

IL28B SNP genotyping

Dr. Shamma Shetye

- HCV ---
- HCV diagnosis—
- HCV viral load---
- HCV genotyping---
- Genotypes 1,4---24-48 weeks
- Genotypes 2,3---12-24 weeks
- SVR, Sustained virologic response
- **Week 12 <2log dec, Week 24 detect**

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Table 2. Monitoring of on-therapy response to PEG IFN plus ribavirin.

Sustained virological response (SVR)	Undetectable HCV RNA level (<50 IU/ml), 24 weeks after treatment
Rapid virological response (RVR)	Undetectable HCV RNA in a sensitive assay (lower limit of detection ≤50 IU/ml) at week 4 of therapy, maintained up to end of treatment
Early virological response (EVR)	HCV RNA detectable at week 4 but undetectable at week 12, maintained up to end of treatment
Delayed virological response (DVR)	More than 2 log ₁₀ drop but detectable HCV RNA at week 12, HCV RNA undetectable at week 24, maintained up to end of treatment
Null response (NR)	Less than 2 log ₁₀ IU/ml decrease in HCV RNA level from baseline at 12 weeks of therapy
Partial nonresponse (PR)	More than 2 log ₁₀ IU/ml decrease in HCV RNA level from baseline at 12 weeks of therapy but detectable HCV RNA at weeks 12 and 24
Breakthrough (BT)	Reappearance of HCV RNA at any time during treatment after virological response

Treatment should be stopped at week 12 if the HCV RNA decrease is less than 2 log₁₀ IU/ml, i.e. if the baseline HCV RNA level is reduced by less than 99% of the baseline value, as the SVR rate in these patients with standard treatment duration is less than 2%. In patients with detectable HCV RNA (≥50 IU/ml) at week 24, treatment should also be stopped due to a minimal chance of SVR (1–3%) [55,78,91,92].

Patients with a more than 2 log₁₀ drop or an undetectable HCV RNA at week 12 can be classified into three groups according to their virological response (Table 2): (1) rapid virologic response

- (iv) In patients infected with HCV genotypes 2 and 3 with an RVR and low baseline viral load (<400,000–800,000 IU/ml), shortening of treatment duration to 16 weeks can be considered at the expense of a slightly higher chance of post-treatment relapse [81,101–104].
- (v) In patients with HCV genotypes 2 and 3 who have advanced fibrosis, cirrhosis or cofactors affecting response (insulin resistance, metabolic syndrome, non-viral steatosis) shortening of treatment duration to 16 weeks should not be considered, even if they have low baseline viral

- Interleukin 28B gene present on chromosome 19.
- SNP single nucleotide polymorphism
- Alleles...C or T,
- CC
- CT
- TT

TEST REPORT

DINESH KUMAR

Reference: Dr. SAMIR SHAH

Sample Collected At:
DR MEETA ADHIYA
410 DOCTOR HOUSE , 14 PEDDER ROAD
MUMBAI 400026
Zone 24

SID: 1116
Collecte
16/11/2011 08:0
Registere
16/11/2011 08:0
Reporte
07/12/2011 05:1

Age:48.00 Years Sex:MALE

IL28B SNP Genotyping Test

TEST NAME:

IL28B SNP Genotyping Test

METHOD:

DNA isolation followed by PCR amplification and SNP genotyping
by direct sequencing

TEST RESULT:

GENOTYPE "TT" DETECTED

TEST INDICATION:

IL28B gene screening in patients with HCV infection

TEST DETAILS:

rs12979860 is a SNP near the IL28B gene, encoding interferon lambda 3 (IFN lambda 3)

IL28B SNP Genotyping Test

TEST NAME:	IL28B SNP Genotyping Test
METHOD:	DNA isolation followed by PCR amplification and SNP genotyping by direct sequencing
TEST RESULT:	GENOTYPE " CC " DETECTED
TEST INDICATION:	IL28B gene screening in patients with HCV infection

TEST DETAILS:

- * **rs12979860 is a SNP near the IL28B gene, encoding interferon-lambda-3 (IFN-lambda-3).**
- * The IL28b SNP genotyping test detects the rs12979860 C/T variant upstream of the IL28B gene.
The presence of cytosine (C) is associated with an approximate two-fold improved response rate across ethnicities compared to thymine (T) at the same position.
- * **rs12979860 is the critical predictor for rapid, early and sustained virological response along with end of treatment in patients of GT1-HCV chronic infection.**
- * **This test has been developed and its performance validated at Molecular Biology Department, Metropolis Healthcare Ltd.**

CLINICAL SIGNIFICANCE:

Single nucleotide polymorphisms (SNPs) of interleukin-28B (IL28B) have received considerable interest for their association with sustained virological response (SVR) when treating patients of genotype-1 hepatitis C virus (GT1-HCV) chronic infection with pegylated interferon and ribavirin (PegIFN/RBV). Patients with rs12979860 SNP genotypes CC respond to treatment at least 2 - 3 fold more likely when compared to patients with CT / TT genotypes.

LIMITATION OF ASSAY:

Presence of PCR inhibitors in the sample prevents DNA amplification for IL28b SNP Genotype.

REFERENCE:

1. Thomas DL, Thio CL, Martin MP, et. al., 2009. Nature 461: 798-801.
2. Alestig E, Arnholm B, Eilard A, et. al., 2011. BMC Infectious Diseases 11:124

TEST REPORT

ASGAR (MAM-1334812)

Reference Dr.

Sample Collected At:
R & D
REX CHEMBURS, ROOM#33 GROUND FLR,
WH MARG MUMBAI 01,
Zone 007

SID: 111850783
Collected On:
07/02/2012 11:18 AM
Registered On:
07/02/2012 11:18 AM
Reported On:
10/02/2012 05:09 PM

Age:35.00 Years Sex:MALE

IL28B SNP Genotyping Test

TEST NAME: IL28B SNP Genotyping Test
METHOD: DNA isolation followed by PCR amplification and SNP genotyping by direct sequencing
TEST RESULT: GENOTYPE "CT" DETECTED
TEST INDICATION: IL28B gene screening in patients with HCV infection

TEST DETAILS:

- rs12979860 is a SNP near the IL28B gene, encoding Interferon-lambda-3 (IFN-lambda-3).
- The IL28B SNP genotyping test detects the rs12979860 C/T variant upstream of the IL28B gene. **The presence of cytosine (C) is associated with an approximate two-fold improved response rate across ethnicities compared to thymine (T) at the same position.**
- **rs12979860 is the critical predictor for rapid, early and sustained virological response along with end of treatment in patients of GT1-HCV chronic infection.**
- **This test has been developed and its performance validated at Molecular Biology Department, Metropolis Healthcare Ltd.**

CLINICAL SIGNIFICANCE:

Single nucleotide polymorphisms (SNPs) of Interleukin-28B (IL28B) have received considerable interest for their association with sustained virological response (SVR) when treating patients of genotype-1 hepatitis C virus (GT1-HCV) chronic infection with pegylated interferon and ribavirin (PegIFN/RBV). Patients with rs12979860 SNP genotypes CC respond to treatment atleast 2 - 3 fold more likely when compared to patients with CT / TT genotypes.

LIMITATION OF ASSAY:

Presence of PCR inhibitors in the sample prevents DNA amplification for IL28B SNP Genotype.

REFERENCE:

1. Thomas DL, Thilo CL, Martin MP, et. al., 2009. Nature 461: 798-801.
2. Aicardi E, Amholm B, Ellard A, et. al., 2011. BMC Infectious Diseases 11:124

End of Report

MAM, Amherst



Refer to conditions of reporting overleaf

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Pravghade
DR. SAVIDEEP WARGHADE
MBA, MD (PATHOLOGY)

**Referenced Test

Results relate only to the sample as received



- **Test Details:**
- **Sample Type:** 2 ml EDTA whole Blood
- **Transportation:** Whole blood is shipped in ambient (18-26 ° C) temperature. Do not freeze whole blood.
Methodology: Conventional PCR followed by sequencing
- **Turn around Time:** Daily batch and reporting 10th Day.

- IL28B is a strong predictor of treatment outcome of chronic hepatitis C.
- Studies have shown that patients with CC genotype have twofold likelihood of SVR as compared to CT and TT.
- CC also correlates with a higher chance of spontaneous virus clearance.
- Most studies are with HCV-1, but also with HCV 2 and 3 (and 4).

Thank you