

Reporting Title: KIT Asp816Val Mutation Analysis, V

Performing Location: Rochester

Specimen Requirements:

Submit only 1 of the following specimens:

Specimen Type: Whole blood

Container/Tube: Lavender top (EDTA) or yellow top (ACD)

Specimen Volume: 3 mL

Collections Instructions:

1. Invert several times to mix blood.
2. Send specimen in original tube. **Do not** aliquot.
3. Label specimen as blood.

Specimen Stability Information: Ambient (preferred) 7 days/Refrigerate 7 days

Specimen Type: Bone marrow

Container/Tube: Lavender top (EDTA) or yellow top (ACD)

Specimen Volume: 2 mL

Collections Instructions:

1. Invert several times to mix bone marrow.
2. Send specimens in original tube. **Do not** aliquot.
3. Label specimen as bone marrow.

Specimen Stability Information: Ambient (preferred) 7 days/Refrigerate 7 days

Specimen Type: Extracted DNA from blood or bone marrow

Container/Tube: 1.5- to 2-mL tube

Specimen Volume: Entire specimen

Collection Instructions:

1. Label specimen as extracted DNA from blood or bone marrow.
2. Provide indication of volume and concentration of DNA.

Specimen Stability Information: Frozen (preferred)/Refrigerated/Ambient

Forms:

1. [Hematopathology Patient Information](#) (T676)
2. If not ordering electronically, complete, print, and send a [Hematopathology/Cytogenetics Test Request](#) (T726)) with the specimen.

Specimen Type	Temperature	Time	Special Container
Varies	Varies		

Ask at Order Entry (AOE) Questions:

Test ID	Question ID	Description	Type	Reportable
KITVS	MP055	Specimen Type: • Peripheral blood	Answer List	Yes

		<ul style="list-style-type: none">• Bone marrow• Extracted DNA from blood• Extracted DNA from bone marrow		
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Result Codes:

Result ID	Reporting Name	Type	Unit	LOINC®
MP055	Specimen Type	Alphanumeric		31208-2
607982	Interpretation	Alphanumeric		69047-9
607983	Signing Pathologist	Alphanumeric		19139-5

LOINC® and CPT codes are provided by the performing laboratory.

Supplemental Report:

No

CPT Code Information:

81273

Reference Values:

An interpretive report will be provided indicating the mutation status as positive or negative.