

Patient ID <b>SA00000808</b>	Patient Name <b>SAMPLEREPDMLM, VLD20150713A0068</b>	Birth Date <b>1981-01-01</b>	Gender <b>M</b>	Age <b>34</b>
Order Number <b>SA00000808</b>	Client Order Number <b>SA00000808</b>	Ordering Physician <b>CLIENT, CLIENT</b>	Report Notes	
Account Information <b>C7028846 DLMP Rochester</b>		Collected <b>12 Jul 2015 14:55</b>		

## Huntington Disease Analysis

### Result Summary

FULL MUTATION IDENTIFIED

MCR

ClinGen. This may not be applicable for all tests.

### Result

CAG repeat: 45 (Full penetrance) and 13

MCR

Test results should be interpreted in the context of clinical findings, family history, and other laboratory data. Misinterpretation of results may occur if the information provided is inaccurate or incomplete.

### Interpretation

1 MCR

This result is consistent with a diagnosis of Huntington disease.

Rare polymorphisms exist that could lead to false-negative or false-positive results. If results obtained do not match the clinical findings, additional testing should be considered.

Identification of a mutation has important implications for family members. Testing of at risk individuals is possible, but it is recommended that predictive testing be performed in conjunction with appropriate pre- and post-test counseling.

Bone Marrow transplants from allogenic donors will interfere with testing. Call Mayo Clinic Laboratories for instructions for testing patients who have received a bone marrow transplant.

A genetic consultation may be of benefit.

Multiple in-silico evaluation tools may have been used to assist in the interpretation of these results. Of note, the sensitivity and specificity of these tools for the determination of pathogenicity is currently unvalidated.

### ADDITIONAL INFORMATION

A PCR-based assay was utilized to detect CAG repeat expansions in exon 1 of the HTT gene. We estimate that the number of CAG repeats is correct within  $\pm 5\%$ .

### Reason for Referral

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Evaluation for a diagnosis of Huntington disease (HD). Test for the presence of an expansion in the HTT gene.

Normal: <27

Intermediate: 27–35

Reduced penetrance: 36–39

Full penetrance: >39

An online research opportunity called GenomeConnect (genomeconnect.org), a project of ClinGen, is available for the recipient of this genetic test. This patient registry collects de-identified genetic and health information to advance the knowledge of genetic variants. Mayo Clinic is a collaborator of

### Specimen

MCR

WB Whole Blood

### Released By

MCR

EMILY LAUER

**Received:** 13 Jul 2015 20:20

**Reported:** 24 Jul 2015 12:47

### Laboratory Notes

- 1 This test was developed and its performance characteristics determined by Mayo Clinic in a manner consistent with CLIA requirements. This test has not been cleared or approved by the U.S. Food and Drug Administration.

### Performing Site Legend

Code	Laboratory	Address	Lab Director	CLIA Certificate
MCR	Mayo Clinic Laboratories - Rochester Main Campus	200 First Street SW, Rochester, MN 55905	William G. Morice M.D. Ph.D	24D0404292