



## FEP Medical Policy Manual

### FEP 2.04.97 Microarray-Based Gene Expression Profile Testing for Multiple Myeloma Risk Stratification

**Annual Effective Policy Date: January 1, 2024**

**Original Policy Date: December 2015**

**Related Policies:**

None

## Microarray-Based Gene Expression Profile Testing for Multiple Myeloma Risk Stratification

### Description

#### Description

Multiple myeloma is a genetically complex and invariably fatal disease. A host of well-characterized factors related to tumor biology, tumor burden, and patient-centered characteristics are used to stratify patients into high-, intermediate-, and standard-risk categories for prognostic purposes, as well as determining treatment intensity. However, clinical outcomes have varied among patients in the same risk category who received similar therapy. Thus, more specific methods have been sought to classify multiple myeloma; one such method being proposed is the utilization of a microarray-based gene expression profile (GEP) analysis, which serves to reveal the underlying activity of cellular biologic pathways. This method lends itself to a variety of benefits including the ability to risk-stratify patients with multiple myeloma, as well as guide treatment decisions.

#### OBJECTIVE

The objective of this evidence review is to determine whether risk stratification using a gene expression profile risk score improves the net health outcome in individuals with multiple myeloma.

## POLICY STATEMENT

Microarray-based gene expression profile testing for multiple myeloma is considered **investigational** for all indications.

## POLICY GUIDELINES

According to Mayo Clinic recommendations, a large number of prognostic factors have been validated and categorized into 3 main groups: tumor biology, tumor burden, and patient-related factors. These factors must be considered to individualize the choice of therapy in individuals with multiple myeloma (Table PG1).

**Table PG1. Prognostic Factors in Multiple Myeloma**

| Tumor Biology  | Tumor Burden   | Patient-Related  |
|--|--|--|
| <ul style="list-style-type: none"> <li>• Ploidy</li> <li>• 17p (p53 deletion)</li> <li>• t(14;16)</li> <li>• t(14;20)</li> <li>• t(4;14)</li> <li>• Deletion 13 on conventional cytogenetics</li> <li>• Alterations in chromosome 1</li> <li>• t(11;14)</li> <li>• t(6;14)</li> <li>• Lactate dehydrogenase levels</li> <li>• Plasma cell proliferative rate</li> <li>• Presentation as plasma cell leukemia</li> <li>• High-risk GEP signature<sup>a</sup></li> </ul> | <ul style="list-style-type: none"> <li>• Durie-Salmon stage</li> <li>• International Staging System stage</li> <li>• Extramedullary disease</li> </ul> | <ul style="list-style-type: none"> <li>• ECOG Performance Status</li> <li>• Age</li> <li>• Renal function</li> </ul> |

Adapted from Mikhael et al (2013).

ECOG: Eastern Cooperative Oncology Group; GEP: gene expression profile.

<sup>a</sup> The Mayo Clinic does not currently recommend or routinely perform GEP analysis in a non-research setting. However, Mikhael et al (2013) have suggested GEP analysis will likely play a greater role in the management of multiple myeloma as evidence develops.

## BENEFIT APPLICATION

Experimental or investigational procedures, treatments, drugs, or devices are not covered (See General Exclusion Section of brochure).

Screening (other than the preventive services listed in the brochure) is not covered. Please see Section 6 General exclusions.

Benefits are available for specialized diagnostic genetic testing when it is medically necessary to diagnose and/or manage a patient's existing medical condition. Benefits are not provided for genetic panels when some or all of the tests included in the panel are not covered, are experimental or investigational, or are not medically necessary.

## FDA REGULATORY STATUS

Clinical laboratories may develop and validate tests in-house and market them as a laboratory service; laboratory-developed tests must meet the general regulatory standards of the Clinical Laboratory Improvement Amendments (CLIA). The MyPRS™/MyPRS Plus™ GEP70 test was acquired by Quest Diagnostics in December 2016. Laboratories that offer laboratory-developed tests 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.

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## RATIONALE

### Summary of Evidence

For individuals who have multiple myeloma who received risk stratification using a gene expression profile (GEP) test, the evidence includes retrospective series that correlate risk scores with survival. Relevant outcomes are progression-free survival, overall survival, disease-specific survival, test validity, and other test performance measures. The microarray-based GEP70 test (MyPRS/MyPRS *Plus*) has been reported to risk-stratify multiple myeloma patients. Some predictive models in the body of evidence combine risk status as determined by the GEP70 test with additional clinical or genetic variables. Patients with a high GEP70 risk score have a substantially increased risk of mortality compared with patients without a high score. However, there is no evidence (from available studies) that this test would add incremental value to existing risk stratification methods; nor have any studies demonstrated the need to prospectively allocate patients to risk-based therapies based on the GEP70 score. The evidence is insufficient to determine that the technology results in an improvement in the net health outcome.

## SUPPLEMENTAL INFORMATION

### Practice Guidelines and Position Statements

Guidelines or position statements will be considered for inclusion in 'Supplemental Information' if they were issued by, or jointly by, a US professional society, an international society with US representation, or National Institute for Health and Care Excellence (NICE). Priority will be given to guidelines that are informed by a systematic review, include strength of evidence ratings, and include a description of management of conflict of interest.

#### Mayo Clinic Stratification of Multiple Myeloma and Risk-Adapted Therapy

Guidelines from the Mayo Clinic (2017) have stated that "if indicated, gene expression profiling may be performed to further understand the behavior of the disease and guide therapy."<sup>30</sup>

#### National Comprehensive Cancer Network

The National Comprehensive Cancer Network practice guidelines (v4.2023 ) on multiple myeloma do not provide recommendations regarding use of gene expression profiling.<sup>31</sup>

#### U.S. Preventive Services Task Force Recommendations

Not applicable.

#### Medicare National Coverage

There is no national coverage determination. In the absence of a national coverage determination, coverage decisions are left to the discretion of local Medicare carriers.

## REFERENCES

1. Fonseca R, Barlogie B, Bataille R, et al. Genetics and cytogenetics of multiple myeloma: a workshop report. *Cancer Res.* Feb 15 2004; 64(4): 1546-58. PMID 14989251
2. Palumbo A, Anderson K. Multiple myeloma. *N Engl J Med.* Mar 17 2011; 364(11): 1046-60. PMID 21410373
3. American Cancer Society. Key Statistics About Multiple Myeloma. 2023; <https://www.cancer.org/cancer/types/multiple-myeloma/about/key-statistics.html>. Accessed September 1, 2023.
4. Brenner H, Gondos A, Pulte D. Recent major improvement in long-term survival of younger patients with multiple myeloma. *Blood.* Mar 01 2008; 111(5): 2521-6. PMID 17901246
5. Marinac CR, Ghobrial IM, Birmann BM, et al. Dissecting racial disparities in multiple myeloma. *Blood Cancer J.* Feb 17 2020; 10(2): 19. PMID 32066732
6. Munshi NC, Anderson KC, Bergsagel PL, et al. Consensus recommendations for risk stratification in multiple myeloma: report of the International Myeloma Workshop Consensus Panel 2. *Blood.* May 05 2011; 117(18): 4696-700. PMID 21292777
7. Kyle RA, Rajkumar SV. Criteria for diagnosis, staging, risk stratification and response assessment of multiple myeloma. *Leukemia.* Jan 2009; 23(1): 3-9. PMID 18971951
8. Blad J, Dimopoulos M, Rosiol L, et al. Smoldering (asymptomatic) multiple myeloma: current diagnostic criteria, new predictors of outcome, and follow-up recommendations. *J Clin Oncol.* Feb 01 2010; 28(4): 690-7. PMID 20026810
9. Morgan GJ, Walker BA, Davies FE. The genetic architecture of multiple myeloma. *Nat Rev Cancer.* Apr 12 2012; 12(5): 335-48. PMID 22495321
10. Kyle RA, Durie BG, Rajkumar SV, et al. Monoclonal gammopathy of undetermined significance (MGUS) and smoldering (asymptomatic) multiple myeloma: IMWG consensus perspectives risk factors for progression and guidelines for monitoring and management. *Leukemia.* Jun 2010; 24(6): 1121-7. PMID 20410922
11. Fonseca R, Bergsagel PL, Drach J, et al. International Myeloma Working Group molecular classification of multiple myeloma: spotlight review. *Leukemia.* Dec 2009; 23(12): 2210-21. PMID 19798094
12. Munshi NC, Avet-Loiseau H. Genomics in multiple myeloma. *Clin Cancer Res.* Mar 15 2011; 17(6): 1234-42. PMID 21411439
13. Mikhael JR, Dingli D, Roy V, et al. Management of newly diagnosed symptomatic multiple myeloma: updated Mayo Stratification of Myeloma and Risk-Adapted Therapy (mSMART) consensus guidelines 2013. *Mayo Clin Proc.* Apr 2013; 88(4): 360-76. PMID 23541011
14. Johnson SK, Heuck CJ, Albino AP, et al. The use of molecular-based risk stratification and pharmacogenomics for outcome prediction and personalized therapeutic management of multiple myeloma. *Int J Hematol.* Oct 2011; 94(4): 321-333. PMID 22002477
15. Zhou Y, Barlogie B, Shaughnessy JD. The molecular characterization and clinical management of multiple myeloma in the post-genome era. *Leukemia.* Nov 2009; 23(11): 1941-56. PMID 19657360
16. Shaughnessy J. Primer on medical genomics. Part IX: scientific and clinical applications of DNA microarrays--multiple myeloma as a disease model. *Mayo Clin Proc.* Sep 2003; 78(9): 1098-109. PMID 12962165
17. Hubank M. Gene expression profiling and its application in studies of haematological malignancy. *Br J Haematol.* Mar 2004; 124(5): 577-94. PMID 14871244
18. Zhou Y, Zhang Q, Stephens O, et al. Prediction of cytogenetic abnormalities with gene expression profiles. *Blood.* May 24 2012; 119(21): e148-50. PMID 22496154
19. Matsui S, Simon R, Qu P, et al. Developing and validating continuous genomic signatures in randomized clinical trials for predictive medicine. *Clin Cancer Res.* Nov 01 2012; 18(21): 6065-73. PMID 22927484
20. Meissner T, Seckinger A, Rme T, et al. Gene expression profiling in multiple myeloma--reporting of entities, risk, and targets in clinical routine. *Clin Cancer Res.* Dec 01 2011; 17(23): 7240-7. PMID 21986844
21. Cowan AJ, Green DJ, Kwok M, et al. Diagnosis and Management of Multiple Myeloma: A Review. *JAMA.* Feb 01 2022; 327(5): 464-477. PMID 35103762
22. Shaughnessy JD, Zhan F, Burington BE, et al. A validated gene expression model of high-risk multiple myeloma is defined by deregulated expression of genes mapping to chromosome 1. *Blood.* Mar 15 2007; 109(6): 2276-84. PMID 17105813
23. Kumar SK, Uno H, Jacobus SJ, et al. Impact of gene expression profiling-based risk stratification in patients with myeloma receiving initial therapy with lenalidomide and dexamethasone. *Blood.* Oct 20 2011; 118(16): 4359-62. PMID 21860025
24. Mohan M, Weinhold N, Schinke C, et al. Daratumumab in high-risk relapsed/refractory multiple myeloma patients: adverse effect of chromosome 1q21 gain/amplification and GEP70 status on outcome. *Br J Haematol.* Apr 2020; 189(1): 67-71. PMID 31820442
25. Papanikolaou X, Alapat D, Rosenthal A, et al. The flow cytometry-defined light chain cytoplasmic immunoglobulin index and an associated 12-gene expression signature are independent prognostic factors in multiple myeloma. *Leukemia.* Aug 2015; 29(8): 1713-20. PMID 25753926
26. Agnelli L, Tassone P, Neri A. Molecular profiling of multiple myeloma: from gene expression analysis to next-generation sequencing. *Expert Opin Biol Ther.* Jun 2013; 13 Suppl 1: S55-68. PMID 23614397
27. Amin SB, Yip WK, Minvielle S, et al. Gene expression profile alone is inadequate in predicting complete response in multiple myeloma. *Leukemia.* Nov 2014; 28(11): 2229-34. PMID 24732597
28. Chng WJ, Dispenzieri A, Chim CS, et al. IMWG consensus on risk stratification in multiple myeloma. *Leukemia.* Feb 2014; 28(2): 269-77. PMID 23974982
29. Fonseca R, Monge J, Dimopoulos MA. Staging and prognostication of multiple myeloma. *Expert Rev Hematol.* Feb 2014; 7(1): 21-31. PMID 24483346
30. Dingli D, Ailawadhi S, Bergsagel PL, et al. Therapy for Relapsed Multiple Myeloma: Guidelines From the Mayo Stratification for Myeloma and Risk-Adapted Therapy. *Mayo Clin Proc.* Apr 2017; 92(4): 578-598. PMID 28291589

31. National Comprehensive Cancer Network (NCCN). NCCN Clinical Practice Guidelines in Oncology: Multiple Myeloma. Version 4.2023. [http://www.nccn.org/professionals/physician\\_gls/pdf/myeloma.pdf](http://www.nccn.org/professionals/physician_gls/pdf/myeloma.pdf). Accessed September 1, 2023.

## **POLICY HISTORY - THIS POLICY WAS APPROVED BY THE FEP® PHARMACY AND MEDICAL POLICY COMMITTEE ACCORDING TO THE HISTORY BELOW:**

| <b>Date</b>   | <b>Action</b>  | <b>Description</b>  |
|---------------|----------------|---|
| December 2015 | New policy     | Microarray-based gene expression profile testing for multiple myeloma is considered investigational for all indications.                                      |
| December 2018 | Replace policy | Policy updated with literature review through August 6, 2018; references 30-31 added. Policy statement unchanged.   |
| December 2019 | Replace policy | Policy updated with literature review through August 6, 2019; no references added, reference on NCCN updated. Policy statement unchanged.                     |
| December 2020 | Replace policy | Policy updated with literature review through August 20, 2020; references added. Policy statement unchanged.  |
| December 2021 | Replace policy | Policy updated with literature review through August 22, 2021; no references added. Policy statement unchanged.   |
| December 2022 | Replace policy | Policy updated with literature review through August 22, 2022; reference added. Minor editorial refinements to Policy Guideline statements; intent unchanged. |
| December 2023 | Replace policy | Policy updated with literature review through September 1, 2023; reference added. Policy statement unchanged.   |

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