

Genetic Testing: Hereditary Breast and Ovarian Cancer

MEDICAL POLICY NUMBER: 143

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INSTRUCTIONS FOR USE: Company Medical Policies serve as guidance for the administration of plan benefits. Medical policies do not constitute medical advice nor a guarantee of coverage. Company Medical Policies are reviewed annually and are based upon published, peer-reviewed scientific evidence and evidence-based clinical practice guidelines that are available as of the last policy update. The Company reserves the right to determine the application of medical policies and make revisions to medical policies at any time. The scope and availability of all plan benefits are determined in accordance with the applicable coverage agreement. Any conflict or variance between the terms of the coverage agreement and Company Medical Policy will be resolved in favor of the coverage agreement. Coverage decisions are made on the basis of individualized determinations of medical necessity and the experimental or investigational character of the treatment in the individual case. In cases where medical necessity is not established by policy for specific treatment modalities, evidence not previously considered regarding the efficacy of the modality that is presented shall be given consideration to determine if the policy represents current standards of care.

SCOPE: Providence Health Plan, Providence Health Assurance, Providence Plan Partners, and Ayin Health Solutions as applicable (referred to individually as “Company” and collectively as “Companies”).

PLAN PRODUCT AND BENEFIT APPLICATION

Commercial

Medicaid/OHP*

Medicare**

*Medicaid/OHP Members

Oregon: Services requested for Oregon Health Plan (OHP) members follow the OHP Prioritized List and Oregon Administrative Rules (OARs) as the primary resource for coverage determinations. Medical policy criteria below may be applied when there are no criteria available in the OARs and the OHP Prioritized List.

**Medicare Members

This *Company* policy may be applied to Medicare Plan members only when directed by a separate *Medicare* policy. Note that investigational services are considered “**not medically necessary**” for Medicare members.

COVERAGE CRITERIA

Note:

- Both maternal and paternal family histories must be considered separately when identifying families with an increased risk of breast and/or ovarian cancer.

Policy Criteria Links

- [Genetic Testing: Family with a Known Breast and/or Ovarian Cancer Mutation](#)
- [Genetic Testing: Personal History of Cancer](#)
- [Genetic Testing: No Personal History of Cancer](#)
- [Genetic Testing: Non-Covered Testing](#)

Genetic Testing: Family with a Known Breast and/or Ovarian Cancer Mutation

- I. Genetic testing for a [hereditary breast and/or ovarian cancer gene mutation](#) may be considered **medically necessary** when **all** of the following criteria (A-C) are met:
 - A. Genetic Counseling general criteria have been met; **and**
 - B. The patient is from a family with a known pathogenic/likely pathogenic variant in a [hereditary breast and/or ovarian cancer gene](#); **and**
 - C. Gene testing is specific to the familial pathogenic/likely pathogenic variant.

Genetic Testing: Personal History of Cancer

- II. Genetic testing for [hereditary breast and/or ovarian cancer gene mutation\(s\)](#) may be considered **medically necessary** when **both** of the following criteria (A. and B.) are met:
 - A. Genetic Counseling general criteria have been met; **and**
 - B. Any **one or more** of the following criteria are met:
 1. The patient has a personal history of breast cancer (including invasive and ductal

carcinoma *in situ*) and any **one or more** of the following criteria are met:

- a. Diagnosed with breast cancer at age 45 or younger; **or**
- b. Diagnosed with breast cancer at age 46-50 with **any** of the following:
 - i. Unknown or limited family history
 - ii. Multiple primary breast cancers (synchronous or metachronous); **or**
 - iii. 1 or more [close blood relative](#) with breast, ovarian, pancreatic, or prostate cancer at any age; **or**
- c. Diagnosed with breast cancer at age 51 or older with 1 or more [close blood relative](#) with **any** of the following:
 - i. Breast cancer at age 50 years or younger or male breast cancer at any age
 - ii. Ovarian or pancreatic cancer at any age
 - iii. Metastatic, intraductal/cribriform histology, or high- or very-high risk group prostate cancer at any age
 - iv. 3 or more total diagnoses of breast cancer in patient and/or [close blood relatives](#)
 - v. 2 or more [close blood relatives](#) with either breast or prostate cancer (any grade) at any age, **or**
- d. Diagnosed with breast cancer **at any age and any one** of the following criteria are met:
 - i. To aid in systemic treatment decisions using PARP inhibitors for breast cancer in the metastatic setting
 - ii. To aid in adjuvant treatment decisions with olaparib for high-risk, HER-2 negative breast cancer
 - iii. Triple-negative breast cancer
 - iv. Lobular breast cancer with personal or family history of diffuse gastric cancer
 - v. 1 or more [close blood relatives](#) with male breast cancer
 - vi. Ethnicity associated with higher mutation frequency or increased risk of founder mutation (e.g., Ashkenazi Jewish, Norwegian, Dutch, or Icelandic descent); **or**

Note: A limited family history occurs when a 1st, 2nd and 3rd generation pedigree cannot be obtained or when there are too few family members in each generation to reasonably see a pattern of inheritance.

2. The patient has ovarian cancer (includes fallopian tube and primary peritoneal cancers); **or**
3. The patient has male breast cancer; **or**
4. The patient has pancreatic cancer; **or**
5. The patient has metastatic prostate cancer (biopsy-proven and/or with radiologic evidence and includes distant metastases and regional bed or nodes); **or**
6. The patient has high-grade (Gleason score ≥ 7) prostate cancer at any age with **any** of the following:
 - a. 1 or more [close blood relative](#) with any of the following:
 - i. Breast cancer at age 50 or younger; **or**
 - ii. Invasive ovarian cancer (including fallopian tube or primary peritoneal cancer) at any age; **or**
 - iii. Pancreatic cancer at any age; **or**

- iv. Metastatic prostate cancer at any age (biopsy-proven and/or with radiologic evidence and includes distant metastases and regional bed or nodes); **or**
- b. Is of Ashkenazi Jewish decent; **or**
- 7. The patient has a personal and/or family history of **3 or more** of the following diagnosis (can include multiple primary cancers in the same individual):
 - a. Breast cancer
 - b. Pancreatic cancer
 - c. Prostate cancer (Gleason score ≥ 7 or metastatic)
 - d. Melanoma
 - e. Sarcoma
 - f. Adrenocortical carcinoma
 - g. Brain tumors
 - h. Leukemia
 - i. Diffuse gastric cancer
 - j. Colon cancer
 - k. Endometrial cancer
 - l. Thyroid cancer
 - m. Kidney cancer
 - n. [Dermatological manifestations](#) of Cowden syndrome
 - o. Macrocephaly
 - p. Gastrointestinal cancer or hamartomatous polyps of GI tract
 - q. Ovarian sex chord tumors
 - r. Testicular sertoli cell tumors
 - s. [Childhood skin pigmentation](#) indicative of Peutz-Jeghers syndrome
- 8. A known pathogenic/likely pathogenic variant in a [hereditary breast and/or ovarian cancer gene](#) detected by tumor profiling on any tumor type in the absence of germline variant analysis.

Genetic Testing: No Personal History of Cancer

- III. Genetic testing for [hereditary breast and/or ovarian cancer gene mutation\(s\)](#) may be considered **medically necessary** when the Genetic Counseling general criteria have been met **and** any one or more of the following criteria (A or B) are met:
 - C. The patient has a first- or second- degree blood relative who meets criterion II.B. above; **or**
 - D. The patient has a first- or second- degree blood relative with at least two [close blood relatives](#) with breast cancer primaries on the same side of the family with at least one diagnosed at age 50 or younger.

Notes:

- Criterion III. may apply to an affected third-degree relative if related through two male relatives (e.g., paternal grandfather's mother or sister).
- When possible, family members with cancer associated with inherited breast and/or ovarian cancer should be tested first, prior to testing unaffected members. Documentation that testing an unaffected member has significant limitations on interpreting test results. Pre-test counseling notes should document the reason why none of these members can be tested prior to testing an unaffected member.

Genetic Testing: Non-covered Testing

- IV. Initial or repeat genetic testing for [hereditary breast and/or ovarian cancer gene mutation\(s\)](#) is considered **not medically necessary** when any of the criteria (I.-III.) above are not met.
- V. Genetic testing for [hereditary breast and/or ovarian cancer gene mutation\(s\)](#) is considered **not medically necessary** in patients who have received an allogeneic bone marrow transplant if only blood or buccal samples are available. Fibroblast culture is the only acceptable source of DNA for these patients.

Note: Per the NCCN, testing from blood and buccal sources may be unreliable due to donor-derived DNA.¹

- VI. Genetic panel testing which include additional genes not identified as [hereditary breast and/or ovarian cancer gene mutation\(s\)](#) is considered **not medically necessary** as a method of cancer risk assessment, unless requested as part of a multi-gene panel that meets criteria per the Medical Policy, "[Next Generation Sequencing for Cancer \(Company\)](#)." Examples of panels considered not medically necessary include, but are not limited to, the following :
- A. +RNAinsight for BreastNext (Ambry Genetics)
 - B. +RNAinsight for OvaNext (Ambry Genetics)
 - C. Breast/Gyn Cancer (GeneDx)
 - D. BreastNext (Sema4/Ambry Genetics)
 - E. myRisk Hereditary Cancer (Myriad Genetics)
 - F. OvaNext (Sema4/Ambry Genetics)
 - G. VistaSeq Breast Cancer Panel (Labcorp)
 - H. VistaSeq Breast and Gyn Panel (Labcorp)
 - I. VistaSeq Hereditary Cancer Panel (Labcorp)
 - J. Common Hereditary Cancers Panel (Invitae)

Link to [Evidence Summary](#)

POLICY CROSS REFERENCES

- [Genetic Counseling](#), MP316
- [Genetic Testing: Non-Covered Genetic Panel Tests](#), MP213
- [Next Generation Sequencing for Cancer](#), MP352

The full Company portfolio of current Medical Policies is available online and can be [accessed here](#).

POLICY GUIDELINES

Hereditary Breast and/or Ovarian Cancer Gene Mutations

The following genes are associated with an increased risk of breast and/or ovarian cancer (not all inclusive):¹

- ATM
- BRIP1
- MLH1
- NF1
- RAD51C
- BARD1
- CDH1
- MSH2
- PALB2
- RAD51D
- BRCA1
- CHEK2
- MSH6
- PMS2
- STK11
- BRCA2
- EPCAM
- NBN
- PTEN
- TP53

Note: BART testing for large rearrangements (e.g., large deletions and/or duplications) may be included and is covered as a component of BRCA 1 and/or 2 testing when medical necessity criteria are met.

DOCUMENTATION REQUIREMENTS

In order to determine the clinical utility of a genetic test, the following documentation must be provided at the time of the request. Failure to submit complete documentation may affect the outcome of the review.

- Specific gene, trade or proprietary name of the test, or if a custom-built test, include every gene(s) and/or component of the test
- Name of laboratory where the testing is being conducted or was conducted
- Clinical notes to include the following:
 - Documentation of genetic counseling as required in the [Genetic Counseling policy](#) which includes how test results will impact clinical decision making
 - If relevant, pre-test counseling notes should document the reason why an affected family member cannot be tested prior to testing an unaffected member
 - Reason (indication) for performing test, including the suspected condition
 - Existing signs and/or symptoms related to reason for current test request
 - Prior test/laboratory results related to reason for current test request
 - Family history, if applicable
 - How results from current test request will impact clinical decision making
- All relevant CPT/HCPCS codes billed

DEFINITIONS

Genetic Panel Testing

The clinical utility of testing has been established for all 19 of the genes listed above. Therefore, when medical necessity criteria are met, genetic panel testing for hereditary breast and ovarian cancer risk may include one or more of **any combination** of the genes listed in this section above. For example, genetic panels, such as the INVITAE Hereditary Breast and Ovarian Cancer Syndrome Panel, which only include select genes from the list above, may be considered medically necessary when criteria are met.

Close Blood Relatives

The National Comprehensive Cancer Network (NCCN) defines close blood relative as first- (parents, siblings and children), second- (grandparents, aunts, uncles, nieces and nephews, grandchildren and

half-siblings), and third degree-relatives (great-grandparents, great-aunts, great uncles, great grandchildren and first cousins) on same side of family.¹

Dermatological manifestations

For a complete list of dermatological manifestations of Cowden syndrome, please see the current NCCN guidelines on "[Genetic/Familial High-Risk Assessment: Breast and Ovarian.](#)"¹

Childhood Skin Pigmentation Indications

For a complete list of childhood skin pigmentation indicative of Peutz-Jeghers syndrome, please see the current NCCN guidelines on "[Genetic/Familial High-Risk Assessment: Colorectal.](#)"²

BACKGROUND

Clinical Utility of Genetic Testing

Establishing the clinical utility of any test is a key component in determining its ultimate usefulness. Clinical utility may be established when published evidence demonstrates test results can be used to:

1. Guide treatment, management, or preventive decisions; and
2. Those decisions lead to improved primary health outcomes.

Genetic Panel Testing

Numerous genetic panels are available which test for various combinations of genes associated with hereditary breast and/or ovarian cancer. Some well-designed genetic panel tests include only the genes which have demonstrated clinical utility and high penetrance. However, other panel tests include a broader range of genes for which penetrance levels are intermediate and low and the clinical utility of testing is not yet established. It is not recommended that patients undergo testing for gene mutations with unknown clinical significance as test results may lead to unnecessary interventions and additional testing which is not supported by current evidence or practice standards. Therefore, the clinical utility of each gene included in the panel must be demonstrated in order to establish the medical necessity of the panel test.

CLINICAL EVIDENCE AND LITERATURE REVIEW

EVIDENCE REVIEW

A review of the ECRI, Hayes, Cochrane, and PubMed databases was conducted regarding the use of genetic testing for hereditary breast and ovarian cancer. Below is a summary of the available evidence identified through March 2023.

Genetic Testing: Personal History of Cancer

In 2016 (archived 2022), Hayes conducted a systematic review evaluating the clinical utility of genetic testing for hereditary breast and ovarian cancer (HBOC) in patients with a personal history and a

suggestive family history of cancer.³ In total, 1159 abstracts were examined, 289 studies were identified for further review, and 24 studies were included for detailed review. Sample size ranged from 26 to 9,982. Genetic tests for *BRCA1*, *BRCA2*, or other genes associated with HBOC syndromes were used as index tests, with comparison to *BRCA* ½ status and HBOC status. Outcomes of interest included diagnostic yield, survival, disease progression, uptake of counseling, testing, surveillance/screening methods, and impact on family members.

Overall, studies indicated that genetic testing of an affected individual is clinically useful, capable of leading to changes in patient management, diagnostic thinking about the hereditary cancer syndrome diagnosis and in the clinical management of at-risk relatives. Genetic testing was also found to inform prognosis and/or predict response to treatment for patients with HBOC. The overall quality of evidence was determined to be “moderate”, comprising 11 “very poor- to poor-quality” studies and 13 “fair quality” studies. Limitations included studies’ retrospective design, small sample sizes, statistical weaknesses, and failure to adjust for potential confounders in analyses.

Hayes ultimately assigned an “A” rating (established benefit) for genetic testing of germline variants in patients with a personal history of breast and/or ovarian cancer and a family history suggestive of HBOC. Investigators noted testing’s ability to reduce subsequent cancer risk through surveillance and risk-reducing procedures, identify variants associated with treatment response, identify at-risk family members, and “balanc[e] this with current limitations such as incomplete scientific understanding of pathogenic variants in specific patient populations.”³

Genetic Testing: No Personal History of Cancer

In 2016 (archived 2022), Hayes conducted a systematic review evaluating the clinical utility of genetic testing for hereditary breast and ovarian cancer (HBOC) in patients with no personal history and a suggestive family history of cancer.⁴ In total, 11 studies were included for detailed review. Sample size ranged from 26 to 2,509 unaffected/asymptomatic individuals with a family history suspected of, or confirmed to be at risk for HBOC. Genetic tests for *BRCA1*, *BRCA2*, other genes associated with hereditary cancer syndromes were employed as index tests; *BRCA1/2* status, HBOC status, study setting, insurance status were used as comparators. Outcomes of interest included the incidence of pathogenic variant(s); recommendation for and uptake of counseling/testing. Screening, prophylactic procedures and psychological impact.

Overall, studies indicated that genetic testing of an unaffected individual can lead to a change in patient management, informing clinical management strategies (e.g., surveillance strategies, prophylactic measures) that would otherwise have been missed by testing only 1 or 2 genes (e.g., *BRCA1/2*). Testing also conferred eligibility for prophylactic procedures, and was informative for family members, who subsequently became eligible for genetic counseling/testing, increased disease surveillance, and/or prophylactic procedures. The overall quality of evidence was determined to be “moderate,” with study quality ranging from “poor” to “fair.” Limitations included studies’ retrospective design, small sample size, survival bias, and insufficient reporting of methods, statistical analysis and results. Despite limitations, Hayes ultimately assigned a “B” rating (some proven benefit) for genetic testing of patients who are unaffected/asymptomatic with a family history of HBOC syndrome.

Genetic Testing: Non-Covered Testing

In 2016 (archived 2020), Hayes conducted a systematic review evaluating the evidence base of the myRisk Hereditary Cancer (Myriad) genetic test.⁵ The panel examines 25 genes using next-generation sequencing (NGS) technology combined with patient clinical factors to assess cancer risk. In total, 3 studies were included for review – 1 analytical validity study and 2 studies that assess the 25-gene panel. No evidence basis was identified supporting the use of the myRisk test due to a lack of evidence. Hayes assigned a “D2” rating (insufficient evidence) for the myRisk Hereditary Cancer test due to a lack of research reporting improvements in patient care, and/or informing patients of familial elevated risk for cancer. Investigators called for additional studies assessing analytical and clinical validity and clinical utility to demonstrate the test’s benefit.

CLINICAL PRACTICE GUIDELINES

The medical necessity criteria within this policy are primarily supported and guided by the National Comprehensive Cancer Network Genetic/Familial High-Risk Assessment: Breast and Ovarian clinical practice guidelines.¹

National Comprehensive Cancer Network (NCCN)

The Version 3.2023 NCCN guidelines regarding hereditary assessment for breast and ovarian cancer recommend the following:¹

NCCN GUIDELINES Breast and Ovarian Management Based on Genetic Test Results		
Gene	Breast Cancer Risk Management	Ovarian Cancer Risk Management
<i>ATM</i>	<u>Increased Risk of Breast Cancer</u> <ul style="list-style-type: none"> Screening: Annual mammogram and consider tomosynthesis and breast MRI with contrast starting at age 40 years. Risk-reducing mastectomy: Consider based on family history. 	Risk-reducing RRSO: Manage based on family history.
<i>BARD1</i>	<ul style="list-style-type: none"> Screening: Annual mammogram with consideration of tomosynthesis and breast MRI with contrast starting at age 40 years. Risk-reducing mastectomy: consider based on family history. 	no evidence for increased risk
<i>BRCA1</i>	<u>Increased Risk of Breast Cancer</u> <ul style="list-style-type: none"> See NCCN guidelines regarding BRCA Pathogenic Variant-Positive Management¹ 	<u>Increased Risk of Ovarian Cancer</u> <ul style="list-style-type: none"> See NCCN guidelines regarding BRCA Pathogenic Variant--Positive Management¹
<i>BRCA2</i>	<u>Increased Risk of Breast Cancer</u> <ul style="list-style-type: none"> See NCCN guidelines regarding BRCA Pathogenic Variant-Positive Management¹ 	<u>Increased Risk of Ovarian Cancer</u> <ul style="list-style-type: none"> See NCCN guidelines regarding BRCA Pathogenic Variant-Positive Management¹

<i>BRIP1</i>	Potential increase in female breast cancer (including triple negative) risk with insufficient evidence for risk management	<u>Increased Risk of Ovarian Cancer</u> <ul style="list-style-type: none"> Consider risk-reducing salpingo-oophorectomy at 45-50 years.
<i>CDH1</i>	<u>Increased Risk of Lobular Breast Cancer</u> <ul style="list-style-type: none"> Screening: Annual mammogram and consider tomosynthesis and breast MRI with contrast starting at age 30 years. Risk-reducing mastectomy: Consider based on family history. 	No increased risk of ovarian cancer
<i>CDKN2A</i>	No increased risk of breast cancer	No increased risk of ovarian cancer
<i>CHEK2</i>	<u>Increased Risk of Breast Cancer</u> <ul style="list-style-type: none"> Screening: Annual mammogram and consider tomosynthesis and breast MRI with contrast age 40 years. Risk-reducing mastectomy: Evidence insufficient, manage based on family history. 	No increased risk of ovarian cancer
<i>MSH2, MLH1, MSH6, PMS2, EPCAM</i>	Unknown or insufficient evidence for breast cancer risk. Manage based on family history.	<u>Increased Risk of Ovarian Cancer</u> <ul style="list-style-type: none"> See NCCN guidelines for Genetic/Familial High-Risk Assessment: Colorectal²
<i>NBN</i>	<ul style="list-style-type: none"> No increased risk of breast cancer 	<u>Potential increase in ovarian cancer risk</u> <ul style="list-style-type: none"> Consider risk-reducing salpingo-oophorectomy based on family history
<i>NF1</i>	<u>Increased Risk of Breast Cancer</u> <ul style="list-style-type: none"> Screening: Annual mammogram starting at age 30 yrs and consider tomosynthesis and breast MRI with contrast from ages 30-50 years. Risk-reducing mastectomy: Evidence insufficient, manage based on family history. 	No increased risk of ovarian cancer
<i>PALB2</i>	<u>Increased Risk of Breast Cancer</u> <ul style="list-style-type: none"> Screening: Annual mammogram and consider tomosynthesis and breast MRI with contrast age 30 years. Risk-reducing mastectomy: Consider based on family history. 	<u>Potential increase in ovarian cancer risk</u> <ul style="list-style-type: none"> Consider risk-reducing salpingo-oophorectomy based on family history, evidence insufficient
<i>PTEN</i>	<u>Increased Risk of Breast Cancer</u> <ul style="list-style-type: none"> See the NCCN guidelines regarding Cowden Syndrome management 	No increased risk of ovarian cancer
<i>RAD51C</i>	Potential increase in triple-negative female breast cancer risk with insufficient evidence for risk management, manage based on family history	<u>Increased Risk of Ovarian Cancer</u> <ul style="list-style-type: none"> Consider risk-reducing salpingo-oophorectomy at 45-50 years.

<i>RAD51D</i>	Potential increase in triple-negative female breast cancer risk with insufficient evidence for risk management, manage based on family history	<u>Increased Risk of Ovarian Cancer</u> <ul style="list-style-type: none"> Consider risk-reducing salpingo-oophorectomy at 45-50 years.
<i>STK11</i>	<u>Increased Risk of Breast Cancer</u> <ul style="list-style-type: none"> Screening: See NCCN guidelines for Genetic/Familial High-Risk Assessment: Colorectal² – Peutz-Jeghers syndrome Risk-reducing mastectomy: Evidence insufficient, manage based on family history. 	<u>Increased Risk of Non-epithelial Ovarian Cancer</u> See NCCN guidelines for Genetic/Familial High-Risk Assessment: Colorectal ² – Peutz-Jeghers syndrome
<i>TP53</i>	<u>Increased Risk of Breast Cancer</u> See NCCN guidelines for Li-Fraumeni Syndrome Management ¹	No increased risk of ovarian cancer

For patients with known familial pathogenic/likely pathogenic variant(s), NCCN recommends testing for the specific familial pathogenic/likely pathogenic variant(s).¹

The American College of Obstetricians and Gynecologists (ACOG)

In 2018 (and reaffirmed in 2023) the ACOG issued a practice bulletin with recommendations for the management of hereditary breast and ovarian cancer syndrome.⁶ The following recommendation were made on the basis of “good and consistent scientific evidence”:

- Genetic counseling is recommended for all women with ovarian epithelial cancer (this includes fallopian tube cancer or primary peritoneal cancer) and for individuals who have a personal or family history of breast cancer or ovarian cancer.
- Women with *BRCA* mutations or who carry another actionable deleterious mutation that is predisposing to breast cancer should be offered risk-reducing bilateral mastectomy.
- Women with *BRCA* mutations or who carry another actionable deleterious mutation predisposing to ovarian cancer should be offered risk-reducing bilateral salpingo-oophorectomy. The timing of risk-reducing bilateral salpingo-oophorectomy can be individualized based on the particular genetic mutation, the patient’s desires for future childbearing, and family history. Typically, risk-reducing salpingo-oophorectomy is recommended at age 35–40 years for *BRCA1* carriers with the highest lifetime risk of ovarian cancer, whereas women with *BRCA2* may consider delaying until age 40–45 years because of later onset of ovarian cancer.
- For a risk-reducing bilateral salpingo-oophorectomy, all tissue from the ovaries and fallopian tubes should be removed. Thorough visualization of the peritoneal surfaces with pelvic washings should be performed. Complete, serial sectioning of the ovaries and fallopian tubes is necessary, with microscopic examination for occult cancer.

In 2017, the ACOG issued a position statement opposing requirements that genetic counseling be provided solely by a “certified” genetic counselor before genetic testing is ordered.⁷ Authors affirmed

that obstetrician-gynecologists are fully trained and qualified to counsel patients regarding genetic issues, and argued that such restrictions would impose unnecessary barriers to care.

EVIDENCE SUMMARY

Evidence is sufficient to support the use of genetic testing for hereditary breast and ovarian cancer among patients with a known breast and/or ovarian cancer mutation, and patients with and without a personal history of cancer. Studies continue to indicate that genetic testing of these populations can lead to a change in patient management, informing clinical management strategies (e.g., surveillance strategies, prophylactic measures) that would otherwise have been missed by testing only 1 or 2 genes (e.g., BRCA1/2). Testing also conferred eligibility for prophylactic procedures and was informative for family members who subsequently became eligible for genetic counseling/testing, increased disease surveillance, and/or prophylactic procedures. In addition, recent evidence-based clinical practice guidelines from the National Comprehensive Cancer Network and the American College of Obstetricians and Gynecology continue to support genetic testing for appropriate patients.

Evidence does not support genetic panel testing that include additional genes not identified as hereditary breast and/or ovarian cancer gene mutation(s) as a method of cancer risk assessment. Additional studies assessing analytical and clinical validity and clinical utility to demonstrate these tests' benefit are needed.

BILLING GUIDELINES AND CODING

BART testing for large rearrangements (e.g., large deletions and/or duplications), billed with 81164 **OR** 81166 and/or 81167, may be denied as not covered when the medical necessity for hereditary breast or ovarian cancer testing (above) is not met.

When no specific CPT or HCPCS code exists for the panel, the provider is required to bill using an unlisted code. It is not appropriate for the provider to bill any of the tests in a panel separately as if they were performed individually. This is a misrepresentation of services performed and is not appropriate based on either CPT or CMS guidelines. In a "Healthcare Fraud Prevention Partnership" white paper published in May, 2018, CMS identified unbundling of lab panels as an example of fraudulent billing.

CODES*		
CPT	0102U	Hereditary breast cancer-related disorders (eg, hereditary breast cancer, hereditary ovarian cancer, hereditary endometrial cancer), genomic sequence analysis panel utilizing a combination of NGS, Sanger, MLPA, and array CGH, with mRNA analytics to resolve variants of unknown significance when indicated (17 genes [sequencing and deletion/duplication])
	0103U	Hereditary ovarian cancer (eg, hereditary ovarian cancer, hereditary endometrial cancer), genomic sequence analysis panel utilizing a combination of NGS, Sanger, MLPA, and array CGH, with mRNA analytics to resolve variants of unknown significance when indicated (24 genes [sequencing and deletion/duplication], EPCAM [deletion/duplication only])
	0129U	Hereditary breast cancer–related disorders (eg, hereditary breast cancer, hereditary ovarian cancer, hereditary endometrial cancer), genomic sequence

		analysis and deletion/duplication analysis panel (ATM, BRCA1, BRCA2, CDH1, CHEK2, PALB2, PTEN, and TP53)
0131U		Hereditary breast cancer-related disorders (eg, hereditary breast cancer, hereditary ovarian cancer, hereditary endometrial cancer), targeted mRNA sequence analysis panel (13 genes) (List separately in addition to code for primary procedure)
0132U		Hereditary ovarian cancer-related disorders (eg, hereditary breast cancer, hereditary ovarian cancer, hereditary endometrial cancer), targeted mRNA sequence analysis panel (17 genes) (List separately in addition to code for primary procedure)
0137U		PALB2 (partner and localizer of BRCA2) (eg, breast and pancreatic cancer) mRNA sequence analysis (List separately in addition to code for primary procedure)
0138U		BRCA1 (BRCA1, DNA repair associated), BRCA2 (BRCA2, DNA repair associated) (eg, hereditary breast and ovarian cancer) mRNA sequence analysis (List separately in addition to code for primary procedure)
0238U		Oncology (Lynch syndrome), genomic DNA sequence analysis of MLH1, MSH2, MSH6, PMS2, and EPCAM, including small sequence changes in exonic and intronic regions, deletions, duplications, mobile element insertions, and variants in non-uniquely mappable regions
81162		BRCA1, BRCA2 (breast cancer 1 and 2) (eg, hereditary breast and ovarian cancer) gene analysis; full sequence analysis and full duplication/deletion analysis
81163		BRCA1 (BRCA1, DNA repair associated), BRCA2 (BRCA2, DNA repair associated) (eg, hereditary breast and ovarian cancer) gene analysis; full sequence analysis
81164		BRCA1 (BRCA1, DNA repair associated), BRCA2 (BRCA2, DNA repair associated) (eg, hereditary breast and ovarian cancer) gene analysis; full duplication/deletion analysis (ie, detection of large gene rearrangements)
81165		BRCA1 (BRCA1, DNA repair associated) (eg, hereditary breast and ovarian cancer) gene analysis; full sequence analysis
81166		BRCA1 (BRCA1, DNA repair associated) (eg, hereditary breast and ovarian cancer) gene analysis; full duplication/deletion analysis (ie, detection of large gene rearrangements)
81167		BRCA2 (BRCA2, DNA repair associated) (eg, hereditary breast and ovarian cancer) gene analysis; full duplication/deletion analysis (ie, detection of large gene rearrangements)
81212		BRCA1, BRCA2 (breast cancer 1 and 2) (eg, hereditary breast and ovarian cancer) gene analysis; 185delAG, 5385insC, 6174delT variants
81215		BRCA1 (breast cancer 1) (eg, hereditary breast and ovarian cancer) gene analysis; known familial variant
81216		BRCA2 (breast cancer 2) (eg, hereditary breast and ovarian cancer) gene analysis; full sequence analysis
81217		BRCA2 (breast cancer 2) (eg, hereditary breast and ovarian cancer) gene analysis; known familial variant
81288		MLH1 (mutL homolog 1, colon cancer, nonpolyposis type 2) (eg, hereditary non-polyposis colorectal cancer, Lynch syndrome) gene analysis; promoter methylation analysis
81292		MLH1 (mutL homolog 1, colon cancer, nonpolyposis type 2) (eg, hereditary non-polyposis colorectal cancer, Lynch syndrome) gene analysis; full sequence analysis

81293	MLH1 (mutL homolog 1, colon cancer, nonpolyposis type 2) (eg, hereditary non-polyposis colorectal cancer, Lynch syndrome) gene analysis; known familial variants
81294	MLH1 (mutL homolog 1, colon cancer, nonpolyposis type 2) (eg, hereditary non-polyposis colorectal cancer, Lynch syndrome) gene analysis; duplication/deletion variants
81295	MSH2 (mutS homolog 2, colon cancer, nonpolyposis type 1) (eg, hereditary non-polyposis colorectal cancer, Lynch syndrome) gene analysis; full sequence analysis
81296	MSH2 (mutS homolog 2, colon cancer, nonpolyposis type 1) (eg, hereditary non-polyposis colorectal cancer, Lynch syndrome) gene analysis; known familial variants
81297	MSH2 (mutS homolog 2, colon cancer, nonpolyposis type 1) (eg, hereditary non-polyposis colorectal cancer, Lynch syndrome) gene analysis; duplication/deletion variants
81298	MSH6 (mutS homolog 6 [E. coli]) (eg, hereditary non-polyposis colorectal cancer, Lynch syndrome) gene analysis; full sequence analysis
81299	MSH6 (mutS homolog 6 [E. coli]) (eg, hereditary non-polyposis colorectal cancer, Lynch syndrome) gene analysis; known familial variants
81300	MSH6 (mutS homolog 6 [E. coli]) (eg, hereditary non-polyposis colorectal cancer, Lynch syndrome) gene analysis; duplication/deletion variants
81307	PALB2 (partner and localizer of BRCA2) (eg, breast and pancreatic cancer) gene analysis; full gene sequence
81308	PALB2 (partner and localizer of BRCA2) (eg, breast and pancreatic cancer) gene analysis; known familial variant
81321	PTEN (phosphatase and tensin homolog) (eg, Cowden syndrome, PTEN hamartoma tumor syndrome) gene analysis; full sequence analysis
81322	PTEN (phosphatase and tensin homolog) (eg, Cowden syndrome, PTEN hamartoma tumor syndrome) gene analysis; known familial variant
81323	PTEN (phosphatase and tensin homolog) (eg, Cowden syndrome, PTEN hamartoma tumor syndrome) gene analysis; duplication/deletion variant
81404	Molecular pathology procedure, Level 5 (eg, analysis of 2-5 exons by DNA sequence analysis, mutation scanning or duplication/deletion variants of 6-10 exons, or characterization of a dynamic mutation disorder/triplet repeat by Southern blot analysis)
81405	Molecular pathology procedure, Level 6 (eg, analysis of 6-10 exons by DNA sequence analysis, mutation scanning or duplication/deletion variants of 11-25 exons, regionally targeted cytogenomic array analysis)
81406	Molecular pathology procedure, Level 7 (eg, analysis of 11-25 exons by DNA sequence analysis, mutation scanning or duplication/deletion variants of 26-50 exons, cytogenomic array analysis for neoplasia)
81432	Hereditary breast cancer-related disorders (eg, hereditary breast cancer, hereditary ovarian cancer, hereditary endometrial cancer); genomic sequence analysis panel, must include sequencing of at least 14 genes, including ATM, BRCA1, BRCA2, BRIP1, CDH1, MLH1, MSH2, MSH6, NBN, PALB2, PTEN, RAD51C, STK11, and TP53

	81433	Hereditary breast cancer-related disorders (eg, hereditary breast cancer, hereditary ovarian cancer, hereditary endometrial cancer); duplication/deletion analysis panel, must include analyses for BRCA1, BRCA2, MLH1, MSH2, and STK11
	81445	Targeted genomic sequence analysis panel, solid organ neoplasm, DNA analysis and RNA analysis when performed, 5-50 genes (EG, ALK, BRAF, CDKN2A, EGFR, ERBB2, KIT, KRAS, MET, NRAS, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, if performed; DNA analysis or combined DNA and RNA analysis
	81455	Targeted genomic sequence analysis panel, solid organ or hemtolymploid neoplasm or disorder, 51 or greater genes (EG, ALK, BRAF, CDKN2A, CEBPA, DNMT3A, EGFR, ERBB2, EZH2, FLT3, IDH1, IDH2, JAK2, KIT, KRAS, MET, MLL, NOTCH1, NPM1, NRAS, MET, NOTCH1, PDGFRA, PDGFRB, PGR, PIK3CA, PTEN, RET), interrogation for sequence variants and copy number variants or rearrangements, or isoform expression or mRNA expression levels, if performed; DNA analysis or combined DNA and RNA analysis
	81479	Unlisted molecular pathology procedure
HCPCS	None	

***Coding Notes:**

- The above code list is provided as a courtesy and may not be all-inclusive. Inclusion or omission of a code from this policy neither implies nor guarantees reimbursement or coverage. Some codes may not require routine review for medical necessity, but they are subject to provider contracts, as well as member benefits, eligibility and potential utilization audit.
- All unlisted codes are reviewed for medical necessity, correct coding, and pricing at the claim level. If an unlisted code is submitted for non-covered services addressed in this policy then it will be **denied as not covered**. If an unlisted code is submitted for potentially covered services addressed in this policy, to avoid post-service denial, **prior authorization is recommended**.
- **See the non-covered and prior authorization lists on the Company [Medical Policy](#), [Reimbursement Policy](#), [Pharmacy Policy](#) and [Provider Information website](#) for additional information.**
- HCPCS/CPT code(s) may be subject to National Correct Coding Initiative (NCCI) procedure-to-procedure (PTP) bundling edits and daily maximum edits known as “medically unlikely edits” (MUEs) published by the Centers for Medicare and Medicaid Services (CMS). This policy does not take precedence over NCCI edits or MUEs. Please refer to the CMS website for coding guidelines and applicable code combinations.

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POLICY REVISION HISTORY

DATE	REVISION SUMMARY
2/2023	Annual review, no changes. Converted to new policy template.
3/2023	Interim update. Denials for non-covered testing changed from “investigational” to “not medically necessary.”
7/2023	Annual Review. Added BARD1 to genes associated with breast cancer. Added +RNAinsight for BreastNext and +RNAinsight for OvaNext by Ambry Genetics to noncovered panels.