

Client: Example Client ABC123
123 Test Drive
Salt Lake City, UT 84108
UNITED STATES

Physician: Doctor, Example

Patient: Patient, Example

DOB: 12/31/1752
Gender: Unknown
Patient Identifiers: 01234567890ABCD, 012345
Visit Number (FIN): 01234567890ABCD
Collection Date: 01/01/2017 12:34

Hemophilia A (F8) 2 Inversions with Reflex to Sequencing and Reflex to Deletion/Duplication

ARUP test code 2001614

F8 COMP Specimen whole Blood

Symptoms for Hemophilia A (F8) No

Family History for Hemophilia A (F8) No

Hemophilia A (F8) Interpretation See Note

H=High, L=Low, *=Abnormal, C=Critical

Unless otherwise indicated, testing performed at:

Section 79-1 of New York State Civil Rights Law requires informed consent be obtained from patients (or their legal guardians) prior to pursuing genetic testing. These forms must be kept on file by the ordering physician. Consent forms for genetic testing are available at www.aruplab.com. Incidental findings are not reported unless clinically significant but are available upon request.

Inversion Analysis: Negative for pathogenic variants, therefore, F8 sequencing was performed.
Sequencing: Negative for pathogenic variants, therefore, F8 deletion/duplication testing was performed.
Deletion/Duplication Analysis: Negative

TEST PERFORMED - 2001614
TEST DESCRIPTION - Hemophilia A (F8) 2 Inversions with Reflex to Sequencing and Reflex to Deletion/Duplication
INDICATION FOR TEST -Carrier Status

RESULTS
No pathogenic variants were detected in the F8 gene.

INTERPRETATION
According to information provided to ARUP, this individual has a family history of hemophilia. No pathogenic variants in the factor 8 (F8) gene were detected by testing for the intron 22-A and intron 1 inversions, gene sequencing or deletion/duplication analysis. Other F8 variants causing hemophilia A (e.g. deep intronic or promoter variants) have not been excluded. Thus, this patient's risk for being a carrier of hemophilia A is reduced, but not eliminated.

RECOMMENDATIONS
Medical management should rely on family history and clinical findings. If this individual's family members have undergone genetic testing, this negative result must be correlated with the result from an affected relative to confirm that the variant causing disease in this family is detectable by this assay. Genetic and hematologic consultations are recommended.

COMMENTS
Reference Sequence: GenBank # NM_000132.3 (F8)
Nucleotide numbering begins at the "A" of the ATG initiation codon.

This result has been reviewed and approved by [REDACTED]

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Unless otherwise indicated, testing performed at:

ARUP LABORATORIES | 800-522-2787 | aruplab.com
500 Chipeta Way, Salt Lake City, UT 84108-1221
Tracy I. George, MD, Laboratory Director

Patient: Patient, Example
ARUP Accession: 19-120-109548
Patient Identifiers: 01234567890ABCD, 012345
Visit Number (FIN): 01234567890ABCD
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4848

BACKGROUND INFORMATION: Hemophilia A (F8) 2
Inversions with Reflex to
Sequencing and Reflex to
Deletion/Duplication

CHARACTERISTICS: Severe deficiency of factor VIII clotting activity is associated with spontaneous joint or deep tissue bleeding. Moderate or mild deficiency is associated with prolonged bleeding after tooth extractions, surgery, or injuries and recurrent or delayed wound healing.
INCIDENCE: 1 in 4,000-5,000 live male births worldwide; rare in females.

INHERITANCE: X-linked recessive. Of simplex cases, 85 percent of mothers are carriers and 10-15 percent of boys have a de novo pathogenic variant.

PENETRANCE: 100 percent in males and 10 percent in females.

CAUSE: Pathogenic F8 gene variants.

CLINICAL SENSITIVITY: 98 percent.

METHODOLOGY FOR INVERSIONS: F8 intron 22-A and intron 1 inversions detected by inverse PCR and electrophoresis.

METHODOLOGY FOR SEQUENCING: Bidirectional sequencing of the F8 coding region and intron-exon boundaries.

METHODOLOGY FOR DELETION/DUPLICATION: Multiplex ligation-dependent probe amplification (MLPA) to detect large deletions/duplications in the F8 coding region.

ANALYTICAL SENSITIVITY AND SPECIFICITY: 99 percent.

LIMITATIONS: Diagnostic errors can occur due to rare sequence variations. Regulatory region and deep intronic variants, repeat element insertions, and rare F8 intron 22-A and intron 1 inversions with different breakpoints, will not be detected. Deletions/duplications in exon 23 will not be detected.

Test developed and characteristics determined by ARUP Laboratories. See Compliance Statement C: aruplab.com/CS

VERIFIED/REPORTED DATES

Procedure	Accession	Collected	Received	Verified/Reported
F8 COMP Specimen	19-120-109548	4/30/2019 11:00:00 AM	4/30/2019 1:46:24 PM	4/30/2019 2:30:00 PM
Symptoms for Hemophilia A (F8)	19-120-109548	4/30/2019 11:00:00 AM	4/30/2019 1:46:24 PM	4/30/2019 2:30:00 PM
Family History for Hemophilia A (F8)	19-120-109548	4/30/2019 11:00:00 AM	4/30/2019 1:46:24 PM	4/30/2019 2:30:00 PM
Hemophilia A (F8) Interpretation	19-120-109548	4/30/2019 11:00:00 AM	4/30/2019 1:46:24 PM	4/30/2019 2:30:00 PM

END OF CHART

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