



LABORATORY REPORT

Example Client, XYZ123
1234 Warde Road
Ann Arbor MI 48108

EXAMPLE, REPORT W
WX0000003827 M 07/08/1978 45 Y

Referral Testing

Collected: 09/22/2023 13:52 Received: 09/22/2023 13:52

Test Name	Result	Flag	Ref-Ranges	Units	Site
CSF3 Mutation Analysis					
Clinical Indication:	Not Given				QCRL
Specimen Source:	EDTA				QCRL
Block/Specimen ID:	1234567890				QCRL
CSF3R Exon 14/17 Mutation	NOT DETECTED				QCRL

Reference Range:
NOT DETECTED

Gene	.				QCRL
Amino Acid	.				QCRL
Mutation Frequency	.				QCRL
Mutation Type	.				QCRL
Exon	.				QCRL
Nucleotide Change:	.				QCRL
Reference	.				QCRL
Interpretation:	SEE NOTE				QCRL

No mutation is detected in exons 14 or 17 of CSF3R.

Reviewed by Pathologist:

Assay Details	SEE NOTE				QCRL
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This PCR-based advanced sequencing assay interrogates DNA from leukocytes for the presence of mutations in exons 14 and 17 of colony stimulating factor 3 receptor (CSF3R). The sensitivity of mutation detection is approximately 5% but may vary depending on the particular mutation type. Insertions up to 30bp and deletions up to 52bp have been successfully detected by the assay. Alterations outside of the tested areas of this gene will not be detected. Synonymous or known non-synonymous polymorphic changes (SNPs) are not reported. Mutations in these two regions of CSF3R are associated with myeloproliferative neoplasms (MPNS), particularly atypical chronic myeloid leukemia (aCML) and chronic neutrophilic leukemia (CNL). Results of this assay should be correlated with morphology and other laboratory testing for final diagnosis and classification. If this test is negative, additional testing that may be useful for workup of MPNs, depending on presenting hematologic features, includes BCR-ABL1 rearrangement (test code 91065 or 12070X) or mutational analysis of JAK2 V617F (polycythemia vera (PV)/essential thrombocythemia (ET)/primary myelofibrosis (PMF), 92473), CALR (ET/PMF, 92475), JAK2 exon 12 (PV, 92474) or MPL (ET/PMF, 92476). Residual material from this sample may

LAB: L - LOW, H - HIGH, AB - ABNORMAL, C - CRITICAL, . - NOT TESTED

F322000016
WX0000003827
Printed D&T: 09/22/23 13:52

Ordered By: KAJAL SITWALA, MD, PhD
WX0000000002365

Kajal V. Sitwala, MD, PhD - Medical Director
Form: MM RL1
PAGE 1 OF 2



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be used except for BCR-ABL1 testing; call lab to add.

DNA was aligned to GRCh37(hg19) for analysis and transcript ID
ENST00000373103 was used as reference for CSF3R sequence.

For additional information, please refer to
http://education.QuestDiagnostics.com/faq/FAQ211
(This link is being provided for informational/educational purposes
only.)

This test was developed and its analytical performance characteristics
have been determined by Quest Diagnostics Nichols Institute San Juan
Capistrano. It has not been cleared or approved by FDA. This assay has
been validated pursuant to the CLIA regulations and is used for
clinical purposes.

Test Performed at:
Quest Diagnostics Nichols Institute
33608 Ortega Highway
San Juan Capistrano, CA 92675-2042 I Maramica MD, PhD, MBA

QCRL: QUEST DIAGNOSTICS REFERENCE LAB CAPISTRANO 33608 Ortega Highway San Juan Capistrano CA 92675

Reported Date: 2023.09.22 13:52 CSMUT

LAB: L - LOW, H - HIGH, AB - ABNORMAL, C - CRITICAL, . - NOT TESTED

F322000016 Ordered By: KAJAL SITWALA, MD, PhD
WX0000003827 WX00000000002365
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Kajal V. Sitwala, MD, PhD - Medical Director
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