

Test Specific Guidelines

Somatic Mutation Testing-Hematological Malignancies

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Introduction

Somatic mutation testing for hematological malignancies is addressed by this guideline.

Procedures Addressed

The inclusion of any procedure code in this table is provided for informational purposes and is not a guarantee of coverage nor an indication that prior authorization is required.

<u>Procedures addressed by this guideline</u>	<u>Procedure codes</u>
<u>ASXL1 Full Gene Sequencing</u>	<u>81175</u>
<u>ASXL1 Mutation Analysis</u>	<u>81176</u>
<u>ABL1 Mutation Analysis</u>	<u>81170</u>
<u>ABL1 Targeted Mutation Analysis</u>	<u>81401</u>
<u>BCR-ABL1 detection, major breakpoint</u>	<u>81206</u>
<u>BCR-ABL1 detection, minor breakpoint</u>	<u>81207</u>
<u>BCR-ABL1 detection, other breakpoint</u>	<u>81208</u>
<u>BCR-ABL1 major and minor breakpoint fusion transcripts</u>	<u>0016U</u>
<u>CALR Exon 9 Mutation Analysis</u>	<u>81219</u>
<u>CCND1/IGH (t(11;14)) Translocation Analysis, Major Breakpoint</u>	<u>81168</u>
<u>CEBPA Full Gene Sequencing</u>	<u>81218</u>
<u>EZH2 Common Variant(s) (e.g. codon 646)</u>	<u>81237</u>
<u>EZH2 Full Gene Sequencing</u>	<u>81236</u>
<u>FISH Analysis for t(9;22) BCR-ABL1</u>	<u>88271</u>
<u>FLT3 internal tandem duplication MRD-Invivoscribe</u>	<u>0046U</u>
<u>FLT3 Mutation Analysis (internal tandem duplication variants)</u>	<u>81245</u>
<u>FLT3 Mutation Analysis (tyrosine kinase domain variants)</u>	<u>81246</u>

<u>Procedures addressed by this guideline</u>	<u>Procedure codes</u>
<u>IDH1 Mutation Analysis</u>	<u>81120</u>
<u>IDH2 Mutation Analysis</u>	<u>81121</u>
<u>IGH@/BCL2 (t(14;18)) Translocation Analysis, Major Breakpoint Region (MBR) and Minor Cluster Region (mcr) Breakpoints</u>	<u>81278</u>
<u>JAK2 Exons 12 to 15 Sequencing</u>	<u>0027U</u>
<u>JAK2 Mutation</u>	<u>0017U</u>
<u>JAK2 Targeted Mutation Analysis (e.g exons 12 and 13)</u>	<u>81279</u>
<u>JAK2 V617F Mutation Analysis</u>	<u>81270</u>
<u>KIT Targeted Mutation Analysis</u>	<u>81272</u>
<u>KIT Mutation Analysis (D816 variants)</u>	<u>81273</u>
<u>MPL Common Variants (e.g. W515A, W515K, W515L, W515R)</u>	<u>81338</u>
<u>MPL Mutation Analysis, Exon 10</u>	<u>81339</u>
<u>MRDx® BCR-ABL Test</u>	<u>0040U</u>
<u>MyAML NGS- Invivoscribe</u>	<u>0050U</u>
<u>NPM1 MRD- Invivoscribe</u>	<u>0049U</u>
<u>NPM1 Mutation Analysis</u>	<u>81310</u>
<u>NRAS Mutation Analysis</u>	<u>81311</u>
<u>RUNX1 Mutation Analysis</u>	<u>81334</u>
<u>TERT Targeted Sequence Analysis</u>	<u>81345</u>
<u>Hematolymphoid Neoplasm Molecular Profiling</u>	<u>81450</u>
<u>SF3B1 Common Variants (e.g. A672T, E622D, L833F, R625C, R625L)</u>	<u>81347</u>
<u>Solid Organ or Hematolymphoid Neoplasm Molecular Profiling - Expanded</u>	<u>81455</u>
<u>SRSF2 Common Variants (e.g. P95H, P95L)</u>	<u>81348</u>
<u>TP53 Sequencing</u>	<u>81351</u>
<u>TP53 Targeted Sequence Analysis</u>	<u>81352</u>
<u>U2AF1 Common Variants (e.g. S34F, S34Y, Q157R, Q157P)</u>	<u>81357</u>
<u>ZRSR2 Common Variants (e.g. E65fs, E122fs, R448fs)</u>	<u>81360</u>
<u>MyMRD NGS Panel</u>	<u>0171U</u>

<u>Procedures addressed by this guideline</u>	<u>Procedure codes</u>
<u>Molecular Tumor Marker Test</u>	<u>81400</u>
	<u>81401</u>
	<u>81402</u>
	<u>81403</u>
	<u>81404</u>
	<u>81405</u>
	<u>81406</u>
	<u>81407</u>
	<u>81408</u>
	<u>81479</u>
<u>Molecular Tumor Marker Test</u>	<u>88271</u>

What Are Somatic Mutation Tests?

Definition

A somatic mutation test for hematologic malignancies is broadly defined here as any test that measures changes in DNA, RNA, or chromosomes and is used to make cancer management decisions.

Somatic mutation tests are increasingly useful for therapy selection. Many cancer therapies are targeted at particular gene functions (therapeutic targets) and some require information about the genetics of the malignancy to use the therapies effectively (companion diagnostics). In these cases, National Comprehensive Cancer Network (NCCN) as well as the U.S. Food and Drug Administration (FDA) have outlined tumor testing that is recommended for specific cancers and the associated treatment implications.¹⁻⁴

Test Information

Somatic Mutation Testing

The specific methodology used to identify somatic mutations is dependent upon the type of mutation being investigated.

DNA mutations are generally detected through direct analysis of individual mutations, portions of a gene, a whole gene, panels of genes, or the entire exome.

Chromosome abnormalities, such as translocations or deletions, may be detected through direct visualization of the chromosomes (karyotyping), in situ hybridization of probes (e.g., FISH) to detect deletions or duplications that are too small to see directly, or by DNA-based methods (hybridization arrays or sequencing) that identify deletions or translocation breakpoints.

Gene expression profiling simultaneously measures the amount of RNA being made by many genes. Expression patterns may be used to predict the type of cancer present, the aggressiveness of the malignancy, and therapies that are likely to be effective.

The efficiency of next generation sequencing (NGS) has led to an increasing number of large, multi-gene somatic mutation panels. Given that malignancies can have multiple and unexpected genetic changes, these panels may provide physicians with information about therapeutic targets that would not otherwise be considered.

Guidelines and Evidence

National Comprehensive Cancer Network

The National Comprehensive Cancer Network (NCCN) provided the following guidance:

NCCN Guidelines for Treatment of Cancer by Site provided detailed guidelines on the use of individual markers for each cancer type addressed.⁴

NCCN also maintains a biomarker compendium and stated “the goal of the NCCN Biomarkers Compendium is to provide essential details for those tests which have been approved by NCCN Guideline Panels and are recommended by the NCCN Guidelines.”³ Biomarkers for specific cancer types that are listed in the NCCN Biomarker Compendium have a level of evidence associated with their clinical utility.

NCCN stated that for individuals with acute lymphoblastic leukemia (ALL), molecular characterization by comprehensive testing by next-generation sequencing (NGS) for gene fusions and pathogenic mutations is recommended for determining risk and planning treatment. Specific biomarkers can be found in the table contained within the policy.⁵

NCCN stated that for individuals with cytopenia when myelodysplasia is suspected, bone marrow or peripheral blood cells should be assayed for MDS-associated gene mutations using gene panels that include ASXL1, BCOR, CALR, CBL, DDX41, DNMT3A, ETV6, EZH2, FLT3, GATA2, IDH1, IDH2, JAK2, MPL, NF1, NPM1, NRAS, PHF6, PPM1D, RUNX1, SETBP1, SF3B1, SRSF2, STAG2, STAT3, TET2, TP53, U2AF1, WT1, and ZRSR2.⁶

NCCN stated that for individuals with acute myeloid leukemia (AML): “Several gene mutations are associated with specific prognoses in a subset of patients

(category 2A) and may guide treatment decisions (category 2B). Presently, c-KIT, FLT3-ITD, FLT3-TKD, NPM1, CEBPA (biallelic), IDH1/IDH2, RUNX1, ASXL1, TP53, BCR-ABL and PML-RAR alpha are included in this group. All patients should be tested for mutations in these genes, and multiplex gene panels and comprehensive next-generation sequencing (NGS) analysis are recommended for the ongoing management of AML and various phases of treatment.”⁷

U.S. Food and Drug Administration

Some FDA labels require results from biomarker tests to effectively or safely use the therapy for a specific cancer type.² A list of all Pharmacogenomic Biomarkers included in FDA labeling and associated implications can be found here. While these tests generally consist of a single biomarker, some larger panels of biomarkers are also included in the FDA labeling.

Criteria

This guideline applies to all molecular somatic mutation testing intended for use in hematological malignancies.

Medical necessity criteria differ based on the type of testing being performed (i.e., tests for individual genes separately chosen based on the cancer type versus pre-defined panels of genes) and how that testing will be billed (one or more individual gene-specific procedure codes, specific panel procedure codes or unlisted procedure codes).

Individual Tumor Markers

When separate procedure codes will be billed for individual biomarkers (e.g., Tier 1 MoPath codes 81200-81355 or Tier 2 MoPath codes 81400-81408), each individually billed test will be evaluated separately. The following criteria will be applied:

The member has a malignancy type that will benefit from information provided by the requested test based on at least one of the following:

All criteria are met from a test-specific guideline, if available, or

An oncology therapy FDA label requires results from the test to effectively or safely use the therapy for the member’s cancer type, or

NCCN guidelines include the test in the management algorithm for that particular cancer type and all other requirements are met (specific pathology findings, staging, etc.); however, the marker must be explicitly recommended in the guidelines and not simply included in a footnote as an intervention that may be considered (See *Common cancer types and associated tumor markers* table below for examples of currently recommended gene tests)

Note If five or more individually billed biomarker tests are under review together (a “panel”) and the member meets the below criteria for a multigene panel, the panel will be approved. However, the laboratory will be redirected to use an appropriate panel CPT code for billing purposes (e.g. 81450).

Companion Diagnostic (CDx) Tumor Marker Panels

Hematological tumor marker companion diagnostic assay panels are considered medically necessary when the member meets ALL of the following criteria:

Member has a diagnosis of cancer, AND

Treatment with a medication for which there is an FDA-approved companion diagnostic assay is being considered, AND

FDA approval for the CDx being requested must include the member's specific cancer type as an approved indication, AND

FDA label for the drug and indication being considered states companion diagnostic testing is necessary for patient selection (See *Common cancer types and associated tumor markers* table below for examples of currently recognized companion diagnostics for available therapies), AND

Member has not had previous somatic and/or germline testing that would have identified the genetic change required to prescribe medication under consideration, AND

Family history:

Member does not have a close (1st or 2nd degree) biological relative with a known germline mutation in a gene that is a target of the requested companion diagnostic test (e.g. known familial mutation in BRCA1/2 and requested test is myChoice CDx), or

Member has a close (1st or 2nd degree) biological relative with a known germline mutation in a gene that is a target of the requested companion diagnostic test (e.g. known familial mutation in BRCA1/2 and requested test is myChoice CDx), and the member's germline test was negative, AND

Rendering laboratory is a qualified provider of service per the Health Plan policy.

Multigene Panel Testing

When a multigene panel is being requested and will be billed with a single panel CPT code (e.g. 81450), the panel will be considered medically necessary when the following criteria are met:

The member has a diagnosis of acute myeloid leukemia (AML), OR

The member has a diagnosis of one of the following cancers, when the panel includes at least five of the genes associated with that cancer type listed in the below table *Common cancer types and associated tumor markers*:

Acute Lymphoblastic Leukemia (ALL)**Hepatosplenic Gamma-Delta T-Cell Lymphoma****Myelodysplastic Syndrome (for cytopenia in which myelodysplasia is suspected),
OR**

All criteria for a multigene panel are met from a test-specific guideline, if available, OR

At least 5 markers included in the panel individually meet criteria for the member's cancer type based on one of the following:

All criteria are met from a test-specific guideline, if available, or

An oncology therapy FDA label requires results from the test to effectively or safely use the therapy for the member's cancer type, or

NCCN guidelines include the test in the management algorithm for that particular cancer type and all other requirements are met (specific pathology findings, staging, etc.); however, the marker must be explicitly included in the guidelines and not simply included in a footnote as an intervention that may be considered.

Note If the member meets criteria for less than 5 of the individual biomarkers in the panel, the panel will not be reimbursed. The laboratory will be redirected to billing for individual tests for which the member meets criteria.

Billing and Reimbursement Consideration

Panels over 50 genes, typically billed with CPT code 81455, are not considered medically necessary, as they are excessive for use in evaluating hematological malignancies and, therefore, are not eligible for reimbursement.

Multigene panels, when medically necessary, will only be considered for reimbursement when billed with an appropriate panel CPT code.

Only one tumor biomarker panel will be considered for reimbursement per occurrence of cancer.

If multiple CDx biomarker panels are ordered simultaneously based on FDA label requirements, only one panel will be considered for reimbursement. Additional unique biomarkers from the second panel may be considered for reimbursement if appropriate single marker or single gene procedure codes are billed.

If a biomarker panel was previously performed and an additional panel is being requested, only testing for the medically necessary, previously untested biomarkers will be reimbursable. Therefore, only the most appropriate procedure codes will be considered for reimbursement.

Other Considerations

For information on somatic mutation testing for solid tumors, please refer to the guideline *Somatic Mutation Testing - Solid Tumors*, as this testing is not addressed here.

For information on tumors markers assayed by liquid biopsy, please refer to the guideline *Liquid Biopsy Testing*, as this testing is not addressed here.

For information on testing for germline (inherited) mutations in genes related to hereditary cancer syndromes (e.g. Hereditary Breast and Ovarian Cancer, Lynch syndrome, etc), please refer to the appropriate test-specific guideline, as this testing is not addressed here. Although some of the same genes may be tested for inherited or acquired (somatic) mutations, this guideline addresses only testing for acquired mutations from hematological malignancies.

Table: Common Cancer Types and Associated Tumor Markers

This list is not all inclusive.

<u>Cancer Type</u>	<u>Tumor Marker</u>	<u>CPT</u>	<u>Indication for Test</u>	<u>Associated Treatments</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>ABL1 Gene Fusion and Kinase Domain Mutation</u>	<u>81170, 81401, 81403</u>	<u>Classification, Diagnostic, Prognostic, Treatment</u>	<u>Bosutinib, Dasatinib, Nilotinib, Ponatinib</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>ABL2 Gene Fusion</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>CRLF2 Gene Fusion</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>CSF1R Gene Fusion</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>EPOR Gene Fusion</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>FLT3 Mutations</u>	<u>81245, 81246</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>

<u>Cancer Type</u>	<u>Tumor Marker</u>	<u>CPT</u>	<u>Indication for Test</u>	<u>Associated Treatments</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>IL7R Mutations</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>JAK1 Mutations</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>JAK2 Gene Fusion and Mutations</u>	<u>81270</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>JAK3 Mutations</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>PDGFRB Gene Fusion</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>Acute Lymphoblastic Leukemia (ALL)</u>	<u>SH2B3 Mutations</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>B-Cell Lymphoblastic Leukemia/Lymphoma</u>	<u>EVT6-RUNX1 Gene Fusion</u>	<u>81401</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>B-Cell Lymphoblastic Leukemia/Lymphoma</u>	<u>IL3-IGH Gene Fusion</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>B-Cell Lymphoblastic Leukemia/Lymphoma</u>	<u>KMT2A Mutations</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>B-Cell Lymphoblastic Leukemia/Lymphoma</u>	<u>TCF3-PBX1 Gene Fusion</u>	<u>81479</u>	<u>Classification, Diagnostic, Prognostic</u>	<u>N/A</u>
<u>Hepatosplenic Gamma-Delta T-Cell Lymphoma</u>	<u>INO80 Mutations</u>	<u>81479</u>	<u>Diagnostic</u>	<u>N/A</u>
<u>Hepatosplenic Gamma-Delta T-Cell Lymphoma</u>	<u>PIK3CD Mutations</u>	<u>81479</u>	<u>Diagnostic</u>	<u>N/A</u>

<u>Cancer Type</u>	<u>Tumor Marker</u>	<u>CPT</u>	<u>Indication for Test</u>	<u>Associated Treatments</u>
<u>Hepatosplenic Gamma-Delta T-Cell Lymphoma</u>	<u>SETD2 Mutations</u>	<u>81479</u>	<u>Diagnostic</u>	<u>N/A</u>
<u>Hepatosplenic Gamma-Delta T-Cell Lymphoma</u>	<u>SMARCA2 Mutations</u>	<u>81479</u>	<u>Diagnostic</u>	<u>N/A</u>
<u>Hepatosplenic Gamma-Delta T-Cell Lymphoma</u>	<u>STAT3 Mutations</u>	<u>81405, 81479</u>	<u>Diagnostic</u>	<u>N/A</u>
<u>Hepatosplenic Gamma-Delta T-Cell Lymphoma</u>	<u>STAT5B Mutations</u>	<u>81479</u>	<u>Diagnostic</u>	<u>N/A</u>
<u>Hepatosplenic Gamma-Delta T-Cell Lymphoma</u>	<u>TET3 Mutations</u>	<u>81479</u>	<u>Diagnostic</u>	<u>N/A</u>
<u>Hepatosplenic Gamma-Delta T-Cell Lymphoma</u>	<u>TCR Mutations</u>	<u>81479</u>	<u>Diagnostic</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>ASXL1 Mutations</u>	<u>81175, 81176</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>BCOR Mutations</u>	<u>81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>CALR Mutations</u>	<u>81219, 81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>CBL Mutations</u>	<u>81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>DDX41 Mutations</u>	<u>81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>DNMT3A Mutations</u>	<u>81403</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>

<u>Cancer Type</u>	<u>Tumor Marker</u>	<u>CPT</u>	<u>Indication for Test</u>	<u>Associated Treatments</u>
<u>Myelodysplastic Syndrome</u>	<u>ETV6 Mutations</u>	<u>81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>EZH2 Mutations</u>	<u>81236, 81237</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>FLT3 Mutations</u>	<u>81245, 81246</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>GATA2 Mutations</u>	<u>81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>IDH1 Mutations</u>	<u>81120, 81403</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>IDH2 Mutations</u>	<u>81121, 81403, 81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>JAK2 Mutations</u>	<u>81270, 81279</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>MPL Mutations</u>	<u>81338, 81339</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>NF1 Mutations</u>	<u>81408</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>NPM1 Mutations</u>	<u>81310</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>NRAS Mutations</u>	<u>81311, 81404, 81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>PHF6 Mutations</u>	<u>81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>

<u>Cancer Type</u>	<u>Tumor Marker</u>	<u>CPT</u>	<u>Indication for Test</u>	<u>Associated Treatments</u>
<u>Myelodysplastic Syndrome</u>	<u>PPM1D Mutations</u>	<u>81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>RUNX1 Mutations</u>	<u>81334, 81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>SETBP1 Mutations</u>	<u>81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>SF3B1 Mutations</u>	<u>81347</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>SRSF2 Mutations</u>	<u>81348</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>STAG2 Mutations</u>	<u>81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>STAT3 Mutations</u>	<u>81405, 81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>TET2 Mutations</u>	<u>81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>TP53 Mutations</u>	<u>81351, 81352</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>U2AF1 Mutations</u>	<u>81357</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>WT1 Mutations</u>	<u>81405, 81479</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>
<u>Myelodysplastic Syndrome</u>	<u>ZRSR2 Mutations</u>	<u>81360</u>	<u>Prognostic (Initial Evaluation)</u>	<u>N/A</u>

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