

# Test Catalog

Diagnostic. Prognostic. Predictive. Predisposition.





# **TP53 Mutation Analysis**

#### **Alternative Name**

TP53 Gene Sequencing

# Methodology

Molecular

# **Test Description**

Bi-directional sequencing of TP53 exons 4-9.

# **Clinical Significance**

The TP53 gene encodes the tumor suppressor p53. TP53 mutations are detected in at least 50% of all adult tumors and are generally associated with a poor prognosis. For patients with chronic lymphocytic leukemia (CLL), TP53 sequencing, in addition to FISH for 17p deletion, aids in prognosis and/or therapy selection. Germline mutations in TP53 are the cause of Li-Fraumeni Syndrome.

# **Specimen Requirements**

- Peripheral blood: 5 mL in EDTA tube.
- Bone marrow: 2 mL in EDTA tube.
- FFPE solid tumor tissue: Paraffin block is preferred. Alternatively, send 1 H&E slide plus 5-10 unstained slides cut at 5 or more microns. Please use positively-charged slides and 10% NBF fixative. Do not use zinc fixatives.

Note: Test is DNA-based. Please select Extract & Hold - DNA 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 <14 days old acceptable.

#### CPT Code(s)\*

81352 (as of 01/01/2021); Prior to CPT Code was 81405

## **New York Approved**

No

### **Level of Service**

Global

#### **Turnaround Time**

7 days

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

<sup>\*</sup>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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