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500 Chipeta Way, Salt Lake City, Utah 84108-1221 phone: 801-583-2787, toll free: 800-522-2787 Jonathan R. Genzen, MD, PhD, Chief Medical Officer

Patient Age/Sex: 8 hours Unknown

Specimen Collected: 07-Nov-22 08:23

CMV Drug Resistance by NGS, Letermovir	Received: 07-Nov-2	2 08:23	Report/Verified: 07-Nov-22 13:50
Procedure	Result	Units	Reference Interval
CMV Drug Resistance by NGS,	See Note ^{fl il}		
Letermovir			

Result Footnote

fl: CMV Drug Resistance by NGS, Letermovir Letermovir,LTV Sensitive

> UL56 drug resistance mutations identified: None UL56 additional mutations identified:V425A, N586D UL56 uncalled mutation sites:None

CMVResistanceCaller software version: 2.0.0.1

CMV_resistance_mutations_20220321.db

Test Information

i1: CMV Drug Resistance by NGS, Letermovir INTERPRETIVE INFORMATION: CMV Drug Resistance by NGS, Letermovir

This assay assesses resistance to letermovir. Resistance-associated mutations in the UL56 gene are sequenced using next generation sequencing. Drug resistance is assigned using an ARUP-developed database of published resistance mutations. For a list of resistance mutations refer to https://ltd.aruplab.com/Tests/Pub/3004509.

This test detects populations down to 10% of the total population which may account for resistance interpretation differences between methods. Some insertions or deletions may be difficult to detect using this software.

Result interpretations are as follows:

-Sensitive indicates no evidence of drug resistance compared with a wild-type virus. -Possible resistance indicates mutations were detected with borderline-level drug resistance or conflicting resistance status reported in the literature.

-Resistant indicates evidence of drug resistance compared with a wild-type virus. -Not determined indicates incomplete sequence coverage across a given gene or genes. -Additional mutations include variants that have not been associated with drug resistance.

-Uncalled mutation sites include drug resistance mutation positions with an inadequate number of sequencing reads.

-Inadequate sequence coverage indicates a low number of sequence reads at a given drug resistance site.

This test was developed, and its performance characteristics determined by ARUP Laboratories. It has not been cleared or approved by the U.S. Food and Drug

*=Abnormal, #=Corrected, C=Critical, f=Result Footnote, H-High, i-Test Information, L-Low, t-Interpretive Text, @=Performing lab

Unless otherwise indicated, testing performed at: ARUP Laboratories 500 Chipeta Way, Salt Lake City, UT 84108 Laboratory Director: Jonathan R. Genzen, MD, PhD
 ARUP Accession:
 22-311-100988

 Report Request ID:
 16971657

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 08-Nov-22 14:46

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Patient Age/Sex: 8 hours Unknown

Test Information

i1: CMV Drug Resistance by NGS, Letermovir Administration. This test was performed in a CLIA-certified laboratory and is intended for clinical purposes.

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