



Units



Name	: XXXXXXXXXX	Referred By	: XXXXXXXXXX
ld	: XXXXXXXXXX	Billed	: XXXXXXXXXX
Age	: XX/Y	Collected On	: XXXXXXXXXX
Gender	: X	Reported	: XXXXXXXXXX
Phone	: XXXXXXXXXX	Vid	: XXXXXXXXXX

Test

Result

Biological Reference Interval

DEPARTMENT OF MOLECULARBIOLOGY - BLOOD (EDTA)

HHH Panel (1455)

(Method: MULTIPLEX RT PCR)

Specimen	EDTA(Whole Blood)	
HIV RNA Qualitative	NEGATIVE	
HCV RNA Qualitative	NEGATIVE	
HBV DNA Qualitative	NEGATIVE	

REMARKS

Clinical Significance: 1. Hepatitis B is a potentially life-threatening liver infection caused by the Hepatitis B virus (HBV). 2. HBV is a major global health problem. It can cause chronic infection and puts people at high risk of death from cirrhosis and liver cancer.

3. Hepatitis C is a liver disease caused by the Hepatitis C virus. The disease can range in severity from a mild illness lasting a few weeks to a serious, lifelong condition that can lead to cirrhosis of the liver or liver cancer.

4. The hepatitis C virus is transmitted through contact with the blood of an infected person. HCV infection is highest among past or present intravenous drug abusers.
5. Hepatitis C is an infectious disease affecting the live, caused by the Hepatitis C virus (HCV). This infection is often asymptomatic, but once established it can progress to chronic scarring of the liver (fibrosis and cirrhosis).
6. HIV-1 is prevalent globally with 2.1 million people living with HIV(PL-HIV). India has largest number of PL-HIV in Asia. Currently three main genetic groups are present for HIV-1: Group M(Main), Group O(Outlier), Group N(Non-M and Non-O).
7. HIV-2 is distinct retrovirus, but has same mode of transmission that of HIV-1.
8. HBV DNA (Genetyme A - G) HCV RNA (Genetyme A - G) HCV RNA (HIV PLAID).

8. HBV DNA (Genotype A - G), HCV RNA (Genotype 1 - 6) and HIV RNA (HIV-1 Group M and O and HIV-2) were detectable by this PCR technology.

9. This test does not discriminate between HIV-1(Group- M and Group-O) and HIV-2.

INTERPRETATION	

RESULT				· · · · · · · · · · · · · · · · · · ·	
NEGATIVE		HBV DNA, HCV RN specimen was not dete limit.	A and HIV RNA ected or less than	h in the given In the detection	
HBV POSITIVE		HBV DNA is Detected in	n the given samp	le.	
HCV POSITIVE		HCV RNA is Detected in	n the given samp	ole.	
HIV POSITIVE		HIV RNA is Detected in	the given sample	e.	
HBV DNA Qualitative (855) (Method: RT-PCR)					
SPECIMEN	WHO	LE BLOOD (EDTA)			
RESULT					
HBV DNA QUALITATIVE	DETE	CTED			
INTERPRETATION					
RESULT					

Disclaimer :

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Test			Result		Units	Biological Reference Interval
Phone	:	XXXXXXXXXX		Vid	:	XXXXXXXXXX
Gender	:	Х		Reported	:	XXXXXXXXXX
Age	:	XX/Y		Collected On	:	XXXXXXXXXX
ld	:	XXXXXXXXXX		Billed	:	XXXXXXXXXX
Name	:	XXXXXXXXXX		Referred By	:	XXXXXXXXXX

NOT DETECTED	HBV DNA in the given specimen was not detected or less than the detection limit.
DETECTED	HBV DNA is detected in the given specimen

PATHOGEN INFORMATION:

HBV Primer and Probe have been designed for the specific and exclusive in-vitro quantification of HBV virus in clinical specimens.

Clinical Significance:

1. Hepatitis B is a potentially life-threatening liver infection caused by the hepatitis B virus (HBV). 2. It is a major global health problem. It can cause chronic infection and puts people at high risk of death from cirrhosis and liver cancer.

HBV DNA QUANTIFICATION (061)

(Method: Real Time RT-PCR)

	- /			
SPECIMEN TYPE		WHOLE BLOOD (EDTA)		
RESULT				
HBV DNA QUANTIFICATION		DETECTED		
		1000	IU/mL	
INTERPRETATION				
RESULT				
NOT DETECTED	HBV DNA in the g	iven specimen was not detected or le	ss than the detection limit.	

DETECTED	HBV DNA is detected in the given specimen
DEIEGIED	HBV DNA IS detected in the diven specimen

REPORTABLE RANGE	30 IU/mL to 3 x 10 ⁹ IU/mL
GENOTYPES	А-Н

PATHOGEN INFORMATION:

HBV primer and Probe have been designed for the specific and exclusive in vitro quantification of HBV virus.

Clinical Significance:

1. Hepatitis B is a potentially life-threatening liver infection caused by the hepatitis B virus (HBV). 2. It is a major global health problem. It can cause chronic infection and puts people at high risk of death from cirrhosis and liver cancer.

HIV RNA Qualitative PCR (854) (Method: Real Time RT-PCR)

SPECIMEN

WHOLE BLOOD (EDTA)

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Test	Result	Units	Biological Reference Interval
RESULT			
HIV RNA Qualitative	NOT DETECTED		

REMARKS

The specificity of the Artus HI Virus-1 RG RT-PCR Kit is first and foremost ensured by the selection of the primers and probes, as well as the selection of stringent reaction conditions. The primers and probes were checked for possible homologies to all in gene banks published sequences by sequence comparison analysis. The detectability of all relevant genotypes has thus been ensured by a database alignment and by a PCR run on Rotor-Gene Instruments.

INTERPRETATION

This assay should not be used as a screening test or primary diagnostic test for HIV-1 infection, except in infants <18 months of age born to HIV-1 infected mothers.

This assay is optimized for the detection of group M subtypes (A to H), N and O, but it may not detect all HIV-1 group N or O strains Diagnosis of HIV-1 infection should not rely solely upon a Detected result for HIV-1 DNA and/or RNA. Such a result should be considered in conjunction with a patient's clinical presentation, physical findings and other diagnostic laboratory tests prior to establishing a diagnosis.

Undetected results should be interpreted with caution, considering the patient's risk factors for HIV-1 infection, the analytical sensitivity of the assay, and the group of the infecting HIV-1 strain. Follow up testing is recommended for high-risk patients with initially Undetected test results.

Undetected result together with repeatedly positive HIV-1 antibody supplemental test results may be observed in HIV-2 infected individuals. For such patients with risk factors for HIV-2 infection, specific testing for HIV-2 antibodies (serologic) and HIV-2 DNA and/or RNA is recommended.

HIV I RNA QUANTITATIVE PCR (660)

(Method: Real Time RT-PCR)

Specimen	WHOLE BLOOD (EDTA)	
RESULT		
HIV I RNA QUANTITAIVE PCR	NOT DETECTED	
	N/A	

REMARKS

The specificity of the Artus HI Virus-1 RG RT-PCR Kit is first and foremost ensured by the selection of the primers and probes, as well as the selection of stringent reaction conditions. The primers and probes were checked for possible homologies to all in gene banks published sequences by sequence comparison analysis. The detectability of all relevant genotypes has thus been ensured by a database alignment and by a PCR run on Rotor-Gene Instruments.

INTERPRETATION

This assay should not be used as a screening test or primary diagnostic test for HIV-1 infection, except in infants <18 months of age born to HIV-1 infected mothers.

This assay is optimized for the detection of group M subtypes (A to H), N and O, but it may not detect all HIV-1 group N or O strains. Diagnosis of HIV-1 infection should not rely solely upon a Detected result for HIV-1 DNA and/or RNA. Such a result should be considered in conjunction with a patient's clinical presentation, physical findings and other diagnostic laboratory tests prior to establishing a diagnosis.

Undetected results should be interpreted with caution, considering the patient's risk factors for HIV-1 infection, the analytical sensitivity of the assay, and the group of the infecting HIV-1 strain. Follow up testing is recommended for high-risk patients with initially Undetected test results.

Undetected result together with repeatedly positive HIV-1 antibody supplemental test results may be observed in HIV-2 infected individuals. For such patients with risk factors for HIV-2 infection, specific testing for HIV-2 antibodies (serologic) and HIV-2 DNA and/or RNA is recommended.

The performance of this test has been validated at department of Molecular Biology Laboratory.

RESULT

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Units

Biological Reference Interval

Name	: XXXXXXXXXX	Referred By	: XXXXXXXXX	
ld	: XXXXXXXXXX	Billed	: XXXXXXXXXX	
Age	: XX/Y	Collected On	: XXXXXXXXXX	
Gender	: X	Reported	: XXXXXXXXXX	
Phone	: XXXXXXXXXX	Vid	: XXXXXXXXXX	

Test

Result

NOT DETECTED	HIV RNA in the given specimen was not detected or less than the detection limit
DETECTED	HIV RNA is detected in the given specimen
REPORTABLE RANGE	120 IU/mL to 1 x 10 ⁸ IU/mL

HCV RNA QUALITATIVE PCR ((Method: Real Time RT-PCR)	262)	
Specimen	WHOLE BLOOD (EDTA)	
RESULT		
HCV RNA	NEGATIVE	
RESULT		

RESOLI		
NEGATIVE	HCV RNA in the given specimen was not detected or less than the detection limit.	
POSITIVE	HCV RNA is detected in the given sample.	

PATHOGEN INFORMATION:

HCV diagnosis is based on the amplification of 3' UTR conserved region of HCV genome covering subtypes 1-6.

Clinical Significance:

Hepatitis C is a liver disease caused by the hepatitis C virus. The disease can range in severity from a mild illness lasting a few weeks to aserious, lifelong condition that can lead to cirrhosis of the liver or liver cancer.

2. The hepatitis C virus is transmitted through contact with the blood of an infected person. HCV infection is highest among past or presentintravenous drug abusers.

Hepatitis C is an infectious disease affecting the live, caused by the Hepatitis C virus (HCV). This infection is often asymptomatic, but once established it can progress to chronic scarring of the liver (fibrosis and cirrhosis).
 Viral load test results have many uses, such as confirming active HCV infection, and measuring HCV treatment response before, during and after therapy. Viral load determination cannot be correlated with the risk of sexual transmission.
 There are 6 genotypes of hepatitis C and they may respond differently to treatment. Careful screening is necessary before starting thetreatment to determine the most appropriate approach for the patient.

HCV RNA QUANTIFICATION (649)

(Method: Real Time RT PCR)

RESULT				
HCV RNA QUANTIFICA	ATION	NOT DETECTED		
		N/A		
INTERPRETATION		· · ·	 ·	
RESULT				
NOT DETECTED	HCV RNA in the given specimen was not detected or less than the detection limit.			

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Units

Name	: XXXXXXXXXX	Referred By	: XXXXXXXXXX
ld	: XXXXXXXXXX	Billed	: XXXXXXXXXX
Age	: XX/Y	Collected On	: XXXXXXXXXX
Gender	: X	Reported	: XXXXXXXXXX
Phone	: XXXXXXXXXX	Vid	: XXXXXXXXX

Test

Result

DETECTED

HCV RNA is detected in the given specimen

REPORTABLE RANGE	30 IU/mL to 10 ⁹ IU/mL
GENOTYPES	1 - 6

PATHOGEN INFORMATION:

HCV diagnosis is based on the amplification of 3' UTR conserved region of HCV genome covering subtypes 1-6.

Clinical Significance:

1. Hepatitis C is a liver disease caused by the Hepatitis C virus. The disease can range in severity from a mild illness lasting a few weeks to a serious lifelong condition that can lead to cirrhosis of the liver or liver cancer.

2. The Hepatitis C virus is transmitted through contact with the blood of an infected person. HCV infection is highest among past or present intravenous drug abusers.

Hepatitis C is an infectious disease affecting the live, caused by the Hepatitis C virus (HCV). This infection is often asymptomatic, but once established it can progress to chronic scarring of the liver (fibrosis and cirrhosis).
 Viral load test results have many uses, such as confirming active HCV infection, and measuring HCV treatment response before, during and after therapy. Viral load determination cannot be correlated with the risk of sexual transmission.

5. There are 6 genotypes of Hepatitis C and they may respond differently to treatment. Careful screening is necessary before starting the treatment to determine the most appropriate approach for the patient.

CMV DNA PCR Qualitative (852)

(Method: RT-PCR)

Specimen	BLOOD	
CMV DNA Qualitative		
Result	NOT DETECTED	

REMARKS

Human cytomegalovirus: Cytomegalovirus (CMV) formally designated as Human Herpes Virus 5 (HHV-5) belongs to the family Herpes viridae. It has a worldwide distribution and infects humans of all ages with no seasonal or epidemic patterns of transmission. Seroprevalence of CMV increases with age ranging from 40-100%; highest being among lower socioeconomic groups. The infections can be congenital, perinatal or postnatal.

CMV quantitative PCR (1483) (Method: RT PCR) **SPECIMEN BLOOD** RESULT **CMV DNA QUANTIFICATION** NOT DETECTED N/A **INTERPRETATION** PATHOGEN INFORMATION:

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Biological Reference Interval





Test		Result	Units	Biological Reference Interval
Phone	: XXXXXXXXXX	Vid		: XXXXXXXXXX
Gender	: X	Rej	ported	: XXXXXXXXXX
Age	: XX/Y	Col	lected On	: XXXXXXXXXX
ld	: XXXXXXXXXXX	Bill	ed	: XXXXXXXXXX
Name	: XXXXXXXXXX	Ref	erred By	: XXXXXXXXXX

Test

CMV primer and Probe have been designed for the specific and exclusive in vitro quantification of CMV virus.

Clinical Significance:

CMV, also known as herpesvirus-5 (HHV-5), belongs to the Herpesviridae family.
 CMV is mostly asymptomatic in healthy people, immunocompromised patients develop a mononucleosis-like syndrome with prolonged fever, mild hepatitis, sore throat and inflammation of the lymph nodes.

REPORTABLE RANGE	77 IU/mL to 7.94×10^7 IU/mL
NOT DETECTED	CMV DNA in the given specimen was not detected or less than the detection limit.
DETECTED	CMV DNA is detected in the given specimen



--- End of the Report ---

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