

Test Catalog

Diagnostic. Prognostic. Predictive. Predisposition.





PML-RARA Translocation, t(15;17)

Alternative Name

PML-RARA Translocation, PML-RARA Fusion

Methodology

Molecular

Test Description

Real-time RT-PCR for quantitative detection of the t(15;17) PML-RARA fusion transcript. Both long and short isoforms of the fusion transcript are detected. Positive results identify the isoform and quantify it as a ratio with the amount of transcript from a normal control gene. Analytical sensitivity is 1 tumor cell in 10,000 normal cells.

Clinical Significance

The (15;17) translocation occurs in nearly all cases of acute promyelocytic leukemia (APL, or AML subtype M3). The translocation is associated with a high rate of complete remission due to sensitivity of leukemic cells to all trans-retinoic acid (ATRA). This assay is recommended for diagnostic confirmation and initiation of ATRA therapy, for monitoring minimal residual disease (MRD), and for detection of relapse

Specimen Requirements

- Bone marrow (preferred): 2 mL in EDTA tube.
- Peripheral blood (acceptable): 5 mL in EDTA tube.

Note: Test is RNA-based. Please select Extract & Hold – RNA if specimen hold service is desired.

Storage & Transportation

Use cold pack for transport, making sure cold pack is not in direct contact with specimen. Ship same day as drawn whenever possible; specimens <7 days old preferred.

CPT Code(s)*

81315

New York Approved

Yes

Level of Service

Global

Turnaround Time

7 days

Please direct any questions regarding coding to the payor being billed.

^{*}The CPT codes provided with our test descriptions are based on AMA guidelines and are for informational purposes only. Correct CPT coding is the sole responsibility of the billing party.

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Committed to research as the means to improve patient care, we provide Pharma Services for pharmaceutical companies, in vitro diagnostic manufacturers, and academic scientist-clinicians. We promote joint publications with our client physicians. NeoGenomics welcomes your inquiries for collaborations. Please contact us for more information.

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