

Gene Expression Profiling for Cutaneous Melanoma

(204146)

Medical Benefit		Effective Date: 10/01/18	Next Review Date: 07/19
Preauthorization	No	Review Dates: 07/18	

This protocol considers this test or procedure investigational. If the physician feels this service is medically necessary, preauthorization is recommended.

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:	Interventions of interest are:	Comparators of interest are:	Relevant outcomes include:
With suspicious	 Gene expression profiling 	 Dermatology exam and 	 Overall survival
pigmented lesions	with the DermTech	dermoscopy	 Disease-specific survival
(based on ABCDE	Pigmented Lesion Assay to		Test accuracy
and/or ugly duckling	determine which lesions		Test validity
criteria) being considered for biopsy	should proceed to biopsy		Resource utilization
Individuals:	Interventions of interest are:	Comparators of interest are:	Relevant outcomes include:
Who have melanocytic	 Gene expression profiling 	 Histopathology alone 	 Overall survival
lesions with indeter-	with the myPath	 Comparative genomic 	 Disease-specific survival
minate histopathologic	Melanoma test added to	hybridization added to	Test accuracy
features	histopathology to aid in	histopathology	Test validity
	diagnosis of melanoma	 Fluorescence in situ 	Change in disease status
		hybridization added to histopathology	 Treatment-related morbidity
Individuals:	Interventions of interest are:	Comparators of interest are:	Relevant outcomes include:
 With American Joint 	 Gene expression profiling 	 Sentinel lymph node biopsy 	 Overall survival
Committee on Cancer	with the DecisionDx-	 Prognostic tools 	 Disease-specific survival
stage I or II cutaneous	Melanoma test to		 Test accuracy
melanoma	determine whether to		Test validity
	perform sentinel lymph		 Change in disease status
	node biopsy		 Resource utilization
			 Treatment-related
			morbidity

DESCRIPTION

Laboratory tests have been developed that detect the expression of different genes in pigmented lesions or melanoma tumor tissue. Test results may help providers and patients decide whether to biopsy suspicious pigmented lesions, aid in diagnosis of lesions with indeterminate histopathologic lesions or determine whether to perform sentinel lymph node biopsy in patients diagnosed with stage I or II cutaneous melanoma. This protocol summarizes the evidence of three tests and is organized by indication.

Last Review Date: 07/18

SUMMARY OF EVIDENCE

For individuals with suspicious pigmented lesions (based on ABCDE and/or ugly duckling criteria) being considered for biopsy who receive gene expression profiling with the DermTech Pigmented Lesion Assay to determine which lesions should proceed to biopsy, the evidence includes observational studies. Relevant outcomes are overall survival, disease-specific survival, test accuracy and validity, and resource utilization. The Pigmented Lesion Assay has one clinical validity study with many methodologic and reporting limitations. Therefore, performance characteristics are not well-characterized. Also, the test has not been compared with dermoscopy, another tool frequently used to make biopsy decisions. No direct evidence of clinical utility was identified. Given that the evidence is insufficient to demonstrate test performance, no inferences can be made about clinical utility through a chain of evidence. The evidence is insufficient to determine the effects of the technology on health outcomes.

For individuals who have melanocytic lesions with indeterminate histopathologic features who receive gene expression profiling with the myPath Melanoma test added to histopathology to aid in the diagnosis of melanoma, the evidence includes observational studies. Relevant outcomes are overall survival, disease-specific survival, test accuracy and validity, change in disease status, treatment-related morbidity. The myPath test has one clinical validity study, which includes long-term follow-up to establish the clinical diagnosis as the reference standard. However, it is not clear if the study population included lesions that were indeterminate following histopathology and the study had other methodologic and reporting limitations. Therefore, performance characteristics are not well-characterized. No direct evidence of clinical utility was identified. Given that the evidence is insufficient to demonstrate test performance, no inferences can be made about clinical utility through a chain of evidence. The evidence is insufficient to determine the effects of the technology on health outcomes.

For individuals with American Joint Committee on Cancer (AJCC) stage I or II cutaneous melanoma who receive gene expression profiling with the DecisionDx-Melanoma test to determine whether to perform sentinel lymph node biopsy (SLNB), the evidence includes observational studies. Relevant outcomes are overall survival, disease-specific survival, test accuracy and validity, change in disease status, resource utilization and treatmentrelated morbidity. The DecisionDx-Melanoma test has two independent clinical validity studies that have reported five year recurrence-free survival (RFS) in AJCC stage I or II patients. Gerami et al (2015) reported RFS rates of 98% in DecisionDx class 1 (low risk) without confidence intervals (Cis), in AJCC stage I or II patients. Zager et al (2017) reported RFS rates of 96% (95% CI, 94% to 99%) for DecisionDx class 1 in patients with AJCC stage I disease; they also reported RFS rates of 74% (95% CI, 60% to 91%) for DecisionDx class 1 in patients with AJCC stage II disease. Although CIs were not available for the first study, RFS does not appear to be well-characterized as evidenced by the variation in estimates across studies. Zager et al (2017) also reported that in 56 patients who were DecisionDx class 1 (low risk) but SLNB-positive, 22 recurrences (39%) occurred over five years. If the DecisionDx test were used as a triage for SLNB, these patients would not undergo SLNB and would likely not receive adjuvant therapy, which has shown to be effective at prolonging time to recurrence in node-positive patients. No direct evidence of clinical utility was identified. Given that the evidence is insufficient to demonstrate test performance, no inferences can be made about clinical utility through a chain of evidence. There is also not an explicated, evidence-based management pathway for the use of the test. The evidence is insufficient to determine the effects of the technology on health outcomes.

POLICY

Gene expression testing, including but not limited to the Pigmented Lesion Assay, in the evaluation of patients with suspicious pigmented lesions is considered **investigational**.

Gene expression testing, including but not limited to the myPath Melanoma test, in the evaluation of patients with melanocytic lesions with indeterminate histopathologic features is considered **investigational**.

Last Review Date: 07/18

Gene expression testing, including but not limited to DecisionDx-Melanoma, in the evaluation of patients with cutaneous melanoma is considered **investigational** for all indications.

POLICY GUIDELINES

GENETIC COUNSELING

Experts recommend formal genetic counseling for patients who are at risk for inherited disorders and who wish to undergo genetic testing. Interpreting the results of genetic tests and understanding risk factors can be difficult for some patients; genetic counseling helps individuals understand the impact of genetic testing, including the possible effects the test results could have on the individual or their family members. It should be noted that genetic counseling may alter the utilization of genetic testing substantially and may reduce inappropriate testing; further, genetic counseling should be performed by an individual with experience and expertise in genetic medicine and genetic testing methods.

BACKGROUND

CUTANEOUS MELANOMA

Cutaneous melanoma accounts for more than 90% of cases of melanoma.¹ For many decades, melanoma incidence was rapidly increasing in the United States. However, recent estimates have suggested the rise may be slowing. In 2018, more than 90,000 new cases of melanoma are expected to be diagnosed, and more than 9000 people are expected to die of melanoma.²

Risk Factors

Exposure to solar ultraviolet radiation is a major risk factor for melanoma. Most melanomas occur on the sunexposed skin, particularly those areas most susceptible to sunburn. Likewise, features that are associated with an individual's sensitivity to sunlight, such as light skin pigmentation, red or blond hair, blue or green eyes, freckling tendency, and poor tanning ability are well-known risk factors for melanoma.^{3, 4} There is also a strong association between high total body nevus counts and melanoma.⁵

Several genes appear to contribute to melanoma predisposition such as tumor suppressor gene CDKN2A, melanocortin-1 receptor (MC1R) gene, and BAP1 variants. Individuals with either familial or sporadic melanoma have a two to three times increased risk of developing a subsequent primary melanoma. Several occupational exposures and lifestyle factors, such as body mass index and smoking, have been evaluated as possible risk factors for melanoma.

Diagnosis

Primary care providers evaluate suspicious pigmented lesions to determine who should be referred to dermatology. Factors considered include both a patient's risk for melanoma as well as a visual examination of the lesion. The visual examination assesses whether the lesion has features suggestive of melanoma.

Criteria for features suggestive of melanoma have been developed. One checklist is the ABCDE checklist¹¹:

- Asymmetry;
- Border irregularities;
- Color variegation;
- Diameter ≥ six mm;
- Evolution.

Last Review Date: 07/18

Another criteria commonly used is the "ugly duckling" sign. ¹² An ugly duckling is a nevus that is obviously different from others in a given patient. Primary care providers generally have a low threshold for referral to dermatology.

Melanoma is difficult to diagnose based on visual examination, and the criterion standard for diagnosis is histopathology. There is a low threshold for excisional biopsy of suspicious lesions for histopathologic examination due to the procedure's ease and low risk as well as the high probability of missing melanoma. However, the yield of biopsy is fairly low. The number of biopsies performed to yield one melanoma diagnosis has been estimated to be about 15 for U.S. dermatologists.¹³ Therefore a test that could accurately identify those lesions not needing a biopsy (i.e., a rule-out test for biopsy) could be clinically useful.

Treatment and Surveillance

Many treatments and surveillance decisions are determined by a patient's prognostic stage group based the American Joint Committee on Cancer tumor, node, metastasis staging system. ¹⁴ The prognostic groups are as follows: stage I, T1a through T2a primary melanomas without evidence of regional or distant metastases; stage II, T2b through T4b primary melanomas without evidence of lymphatic disease or distant metastases; stage III, pathologically documented involvement of regional lymph nodes or in transit or satellite metastases (N1 to N3); stage IV, distant metastases.

Patients may also undergo sentinel lymph node biopsy to gain more definitive information about the status of the regional nodes.

Wide local excision is the definitive surgical treatment of melanoma. Following surgery, patients with American Joint Committee on Cancer stage I or II (node-negative) melanoma do not generally receive adjuvant therapy. Patients with higher risk melanoma receive adjuvant immunotherapy or targeted therapy. Ipilimumab has been shown to prolong recurrence-free survival by approximately 25% compared with placebo at a median of 5.3 years in patients with resected, stage III disease. ¹⁵ Nivolumab has been shown to further prolong survival compared with ipilimumab by approximately 35% at 18 months. ¹⁶ For patients who are BRAF V600 variant-positive with stage III melanoma, the combination of dabrafenib plus trametinib has been estimated to prolong relapse-free survival by approximately 50% over three years. ¹⁷

Patients with stage I and II disease should undergo an annual routine physical and dermatologic examination. However, follow-up strategies and intervals have not been standardized or tested, and there is no consensus. These patients typically do not receive surveillance imaging. Patients with stage III melanoma may be managed with more frequent follow-up and imaging surveillance following therapy.

Gene Expression Profiling

Gene expression profiling measures the activity of thousands genes simultaneously and creates a snapshot of cellular function. Data for gene expression profiles are generated by several molecular technologies including DNA microarrays that measures activity relative to previously identified genes and RNA-Seq that directly sequences and quantifies RNA molecules. Clinical applications of gene expression profiling include disease diagnosis, disease classification, prediction of drug response, and prognosis.

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). The Pigmented Lesion Assay, myPath Melanoma, and DecisionDx-Melanoma tests are available under the auspices of the CLIA. Laboratories that offer LDTs must be licensed by the CLIA for high-complexity testing. To date, the U.S. Food and Drug Administration has chosen not to require any regulatory review of this test.

Protocol

Gene Expression Profiling for Cutaneous Melanoma

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.

- Chang AE, Karnell LH, Menck HR. The National Cancer Data Base report on cutaneous and noncutaneous melanoma: a summary of 84,836 cases from the past decade. The American College of Surgeons Commission on Cancer and the American Cancer Society. Cancer. Oct 15 1998;83(8):1664-1678. PMID 9781962
- 2. Siegel RL, Miller KD, Jemal A. Cancer statistics, 2018. Jan 2018;68(1):7-30. PMID 29313949
- 3. Gilchrest BA, Eller MS, Geller AC, et al. The pathogenesis of melanoma induced by ultraviolet radiation. N Engl J Med. Apr 29 1999;340(17):1341-1348. PMID 10219070
- 4. Gandini S, Sera F, Cattaruzza MS, et al. Meta-analysis of risk factors for cutaneous melanoma: III. Family history, actinic damage and phenotypic factors. Eur J Cancer. Sep 2005;41(14):2040-2059. PMID 16125929
- 5. Caini S, Gandini S, Sera F, et al. Meta-analysis of risk factors for cutaneous melanoma according to anatomical site and clinico-pathological variant. Eur J Cancer. Nov 2009;45(17):3054-3063. PMID 19545997
- Goldstein AM, Chan M, Harland M, et al. Features associated with germline CDKN2A mutations: a GenoMEL study of melanoma-prone families from three continents. J Med Genet. Feb 2007;44(2):99-106. PMID 16905682
- 7. Wendt J, Rauscher S, Burgstaller-Muehlbacher S, et al. Human determinants and the role of melanocortin-1 receptor variants in melanoma risk independent of UV radiation exposure. JAMA Dermatol. Jul 1 2016; 152(7):776-782. PMID 27050141
- 8. Wiesner T, Obenauf AC, Murali R, et al. Germline mutations in BAP1 predispose to melanocytic tumors. Nat Genet. Aug 28 2011;43(10):1018-1021. PMID 21874003
- 9. Chen T, Fallah M, Forsti A, et al. Risk of next melanoma in patients with familial and sporadic melanoma by number of previous melanomas. JAMA Dermatol. Jun 2015;151(6):607-615. PMID 25671687
- 10. Jiang AJ, Rambhatla PV, Eide MJ. Socioeconomic and lifestyle factors and melanoma: a systematic review. Br J Dermatol. Apr 2015;172(4):885-915. PMID 25354495
- 11. Abbasi NR, Shaw HM, Rigel DS, et al. Early diagnosis of cutaneous melanoma: revisiting the ABCD criteria. Jama. Dec 8 2004;292(22):2771-2776. PMID 15585738
- 12. Grob JJ, Bonerandi JJ. The 'ugly duckling' sign: identification of the common characteristics of nevi in an individual as a basis for melanoma screening. Arch Dermatol. Jan 1998;134(1):103-104. PMID 9449921
- 13. Wilson RL, Yentzer BA, Isom SP, et al. How good are US dermatologists at discriminating skin cancers? A number-needed-to-treat analysis. J Dermatolog Treat. Feb 2012;23(1):65-69. PMID 21756146
- 14. Gershenwald JES, R.A.; Hess, K.R.; et al. Melanoma of the Skin. Chicago, IL: American Joint Committee on Cancer; 2017.

Last Review Date: 07/18

ge III melanoma with inilimumah

Last Review Date: 07/18

- 15. Eggermont AM, Chiarion-Sileni V, Grob JJ, et al. Prolonged survival in stage III melanoma with ipilimumab adjuvant therap. N Engl J Med. Nov 10 2016;375(19):1845-1855. PMID 27717298
- 16. Weber J, Mandala M, Del Vecchio M, et al. Adjuvant nivolumab versus ipilimumab in resected stage III or IV melanoma. N Engl J Med. Nov 9 2017;377(19):1824-1835. PMID 28891423
- 17. Long GV, Hauschild A, Santinami M, et al. Adjuvant dabrafenib plus trametinib in stage III BRAF-mutated melanomas. N Engl J Med. Nov 9 2017;377(19):1813-1823. PMID 28891408
- 18. National Center for Biotechnology Information. PRAME preferentially expressed antigen in melanoma. 2018; https://www.ncbi.nlm.nih.gov/gene/23532. Accessed March 30, 2018.
- 19. DermTech. Pigmented Lesion Assay: Non-invasive gene expression analysis of pigmented skin lesions. Performance and Development Notes. 2015; http://dermtech.com/wp-content/uploads/2015/10/White-Paper-DermTech-Melanoma-Assay-.pdf. Accessed March 16, 2018.
- 20. Wachsman W, Morhenn V, Palmer T, et al. Noninvasive genomic detection of melanoma. Br J Dermatol. Apr 2011;164(4):797-806. PMID 21294715
- 21. Gerami P, Alsobrook JP, 2nd, Palmer TJ, et al. Development of a novel noninvasive adhesive patch test for the evaluation of pigmented lesions of the skin. J Am Acad Dermatol. Aug 2014;71(2):237-244. PMID 24906614
- 22. Gerami P, Yao Z, Polsky D, et al. Development and validation of a noninvasive 2-gene molecular assay for cutaneous melanoma. J Am Acad Dermatol. Jan 2017;76(1):114-120 e112. PMID 27707590
- 23. Vestergaard ME, Macaskill P, Holt PE, et al. Dermoscopy compared with naked eye examination for the diagnosis of primary melanoma: a meta-analysis of studies performed in a clinical setting. Br J Dermatol. Sep 2008;159(3):669-676. PMID 18616769
- 24. Murzaku EC, Hayan S, Rao BK. Methods and rates of dermoscopy usage: a cross-sectional survey of US dermatologists stratified by years in practice. J Am Acad Dermatol. Aug 2014;71(2):393-395. PMID 25037790
- 25. Engasser HC, Warshaw EM. Dermatoscopy use by US dermatologists: a cross-sectional survey. J Am Acad Dermatol. Sep 2010;63(3):412-419, 419.e411-412. PMID 20619490
- 26. Bossuyt PM, Irwig L, Craig J, et al. Comparative accuracy: assessing new tests against existing diagnostic pathways. Bmj. May 6 2006;332(7549):1089-1092. PMID 16675820
- 27. Ferris LK, Jansen B, Ho J, et al. Utility of a noninvasive 2-gene molecular assay for cutaneous melanoma and effect on the decision to biopsy. JAMA Dermatol. Jul 1 2017;153(7):675-680. PMID 28445578
- 28. Myriad. n.d. Understanding the myPath® Melanoma Results; https://mypathmelanoma.com/about-mypath-melanoma/understanding-the-mypath-melanoma-results/. Accessed March 30, 2018.
- 29. Clarke LE, Warf MB, Flake DD, 2nd, et al. Clinical validation of a gene expression signature that differentiates benign nevi from malignant melanoma. J Cutan Pathol. Apr 2015;42(4):244-252. PMID 25727210
- 30. Gaiser T, Kutzner H, Palmedo G, et al. Classifying ambiguous melanocytic lesions with FISH and correlation with clinical long-term follow up. Mod Pathol. Mar 2010;23(3):413-419. PMID 20081813
- 31. Vergier B, Prochazkova-Carlotti M, de la Fouchardiere A, et al. Fluorescence in situ hybridization, a diagnostic aid in ambiguous melanocytic tumors: European study of 113 cases. Mod Pathol. May 2011;24(5):613-623. PMID 21151100
- 32. Clarke LE, Flake DD, 2nd, Busam K, et al. An independent validation of a gene expression signature to differentiate malignant melanoma from benign melanocytic nevi. Cancer. Feb 15 2017;123(4):617-628. PMID 27768230
- 33. Ko JS, Matharoo-Ball B, Billings SD, et al. Diagnostic distinction of malignant melanoma and benign nevi by a gene expression signature and correlation to clinical outcomes. Cancer Epidemiol Biomarkers Prev. Jul 2017; 26(7):1107-1113. PMID 28377414
- 34. Cockerell C, Tschen J, Billings SD, et al. The influence of a gene-expression signature on the treatment of diagnostically challenging melanocytic lesions. Per Med. Mar 2017;14(2):123-130. PMID 28757886
- 35. Cockerell CJ, Tschen J, Evans B, et al. The influence of a gene expression signature on the diagnosis and recommended treatment of melanocytic tumors by dermatopathologists. Medicine (Baltimore). Oct 2016; 95(40):e4887. PMID 27749545

- 36. National Comprehensive Cancer Network (NCCN). NCCN Clinical Practice Guidelines in Oncology: Melanoma. Version 2.2018. 2018; https://www.nccn.org/professionals/physician_gls/pdf/melanoma.pdf. Accessed March 23, 2018.
- 37. Castle Biosciences. Cutaneous Melanoma: DecisionDx-Melanoma Overview. n.d.; http://castlebiosciences. com/tests/cutaneous-melanoma/. Accessed March 30, 2018.
- 38. Gerami P, Cook RW, Wilkinson J, et al. Development of a prognostic genetic signature to predict the metastatic risk associated with cutaneous melanoma. Clin Cancer Res. Jan 1 2015;21(1):175-183. PMID 25564571
- 39. Zager JS, Gastman BR, Leachman S, et al. Performance of a prognostic 31-gene expression profile in an independent cohort of 523 cutaneous melanoma patients. BMC Cancer. Feb 5 2018;18(1):130. PMID 29402264
- 40. Wong SL, Balch CM, Hurley P, et al. Sentinel lymph node biopsy for melanoma: American Society of Clinical Oncology and Society of Surgical Oncology joint clinical practice guideline. J Clin Oncol. Aug 10 2012;30(23): 2912-2918. PMID 22778321
- 41. Wrightson WR, Wong SL, Edwards MJ, et al. Complications associated with sentinel lymph node biopsy for melanoma. Ann Surg Oncol. Jul 2003;10(6):676-680. PMID 12839853
- 42. Soong SJ, Ding S, Coit DG, et al. AJCC: Individualized melanoma patient outcome prediction tools. n.d.; http://www.melanomaprognosis.net/. Accessed March 21, 2018.
- 43. Callender GG, Gershenwald JE, Egger ME, et al. A novel and accurate computer model of melanoma prognosis for patients staged by sentinel lymph node biopsy: comparison with the American Joint Committee on Cancer model. J Am Coll Surg. Apr 2012;214(4):608-617; discussion 617-609. PMID 22342785
- 44. Dicker TJ, Kavanagh GM, Herd RM, et al. A rational approach to melanoma follow-up in patients with primary cutaneous melanoma. Scottish Melanoma Group. Br J Dermatol. Feb 1999;140(2):249-254. PMID 10233217
- 45. Garbe C, Paul A, Kohler-Spath H, et al. Prospective evaluation of a follow-up schedule in cutaneous melanoma patients: recommendations for an effective follow-up strategy. J Clin Oncol. Feb 1 2003;21(3):520-529. PMID 12560444
- 46. Faries MB, Steen S, Ye X, et al. Late recurrence in melanoma: clinical implications of lost dormancy. J Am Coll Surg. Jul 2013;217(1):27-34; discussion 34-26. PMID 23643694
- 47. Hsueh EC, DeBloom JR, Lee J, et al. Interim analysis of survival in a prospective, multi-center registry cohort of cutaneous melanoma tested with a prognostic 31-gene expression profile test. J Hematol Oncol. Aug 29 2017;10(1):152. PMID 28851416
- 48. Gerami P, Cook RW, Russell MC, et al. Gene expression profiling for molecular staging of cutaneous melanoma in patients undergoing sentinel lymph node biopsy. J Am Acad Dermatol. May 2015;72(5):780-785 e783. PMID 25748297
- 49. Ferris LK, Farberg AS, Middlebrook B, et al. Identification of high-risk cutaneous melanoma tumors is improved when combining the online American Joint Committee on Cancer Individualized Melanoma Patient Outcome Prediction Tool with a 31-gene expression profile-based classification. J Am Acad Dermatol. May 2017;76(5):818-825.e813. PMID 28110997
- 50. Berger AC, Davidson RS, Poitras JK, et al. Clinical impact of a 31-gene expression profile test for cutaneous melanoma in 156 prospectively and consecutively tested patients. Curr Med Res Opin. Sep 2016;32(9):1599-1604. PMID 27210115
- 51. Farberg AS, Glazer AM, White R, et al. Impact of a 31-gene expression profiling test for cutaneous melanoma on dermatologists' clinical management decisions. J Drugs Dermatol. May 1 2017;16(5):428-431. PMID 28628677
- 52. Schuitevoerder D, Heath M, Cook RW, et al. Impact of gene expression profiling on decision-making in clinically node negative melanoma patients after surgical staging. J Drugs Dermatol. Feb 1 2018;17(2):196-199. PMID 29462228
- 53. Dillon LD, Gadzia JE, Davidson RS, et al. Prospective, multicenter clinical impact evaluation of a 31-gene expression profile test for management of melanoma patients. Skin. 2018;2(2):111-121. PMID

Protocol

Gene Expression Profiling for Cutaneous Melanoma

Last Review Date: 07/18

- 54. PROPOSED/DRAFT Local Coverage Determination (LCD): MoIDX: Pigmented Lesion Assay (PLA) (DL37715). 2018; https://www.cms.gov/medicare-coverage-database/details/lcd-details.aspx?LCDId=37714&ver=3 &DocType=All&bc=AAlAAAAAAAA. Accessed Apr 25, 2018.
- 55. PROPOSED/DRAFT Local Coverage Determination (LCD): MoIDX: Pigmented Lesion Assay (PLA) and PLA Score (DL37669). 2018; https://www.cms.gov/medicare-coverage-database/details/lcd-details.aspx?LCDId= 37668&ver=2&DocType=All&bc=AAIAAAAAAAAA. Accessed Apr 25, 2018.
- 56. PROPOSED/DRAFT Local Coverage Determination (LCD): MoIDX: Pigmented Lesion Assay (PLA) and PLA Score (DL37533). 2018; https://www.cms.gov/medicare-coverage-database/details/lcd-details.aspx?LCDId= 37532&ver=3&DocType=All&bc=AAIAAAAAAAAA. Accessed Apr 25, 2018.
- 57. PROPOSED/DRAFT Local Coverage Determination (LCD): MoIDX: DecisionDx-Melanoma (DL37725). 2018; https://www.cms.gov/medicare-coverage-database/details/lcd-details.aspx?LCDId=37724&ver=7&DocID=DL37725&bc=gAAAABAAAAA&. Accessed Apr 25, 2018.