzal F, Filipits M, Rudas M, et al. The genomic expression test EndoPredict is a prognostic tool for identifying risk of local recurrence in postmenopausal endocrine receptor-positive, her2neu-negative breast cancer patients randomised within the prospective ABCSG 8 trial. *Br J Cancer.* Apr 14 2015; 112(8):1405-1410. PMID 25867274
43. National Comprehensive Cancer Network (NCCN). NCCN Clinical Practice Guidelines in Oncology: Breast Cancer. Version 2.2016. http://www.nccn.org/professionals/physician_gls/pdf/breast.pdf. Accessed July, 2016.
44. Harris LN, Ismaila N, McShane LM, et al. Use of biomarkers to guide decisions on adjuvant systemic therapy for women with early-stage invasive breast cancer: American Society of Clinical Oncology Clinical Practice Guideline. *J Clin Oncol.* Apr 1 2016; 34(10):1134-1150. PMID 26858339
45. Coates AS, Winer EP, Goldhirsch A, et al. Tailoring therapies--improving the management of early breast cancer: St Gallen International Expert Consensus on the Primary Therapy of Early Breast Cancer 2015. *Ann Oncol.* Aug 2015; 26(8):1533-1546. PMID 25939896
46. Centers for Medicare & Medicaid Services. Local Coverage Determination (LCD): MoIDX: Breast Cancer IndexSM Genetic Assay (L35631), effective 10/01/2015. <https://www.cms.gov/medicare-coverage->

database/details/lcd-

details.aspx?LCDId=35631&ver=9&CoverageSelection=Local&ArticleType=All&PolicyType=Final&s=All&KeyWord=Breast+Cancer+Index&KeyWordLookUp=Title&KeyWordSearchType=And&bc=gAAAAACAAAAAAA%3d%3d&. Accessed November 7, 2016.

47. Zujewski JA, Kamin L. Trial assessing individualized options for treatment for breast cancer: the TAILORx trial. *Future Oncol.* Oct 2008; 4(5):603-610. PMID 1892211
48. Eiermann W, Rezai M, et al. The 21-gene recurrence score assay impacts adjuvant therapy recommendations for ER-positive, node-negative and node-positive early breast cancer resulting in a risk-adapted change in chemotherapy use. *Annals of Oncology.* March 2013; 24(3):618–624.
49. Gyorkffy B, Hatzis C, et al. Multigene prognostic tests in breast cancer: past, present, future. *Breast Cancer Research.* 2015 17:11.
50. Van Poznak C, Somerfield MR, et al. Use of Biomarkers to Guide Decisions on Systemic Therapy for Women With Metastatic Breast Cancer: American Society of Clinical Oncology Clinical Practice Guideline. *Journal of Clinical Oncology.* 2015. www.jco.org
51. DeFazio-Eli L, Strommen K, Dao-Pick T, et al. Quantitative assays for the measurement of HER1-HER2 heterodimerization and phosphorylation in cell lines and breast tumors: applications for diagnostics and targeted drug mechanism of action. *Breast Cancer Res.* 2011; 13(2):R44. PMID 21496232
52. Shi Y, Huang W, Tan Y, et al. A novel proximity assay for the detection of proteins and protein complexes: quantitation of HER1 and HER2 total protein expression and homodimerization in formalin-fixed, paraffin embedded cell lines and breast cancer tissue. *Diagn Mol Pathol.* 2009; 18(1):11-21.
53. Huang W, Reinholz M, Weidler J, et al. Comparison of central HER2 testing with quantitative total HER2 expression and HER2 homodimer measurements using a novel proximity-based assay. *Am J Clin Pathol.* 2010; 134(2):303-311.
54. Wolff AC, Hammond ME, Schwartz JN, et al. American Society of Clinical Oncology/College of American Pathologists guideline recommendations for human epidermal growth factor receptor 2 testing in breast cancer. *Arch Pathol Lab Med.* 2007; 131(1):18-43.
55. Bates M, Sperinde J, Köstler WJ, et al. Identification of a subpopulation of metastatic breast cancer patients with very high HER2 expression levels and possible resistance to trastuzumab. *Ann Oncol.* 2011; 22(9):2014-2020.
56. Joensuu H, Sperinde J, Leinonen M, et al. Very high quantitative tumor HER2 content and outcome in early breast cancer. *Ann Oncol.* 2011; 22(9):2007-2013.
57. Toi M, Sperinde J, Huang W, et al. Differential survival following trastuzumab treatment based on quantitative HER2 expression and HER2 homodimers in a clinic-based cohort of patients with metastatic breast cancer. *BMC Cancer.* 2010; 10:56. PMID 20178580
58. Lipton A, Köstler WJ, Leitzel K, et al. Quantitative HER2 protein levels predict outcome in fluorescence in situ hybridization-positive patients with metastatic breast cancer treated with trastuzumab. *Cancer.* 2010; 116(22):5168-5178.
59. Lipton A, Goodman L, Leitzel K, et al. HER3, p95HER2, and HER2 protein expression levels define multiple subtypes of HER2-positive metastatic breast cancer. *Breast Cancer Res Treat.* Aug 20 2013. PMID 23959396
60. Duchnowska R, Sperinde J, Chenna A, et al. Quantitative measurements of tumoral p95HER2 protein expression in metastatic breast cancer patients treated with trastuzumab: independent validation of the p95HER2 clinical cutoff. *Clin Cancer Res.* May 15, 2014; 20(10):2805-2813. PMID 24668646
61. Han SW, Cha Y, Paquet A, et al. Correlation of HER2, p95HER2 and HER3 expression and treatment outcome of lapatinib plus capecitabine in her2-positive metastatic breast cancer. *PLoS One.* 2012; 7(7):e39943. PMID 22848366
62. Duchnowska R, Biernat W, Szostakiewicz B, et al. Correlation between quantitative HER-2 protein expression and risk for brain metastases in HER-2+ advanced breast cancer patients receiving trastuzumab-containing therapy. *Oncologist.* 2012; 17(1):26-35. PMID 22234627

63. Barros FF, Abdel-Fatah TM, Moseley P, et al. Characterisation of HER heterodimers in breast cancer using in situ proximity ligation assay. *Breast Cancer Res Treat.* Apr 2014; 144(2):273-285. PMID 24557338
64. Camp RL, Dolled-Filhart M, Rimm DL. X-tile: a new bio-informatics tool for biomarker assessment and outcome based cut-point optimization. *Clin Cancer Res.* Nov 1 2004; 10(21):7252-7259. PMID 15534099
65. Green AR, Barros FF, Abdel-Fatah TM, et al. HER2/HER3 heterodimers and p21 expression are capable of predicting adjuvant trastuzumab response in HER2+ breast cancer. *Breast Cancer Res Treat.* May 2014; 145(1):33-44. PMID 24706169
66. National Comprehensive Cancer Network (NCCN). Clinical practice guidelines in oncology: breast cancer, version 3.2015. http://www.nccn.org/professionals/physician_gls/pdf/breast.pdf. Accessed November 12, 2015.
67. Noridian Healthcare Solutions, LLC, (Jurisdiction - California - Entire State, American Samoa, Guam, Hawaii, Northern Mariana Islands, Nevada) Local Coverage Determination (LCD): MoIDX: Molecular Diagnostic Tests (MDT) (L35160), Revision Effective Date for services performed on or after 01/01/2017.
68. National Government Services, Inc. Local Coverage Determination (LCD): Molecular Pathology Procedures (L35000), Revision Effective Date for services performed on or after 01/01/2017.
69. Noridian Healthcare Solutions, LLC, (Jurisdiction - California - Entire State, American Samoa, Guam, Hawaii, Northern Mariana Islands, Nevada) PROPOSED/DRAFT Local Coverage Determination (LCD): MoIDX - CDD: ONCOTYPE DX[®] Breast Cancer for DCIS (Genomic Health[™]) (DL36941), Comment Period Start Date 10/06/2016, Comment Period End Date 12/30/2016, Released to Final LCD Date 01/17/2017.
70. Palmetto GBA (North Carolina, South Carolina, West Virginia, Virginia) Local Coverage Determination (LCD): MoIDX: BREAST CANCER Assay: Prosigna (L36125), Revision Effective Date for services performed on or after 10/17/2016.
71. Palmetto GBA (North Carolina, South Carolina, West Virginia, Virginia) Local Coverage Determination (LCD): MoIDX: BREAST CANCER IndexSM GENETIC Assay (L35631), Revision Effective Date for services performed on or after 01/25/2016.