

### Update Notes

Due to reagent shortage Warde will be temporarily sending the tests in this update to Quest. We are providing the test build for the Quest tests during the interim period. We will inactivate the Warde codes (F5L, MTHFR, P2021) on 8/26/20. The Quest code is available to order 8/18/2020. If you have any questions, please contact Warde Client Services.

### Update Summary

<b>Temporarily Inactivate Test With Replacement</b>	8/18/2020	<a href="#">F5L - "Factor V Leiden Mutation Analysis" replaced by FVLMA - "Factor V Leiden Mutation Analysis"</a>
<b>Temporarily Inactivate Test With Replacement</b>	8/18/2020	<a href="#">MTHFR - "MTHFR Genotyping C677T and A1298C" replaced by MTHFA - "Methylenetetrahydrofolate Reductase"</a>
<b>Temporarily Inactivate Test With Replacement</b>	8/18/2020	<a href="#">P2021 - "Prothrombin 20210A Mutation Analysis" replaced by P202A - "Prothrombin (Factor II) 20210G&gt;A"</a>

Inactivate Test With Replacement			
<b>Effective Date</b>	8/18/2020		
Inactivated Test			
<b>Name</b>	Factor V Leiden Mutation Analysis		
<b>Code</b>	F5L		
<b>Legacy Code<sup>1</sup></b>	F5L		
<b>Interface Order Code</b>	3091240		
<b>Notes</b>			
Replacement Test			
<b>Name</b>	Factor V Leiden Mutation Analysis		
<b>Code</b>	FVLMA		
<b>CPT Code(s)</b>	81241 ZB0WL		
<b>Notes</b>			
Specimen Requirements			
<b>Specimen Required</b>	Draw blood in a lavender EDTA. Send 5.0 mL (3.0 mL minimum) in original tube at room temperature.		
<b>Alternate Specimen</b>	Whole blood: Dark blue EDTA, sodium or lithium heparin, and ACD A		
<b>Stability</b>	Room temperature: 8 days; Refrigerated: 8 days; Frozen: 30 days		
Performing Information			
<b>Methodology</b>	Polymerase Chain Reaction and Detection		
<b>Reference Range</b>	See report		
<b>Performed Days</b>	Sunday - Saturday		
<b>Turnaround Time</b>	5 - 7 days		
<b>Performing Laboratory</b>	Quest SJC		
Interface Information			
<b>Legacy Code<sup>1</sup></b>	FVLMA		
<b>Interface Order Code</b>	3400370		
<b>Result Code</b>	<b>Name</b>	<b>LOINC Code</b>	<b>AOE/Prompt<sup>2</sup></b>
3400371	Factor V Leiden Mutation	21668-9	No
3400372	Interpretation	21667-1	No



LABORATORY REPORT

Example Client, XYZ123
1234 Warde Road
Ann Arbor MI 48108

EXAMPLE, REPORT

WX0000003039 M 12/05/1988 31 Y

Referral Testing

Collected: 08/18/2020 10:36 Received: 08/18/2020 10:36

Test Name Result Flag Ref-Ranges Units Site

Factor V Leiden Mutation Analysis

Factor V Leiden Mutation SEE NOTE QCRL

RESULT: FACTOR V LEIDEN (R506Q) VARIANT NOT DETECTED

Interpretation SEE NOTE QCRL

INTERPRETATION: This individual is negative (normal) for the Factor V Leiden (R506Q) variant in the Factor V gene. Increased risk of thrombophilia can be caused by a variety of genetic and non-genetic factors not screened for by this assay.

Laboratory testing supervised and results monitored by Qing Zhang, MD, PhD, FACMG, CGMBS.

MUTATION ANALYSIS:

The Factor V Leiden (R506Q) mutation [NM 000130.2: c.1601G>A (p.R534Q)] in the Factor V gene is one of the most common causes of inherited thrombophilia. This mutation causes resistance to degradation of activated Factor V protein by activated protein C (APC). The Factor V Leiden (R506Q) mutation is detected by amplification of the selected region of Factor V gene by polymerase chain reaction (PCR) and fluorescent probe hybridization to the targeted region, followed by melting curve analysis with a real time PCR system. Although rare, false positive or false negative results may occur. All results should be interpreted in context of clinical findings, relevant history, and other laboratory data.

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.

Health care providers, please contact your local Quest Diagnostics' genetic counselor or call 1-866-GENEINFO (866-436-3463) for assistance with interpretation of these results.

Test Performed at:

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

Performing Site:

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

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

B618000001 Ordered By: CLIENT CLIENT
WX0000003039 WX00000000001595
Printed D&T: 08/18/20 10:37

William G. Finn, M.D. - Medical Director
Form: MM RL1
PAGE 1 OF 1

Inactivate Test With Replacement			
<b>Effective Date</b>	8/18/2020		
Inactivated Test			
<b>Name</b>	MTHFR Genotyping C677T and A1298C		
<b>Code</b>	MTHFR		
<b>Legacy Code<sup>1</sup></b>	MTHFR		
<b>Interface Order Code</b>	3041200		
<b>Notes</b>			
Replacement Test			
<b>Name</b>	Methylenetetrahydrofolate Reductase Mutation		
<b>Code</b>	MTHFA		
<b>CPT Code(s)</b>	81291 ZB0WQ		
<b>Notes</b>			
Specimen Requirements			
<b>Specimen Required</b>	Draw blood in a lavender EDTA. Send 5.0 mL (3.0 mL minimum) in original tube at room temperature.		
<b>Alternate Specimen</b>	Whole blood: Dark blue EDTA, sodium or lithium heparin, and ACD A		
<b>Stability</b>	Room temperature: 8 days; Refrigerated: 8 days; Frozen: 30 days		
Performing Information			
<b>Methodology</b>	Polymerase Chain Reaction and Detection		
<b>Reference Range</b>	See report		
<b>Performed Days</b>	Sunday - Saturday		
<b>Turnaround Time</b>	5 -7 days		
<b>Performing Laboratory</b>	Quest SJC		
Interface Information			
<b>Legacy Code<sup>1</sup></b>	MTHFA		
<b>Interface Order Code</b>	3400501		
Result Code	Name	LOINC Code	AOE/Prompt <sup>2</sup>
3400502	MTHFR Mutation	21709-1	No
3400503	Interpretation	38415-6	No



# LABORATORY REPORT

Example Client, XYZ123  
1234 Warde Road  
Ann Arbor MI 48108

## EXAMPLE, REPORT

WX0000003039 M 12/05/1988 31 Y

### Referral Testing

Collected: 08/18/2020 10:38

Received: 08/18/2020 10:38

Test Name	Result	Flag	Ref-Ranges	Units	Site
-----------	--------	------	------------	-------	------

### Methylenetetrahydrofolate Reductase

MTHFR Mutation	SEE NOTE				QCRL
----------------	----------	--	--	--	------

RESULT: NO VARIANT DETECTED

Interpretation	SEE NOTE				QCRL
----------------	----------	--	--	--	------

INTERPRETATION: This individual is negative (normal) for the variants, C677T and A1298C, in the MTHFR gene. This result is not associated with an increased risk for coronary artery disease, venous thromboembolism, or adverse pregnancy outcome.

Laboratory testing supervised and results monitored by Bernard Joseph Ilagan, MD, MHA, FACMG, CGMBS.

Reduced methylenetetrahydrofolate reductase (MTHFR) enzyme activity is a genetic risk factor for hyperhomocysteinemia, especially when present with low serum folate levels. Two common variants in the MTHFR gene result in reduced enzyme activity. The "thermolabile" variant C677T [NM 005957.3: c.665C>T (p.A222V)] and A1298C [c. 1286A>C (p.E429A)] occur frequently in the general population.

Mild to moderate hyperhomocysteinemia has been identified as a risk factor for coronary artery disease and venous thromboembolism. Hyperhomocysteinemia is multifactorial, involving a combination of genetic, physiologic and environmental factors. Recent studies do not support the previously described association of increased risk for coronary artery disease and venous thromboembolism with mild hyperhomocysteinemia caused by reduced MTHFR activity. Therefore, the utility of MTHFR variant testing is uncertain and is not recommended by The American College of Medical Genetics and Genomics (ACMG) or the American Congress of Obstetricians and Gynecologists (ACOG) in the evaluation of venous thromboembolism or adverse pregnancy outcome.

Modest positive association has also been found between the "thermolabile" variant of the MTHFR gene and many other medical complications, such as recurrent pregnancy loss, risk of offspring with neural tube defects, neuropsychiatric disease, and chemotherapy toxicity. Increased risk of coronary artery disease, venous thromboembolism and increased plasma homocysteine can be caused by a variety of genetic and non-genetic factors not screened for by this assay. If indicated by personal or family history of thromboembolism, consider additional testing such as plasma homocysteine levels, factor V Leiden and prothrombin gene mutations.

The C677T and A1298C variants are detected by amplification of the selected regions of MTHFR gene by polymerase chain reaction (PCR) and

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

B61800002  
WX0000003039  
Printed D&T: 08/18/20 10:40

Ordered By: CLIENT CLIENT  
WX00000000001595

William G. Finn, M.D. - Medical Director  
Form: MM RL1  
PAGE 1 OF 2



LABORATORY REPORT

Example Client, XYZ123
1234 Warde Road
Ann Arbor MI 48108

EXAMPLE, REPORT

WX0000003039 M 12/05/1988 31 Y

Referral Testing

Collected: 08/18/2020 10:38

Received: 08/18/2020 10:38

Test Name Result Flag Ref-Ranges Units Site

fluorescent probes hybridization to the targeted regions, followed by melting curve analysis with a real time PCR system. Although rare, false positive or false negative results may occur. All results should be interpreted in context of clinical findings, relevant history, and other laboratory data. Health care providers, please contact your local Quest Diagnostics' genetic counselor or call 866-GENEINFO (866-436-3463) for assistance with interpretation of these results.

For additional information, please refer to:
http://education.QuestDiagnostics.com/faq/FAQ66
(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

Performing Site:

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

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

B61800002 Ordered By: CLIENT CLIENT
WX0000003039 WX00000000001595
Printed D&T: 08/18/20 10:40

William G. Finn, M.D. - Medical Director
Form: MM RL1
PAGE 2 OF 2

Inactivate Test With Replacement			
<b>Effective Date</b>	8/18/2020		
Inactivated Test			
<b>Name</b>	Prothrombin 20210A Mutation Analysis		
<b>Code</b>	P2021		
<b>Legacy Code<sup>1</sup></b>	PRO2021		
<b>Interface Order Code</b>	3091280		
<b>Notes</b>			
Replacement Test			
<b>Name</b>	Prothrombin (Factor II) 20210G>A Mutation		
<b>Code</b>	P202A		
<b>CPT Code(s)</b>	81240 ZB0WB		
<b>Notes</b>			
Specimen Requirements			
<b>Specimen Required</b>	Draw blood in a lavender EDTA. Send 5.0 mL (3.0 mL minimum) in original tube at room temperature.		
<b>Alternate Specimen</b>	Whole blood: Dark blue EDTA, sodium or lithium heparin, and ACD A		
<b>Stability</b>	Room temperature: 8 days; Refrigerated: 8 days; Frozen: 30 days		
Performing Information			
<b>Methodology</b>	Polymerase Chain Reaction and Detection		
<b>Reference Range</b>	See report		
<b>Performed Days</b>	Sunday - Saturday		
<b>Turnaround Time</b>	5 -7 days		
<b>Performing Laboratory</b>	Quest SJC		
Interface Information			
<b>Legacy Code<sup>1</sup></b>	P202A		
<b>Interface Order Code</b>	3400504		
<b>Result Code</b>	<b>Name</b>	<b>LOINC Code</b>	<b>AOE/Prompt<sup>2</sup></b>
3400505	Prothrombin (Factor II)	24475-6	No
3400506	Interpretation	24476-4	No



LABORATORY REPORT

Example Client, XYZ123
1234 Warde Road
Ann Arbor MI 48108

EXAMPLE, REPORT

WX0000003039 M 12/05/1988 31 Y

Referral Testing

Collected: 08/18/2020 10:41 Received: 08/18/2020 10:41

Test Name Result Flag Ref-Ranges Units Site

Prothrombin (Factor II) 20210G>A

Prothrombin (Factor II) SEE NOTE QCRL

RESULT: G20210A variant not detected

Interpretation SEE NOTE QCRL

INTERPRETATION: This individual is negative (normal) for the G20210A variant in the Prothrombin/Factor II gene. Increased risk of thrombophilia can be caused by a variety of genetic and non-genetic factors not screened for by this assay.

Laboratory testing supervised and results monitored by Carole Oddoux, Ph.D., DABMGG, CGMBS.

The G20210A mutation [AF478696.1: g.21538G>A (c.\*97G>A)] in the Prothrombin/Factor II gene is the second most common inherited risk factor for thrombosis occurring in approximately 2% of Caucasians. Presence of the mutation is associated with an elevation of prothrombin levels to about 30% above normal in heterozygotes and to 70% above normal in homozygotes.

Prothrombin (G20210A) mutations are detected by amplification of their selected gene regions by polymerase chain reaction (PCR) and fluorescent probe hybridization to the targeted region, followed by melting curve analysis with a real time PCR system. Although rare, false positive or false negative results may occur. All results should be interpreted in context of clinical findings, relevant history, and other laboratory data.

Health care providers, please contact your local Quest Diagnostics' genetic counselor or call 1-866-GENEINFO (866-436-3463) for assistance with interpretation of these results.

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

Performing Site:

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

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

B61800003 Ordered By: CLIENT CLIENT
WX0000003039 WX00000000001595
Printed D&T: 08/18/20 10:42

William G. Finn, M.D. - Medical Director
Form: MM RL1
PAGE 1 OF 1