

mSMART Plasma Cell Proliferative Disorder, Pre-Analysis Cell Sorting, Bone Marrow

#### Overview

#### **Useful For**

Risk stratification of patients with multiple myeloma, which can assist in determining treatment and management decisions

Sorting plasma cells for fluorescence in situ hybridization analysis

Risk stratification of patients with newly diagnosed multiple myeloma

#### **Method Name**

Only orderable as a reflex. For more information see MSMRT / Mayo Algorithmic Approach for Stratification of Myeloma and Risk-Adapted Therapy Report, Bone Marrow or MSMRD / Myeloma Stratification and Risk-Adapted Therapy with Reflex to Minimal Residual Disease, Bone Marrow

Flow Cytometric Cell Selection

#### **NY State Available**

Yes

## Specimen

#### Specimen Type

**Bone Marrow** 

#### Specimen Required

Only orderable as a reflex. For more information see MSMRT / Mayo Algorithmic Approach for Stratification of Myeloma and Risk-Adapted Therapy Report, Bone Marrow or MSMRD / Myeloma Stratification and Risk-Adapted Therapy with Reflex to Minimal Residual Disease, Bone Marrow

**Specimen Type:** Redirected bone marrow **Preferred:** Yellow top (ACD solution A or B)

Acceptable: Lavender top (EDTA) or green top (heparin)

Specimen Volume: 4 mL

#### **Specimen Minimum Volume**

1 mL

## Reject Due To



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Gross	Reject
hemolysis	
Other	Fully clotted

## **Specimen Stability Information**

Specimen Type	Temperature	Time	Special Container
Bone Marrow	Ambient (preferred)	4 days	
	Refrigerated	4 days	

## **Clinical & Interpretive**

#### **Clinical Information**

Multiple myeloma is increasingly recognized as a disease characterized by marked cytogenetic, molecular, and proliferative heterogeneity. This heterogeneity is manifested clinically by varying degrees of disease aggressiveness. Multiple myeloma patients with more aggressive disease experience suboptimal responses to some therapeutic approaches; therefore, identifying these patients is critically important for selecting appropriate treatment options.

MSMRT / Mayo Algorithmic Approach for Stratification of Myeloma and Risk-Adapted Therapy Report, Bone Marrow classifies patients into either standard or high-risk categories based on the results of 2 assays: plasma cell proliferation and FISH for specific multiple myeloma-associated abnormalities.

#### **Reference Values**

Only orderable as a reflex. For more information see MSMRT / Mayo Algorithmic Approach for Stratification of Myeloma and Risk-Adapted Therapy Report, Bone Marrow or MSMRD / Myeloma Stratification and Risk-Adapted Therapy with Reflex to Minimal Residual Disease, Bone Marrow

An interpretive report will be provided.

#### Interpretation

Correlation with clinical, histopathologic and additional laboratory findings is required for final interpretation of these results. The final interpretation of results for clinical management of the patient is the responsibility of the managing physician.

#### **Cautions**

No significant cautionary statements

#### **Clinical Reference**

- Rajkumar SV, Greipp PR. Prognostic factors in multiple myeloma. Hematol Oncol Clin North Am. 1999;13(6):1295-1314
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- 5. Gonsalves WI, Buadi FK, Ailawadhi S, et al. Utilization of hematopoietic stem cell transplantation for the treatment of multiple myeloma: a Mayo Stratification of Myeloma and Risk-Adapted Therapy (mSMART) consensus statement. Bone Marrow Transplant. 2019;54(3):353–367. doi:10.1038/s41409-018-0264-8
- 6. Kapoor P, Ansell SM, Fonseca R, et al. Diagnosis and management of waldenstrom macroglobulinemia: Mayo stratification of macroglobulinemia and risk-adapted therapy (mSMART) guidelines 2016. JAMA Oncol. 2017;3(9):1257–1265. doi:10.1001/jamaoncol.2016.5763
- 7. 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. 2013;88(4):360–376. doi:10.1016/j.mayocp.2013.01.019
- 8. Swerdlow S, Campo E, Harris NL, et al. WHO Classification of Tumours of Haematopoietic and Lymphoid Tissues. IARC Press: Lyon. 2017
- 9. Kumar SK, Rajkumar SV. The multiple myelomas-current concepts in cytogenetic classification and therapy. Nat Rev Clin Oncol. 2018;15(7):409-421. doi:10.1038/s41571-018-0018-y
- 10. Rajkumar SV, Landgren O, Mateos MV. Smoldering multiple myeloma. Blood. 2015;125(20):3069-3075. doi:10.1182/blood-2014-09-568899

## **Performance**

#### **Method Description**

Selection of plasma cells using fluorescence activated cell sorting is the most direct and robust method of obtaining relatively pure plasma cell populations for fluorescence in situ hybridization (FISH) assessment. (Instruction manual: BD FACSMelody Cell Sorter User's Guide. Revision 3. BD Biosciences; 03/2020)

## **PDF Report**

No

#### Day(s) Performed

Specimens processed Monday through Sunday Results reported Monday through Friday

#### Report Available

1 to 11 days

## **Specimen Retention Time**

14 days

#### Performing Laboratory Location

Mayo Clinic Laboratories - Rochester Main Campus



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## **Fees & Codes**

#### **Fees**

- Authorized users can sign in to <u>Test Prices</u> for detailed fee information.
- Clients without access to Test Prices can contact <u>Customer Service</u> 24 hours a day, seven days a week.
- Prospective clients should contact their account representative. For assistance, contact <u>Customer Service</u>.

#### **Test Classification**

This test was developed and its performance characteristics determined by Mayo Clinic in a manner consistent with CLIA requirements. It has not been cleared or approved by the US Food and Drug Administration.

#### **CPT Code Information**

88184-Flow Cytometry; first cell surface, cytoplasmic or nuclear marker 88185 x 5-Flow Cytometry, additional cell surface, cytoplasmic or nuclear marker (each)

#### **LOINC®** Information

Test ID	Test Order Name	Order LOINC® Value
CSMRT	MPCDS Pre-Analysis Cell Sorting, BM	No LOINC Needed

Result ID	Test Result Name	Result LOINC® Value
607682	MPCDS Pre-Analysis Cell Sort	No LOINC Needed
607688	Final Diagnosis	No LOINC Needed