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# Role of IL-28B polymorphisms in virologic response to combined pegylated interferon and ribavirin therapy in genotype 4 chronic HCV infected patients with and without cirrhosis



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# **KEYWORDS**

HCV; IL28B; Polymorphism; Interferon; Schistosomiasis; Response **Abstract** *Background:* Chronic hepatitis C virus (HCV) represents one of the common causes of chronic liver disease worldwide with Egypt having the highest prevalence, namely genotype 4. The rs12979860 CC genotype of the interleukin 28B (IL28B) polymorphisms is associated with high rates of sustained virological response to pegylated interferon and ribavirin in HCV genotype-1 patients. Data on other genotypes are more limited.

*Objective:* We aim to evaluate the predictive power of the rs12979860 IL28B single nucleotide polymorphisms for treatment response at 3 and 6 months in chronic HCV genotype 4 Egyptian patients in relation to other predictors.

Patients and methods: The study included 60 chronic HCV Egyptian patients receiving pegylated interferon and ribavirin therapy. Patients were classified into 2 groups; 30 patients with compensated cirrhosis, and 30 patients without cirrhosis. We analyzed selected pretreatment factors such as age, sex, HCV viral load, anti-schistosomal antibodies, insulin resistance, alpha fetoprotein, low and high density lipoproteins and single nucleotide polymorphisms of IL28B and tried to find out which of them influence sustained virological response.

Results: In univariate analysis, CC genotype showed a significant association with sustained virological response at 6 months among the cirrhotic patients (81.8% responders had the CC genotype, 58.3% had the CT/TT genotype) (p = 0.009). While in multivariate analysis, the presence

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of cirrhosis showed higher risk of failed response at 3 and 6 months (p = 0.016 and 0.020 respectively). Also, positive schistosoma serology was an important negative predictor of response at 3 and 6 months in both groups (p = 0.003 and 0.001 respectively).

Conclusion: In Egypt, where chronic HCV genotype 4 and schistosoma coinfection predominate, both schistosoma infection and cirrhosis are more potent than IL28B polymorphisms as strong baseline negative predictors of hepatitis C treatment response.

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### 1. Introduction

Hepatitis C virus (HCV) infection is one of the main causes of chronic liver disease worldwide. The long-term impact of HCV infection is highly variable, from minimal changes to extensive fibrosis and cirrhosis with or without hepatocellular carcinoma (HCC). The number of chronically infected persons worldwide is estimated to be about 160 million, but most of them are unaware of their infection. <sup>1</sup>

In fact, Egypt has the largest epidemic of HCV in the world with an overall serum positive prevalence of 14.7% as reported by the Egyptian demographic health survey, with >90% of cases having HCV genotype 4.<sup>2,3</sup> Hepatitis C virus (HCV) genotype 4 is the most frequent cause of chronic hepatitis C in the Middle East, North Africa, and sub-Saharan Africa.<sup>4</sup> It has been reported to be frequently associated with cirrhosis and a poor response to interferon (IFN).<sup>5,6</sup> Furthermore, epidemiological reports indicate that HCV genotype 4 is beginning to spread from its native African and Middle Eastern origins to countries of Southern Europe such as France, Italy and Spain and in some foci in the United States, particularly among intravenous drug users.<sup>7–9</sup>

Chronic hepatitis C proceeds toward cirrhosis over several decades. On average, 10 to 20% of patients develop cirrhosis over 20–30 years of infection. Until 2011, treatment with pegylated interferon (PEG-IFN) and ribavirin (RBV) was the standard of care (SOC) for all HCV genotypes, with HCV clearance depending both on virus and host-related factors.

It has been reported that treatment with conventional IFN is less effective in patients with genotypes 1 and 4 than in patients with genotypes 2 and 3. 11 European Association for the Study of the Liver recommended that liver disease severity should be assessed prior to therapy. Identifying patients with cirrhosis is of particular importance, as their prognosis, their likelihood of response and the duration of therapy are altered. 1

Currently the treatment of chronic hepatitis is inclined toward individualized therapy, based on the knowledge of factors predicting response to treatment. All treatment-naïve patients with compensated disease due to HCV should be considered for therapy. The goal of therapy is to eradicate HCV infection to prevent liver cirrhosis, HCC, and death. The endpoint of therapy is undetectable HCV RNA in a sensitive assay (<15 IU/ml) 12 and 24 weeks after the end of treatment (i.e. sustained virologic response SVR12 and SVR24). In patients with cirrhosis, HCV eradication reduces the rate of decompensation and will reduce, albeit not abolish the risk of HCC. In these patients screening for HCC should be continued.

Several independent genome-wide association studies (GWAS) reported single nucleotide polymorphisms (SNPs) rs12979860, located 3 kilobases upstream of the IL28B gene,

encodes a type III IFN (IFN-3) is associated with more than a 2-fold difference in the rate of SVR. 6 CC genotype of IL-28B is associated with a 2–3-fold increase in sustained virologic response (SVR) compared to CT or TT genotype. 13,14

The mechanism by which SNPs influence the outcome of HCV infection and its treatment is not clear. It is suggested that regulation of the promoter region of IL28B in antiviral activity may also affect two other genes belonging to interferon (IFN)  $\lambda$  family encoded in this region. There are a few data so far regarding the role of IL28B polymorphism in HCV-4 patients with respect to response to antiviral therapy or fibrosis progression. The use of these genetic markers may help us to select patients who are more or less prone to respond to pegylated interferon plus ribavirin.

### 2. Aim of the work

This study assessed the predictive power of rs12979860 IL-28B variations on the response to PEG-IFN/RBV therapy in a group of Egyptian patients infected with chronic hepatitis C (CHC) genotype 4 with and without cirrhosis, in relation to other predictors of response.

# 3. Patients and methods

# 3.1. Ethical aspects

The study was approved by the local ethics committee of Alexandria University. All patients provided written informed consent to participate in this study.

### 3.2. Patients and clinical data

Chronic infection was defined as a detectable HCV RNA for at least 6 months. <sup>17</sup> The current study included 60 patients with chronic HCV infection classified according to liver biopsy and laboratory parameters into 2 groups: 30 patients with chronic HCV and compensated cirrhosis, and 30 chronic HCV infected patients without cirrhosis. All patients underwent treatment under a standard protocol with PEG-IFN and weight-based ribavirin (RBV) for 48 weeks. Patients were recruited from the Tropical department of Alexandria Main University Hospital.

Patients were subjected to thorough history taking, clinical examination and routine pre-treatment work up. The baseline HCV RNA load and subsequent viral concentrations in response to treatment were measured using a quantitative polymerase chain reaction (PCR) assay according to the available technique (Applied Biosystem, USA), with a detection

limit of 30 IU/ml. The HCV genotype was determined using the INNO-LiPA v2.0 (Innogenetics, Zwijndrecht, Belgium) HCV assav.

Histopathological examination of histological activity and degree of hepatic fibrosis, of ultrasound guided percutaneous liver biopsy, was performed according to the Metavir score. <sup>18</sup> The exclusion criteria included any cause of liver disease other than chronic HCV based on the patient history, laboratory or liver biopsy findings as: HBV/HIV coinfection, autoimmune hepatitis, hemochromatosis, Wilson's disease, Alpha 1-antitrypsin deficiency, alcoholic liver disease, drug induced liver disease, decompensated liver disease (Total serum bilirubin >1.5 g/dL; INR > 1.5; serum albumin < 3.4; platelet count < 75,000 mm) and evidence of hepatic decompensation (hepatic encephalopathy or ascites), hepatic tumors, pregnancy or breast feeding, advanced ischemic heart disease and uncontrolled diabetes (Hb A1C > 8.5%).

The outcome variable in this study was sustained virologic response, defined as non detectable HCV RNA in serum 24 weeks after the completion of HCV therapy (SVR24). Sustained virologic response is defined as non detectable HCV RNA in serum 12 weeks after the completion of HCV therapy (SVR12). Patients with detectable plasma HCV RNA at week 24, were considered to be non responders. <sup>20</sup>

# 3.3. Determination of the IL28B genotype

Genomic DNA are isolated from 200 μl of blood using the QIAamp DNA Blood Mini Kit (Qiagen) according to the manufacturer's protocol, and preserved at -70 °C for genetic determinations. SNPs near *IL28B* (marker *rs12979860*) were genotyped using TAQMAN genotyping assay (Applied Biosystems). Primer and probe sequences were: Forward Primer 5.-GCCTGTCGTGTACTGAACCA, Reverse Primer 5.-GCGCGGAGTGCAATTCAAC, Probe (C allele) 5.-VICTGGTTCGCGCCTTC, Probe (T allele) 5.-FAM-CTGGTT CACGCCTTC. Genotyping was performed in 25 μl reaction volume containing 10 ng DNA, 12.5.1 TaqMan® Universal PCR Master Mix and 1.25.1 (40×) Custom® SNP Genotyping Assays.

# 3.4. Statistical analysis

Data were analyzed using SPSS version 16.0, the 0.05 level was used as the cut off value for statistical significance. Counts and percentage were used for describing and summarizing qualitative data, the arithmetic mean  $(\overline{X})$  and the standard deviation (SD) were used as measures of central tendency and dispersion respectively for normally distributed quantitative data. The association between two qualitative variables was done using Chi square test ( $\chi^2$ ) and Fisher Exact test (FET), and Monte Carlo Exact probability (MCP) was used when  $\chi^2$  is not valid (>20% of the expected cells have count less than 5), Mann—whitney Z test was done for comparing two independent quantitative non-normally distributed variables. A stepwise logistic regression model to extract the most significant factors contributing in the response of the patient in 3 and 6 months was done.

# 4. Results

### 4.1. Features of the study population

The study group consisted of 60 chronic HCV infected patients, out of whom 30 were cirrhotic. Clinical and demographic characteristics of the patients enrolled in this study are summarized in Table 1. All patients were infected with HCV genotype 4 and were on anti-viral therapy. These characteristics were comparable in the 2 groups with no statistical significance except for the mean age which was slightly higher among the HCV group as compared to the group of HCV combined with cirrhosis, this difference although very small showed to be significant. The AST showed a significantly higher mean among the chronic HCV and cirrhosis group as compared to the other group. Schistosomal serology showed significant association, denoting a more frequent schistosoma infection among chronic HCV cirrhotic patients. All other studied lab parameters showed some difference but yet not statistically significant. The genotype frequency of rs12979860 CT was the most common (56.7%), occurring more frequently in non-cirrhotic patients (60%), than in chronic HCV with the cirrhosis group. This was statistically non significant.

# 4.2. Outcome of antiviral therapy (Table 1)

Cirrhotic patients achieved significantly lower SVR12 and SVR24 rates than non cirrhotics (12/30 = 40% and 14/30 = 46.7%) versus (23/30 = 76.7% and 24/30 = 80%) (p = 0.004 and 0.007) respectively.

## 4.3. Baseline predictors of treatment outcome

# 4.3.1. SVR12

Higher BMI was significantly associated with failure of response among the non cirrhotic group, when compared to responders (30.7  $\pm$  1.9 and 27.7  $\pm$  2.8 respectively) (p = 0.015) while higher GGTP, INR, PT, AFP, insulin resistance index (IR) and lower serum albumin levels, as well as a higher rate of positive schistosomal serology, were significantly associated with failure of response in the cirrhotic group (p < 0.05) (Table 2).

In the logistic regression model, we evaluated the odds ratio (OR) and the 95% confidence interval (CI) of SVR12, depending on the most significant predictors of response after 3 months.

The OR revealed that cirrhotic patients had a 6.39 higher risk of non response than non cirrhotic patients (p = 0.016). Presence of schistosomal antibodies had nearly 12 times greater probability of achieving non response compared with negative schistosomal serology (p = 0.003). Also patients with higher BMI had 19-fold higher chance more than non obese (Normal and Overweight) to be non responders (p = 0.001) (Table 3).

# 4.3.2. SVR24

In the non cirrhotic group, higher mean of cholesterol and LDL levels, lower mean INR, PT and negative serology for schistosomiasis demonstrated a significant association with response.

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Parameter	Group	Test of significance (p Value)		
	Chronic HCV + cirrhosis $(n = 30)$	Chronic HCV $(n = 30)$		
Age (years)	45.3 ± 7.7	49 ± 7.7	Z = 2.01 (0.045)	
Female $(N/\%)$ (17/28.3)	8 (26.7)	9 (30)	$X^2 = 0.08$	
Male $(N/\%)$ (43/71.7)	22 (73.3)	21 (70)	p = 0.774	
BMI $(kg/m^2)$	$28 \pm 3.1$	$28.4 \pm 2.9$	Z = 0.75 (0.451)	
AST(IU/L)	$53 \pm 23.7$	$47.3 \pm 38.6$	Z = 2.20 (0.028)	
Alb (g/dl)	$3.4 \pm 0.4$	$4.1 \pm 0.53$	Z = 1.19 (0.233)	
GGTP (IU/L)	$60.6 \pm 47.3$	$72.8 \pm 92.5$	Z = 0.34 (0.734)	
TLC count ( $\times 10^3/\text{mm}^3$ )	$4.50 \pm 1.43$	$4.89 \pm 1.61$	Z = 0.98 (0.328)	
Hb (g/dl)	$12.7 \pm 1.3$	$12.7 \pm 1.6$	Z = 0.26 (0.796)	
Platelets (×10 <sup>3</sup> /mm <sup>3</sup> )	$179.8 \pm 63$	$183.5 \pm 60.1$	Z = 0.18 (0.859)	
INR	$1.1 \pm 0.1$	$1.1 \pm 0.0$	Z = 1.61 (0.107)	
PT	$13.06 \pm 1.37$	$12.29 \pm 0.41$	Z = 1.82 (0.069)	
PCR-HCV (×10 <sup>6</sup> IU/ml)	$1.3 \pm 1.4$	$1.8 \pm 3.4$	Z = 0.03 (0.976)	
AFP(ng/ml)	$9.1 \pm 22.1$	$10.6 \pm 22.2$	Z = 0.26 (0.796)	
Schistosomal serology $(N/\%)$				
Negative (43/71.7)	17 (56.7)	26 (86.7)	$X^2 = 6.65$	
Positive (17/28.3)	13 (43.3)	4 (13.3)	p = 0.010	
Cholesterol (mg/dl)	$149.4 \pm 38.9$	$151.6 \pm 30.6$	$Z = 0.40 \ (0.690)$	
HDL (mg/dl)	$48.2 \pm 18.2$	$53.9 \pm 18.8$	Z = 1.37 (0.171)	
LDL (mg/dl)	$78.9 \pm 27.0$	$75.9 \pm 29.8$	Z = 0.53 (0.600)	
IR	$4.8 \pm 10.2$	$2.4 \pm 1.4$	Z = 0.27 (0.784)	
IL28B genotypes (N/%)				
CC (18/30)	11 (36.7)	7 (23.3)	MCp = 0.471	
CT (34/56.7)	16 (53.3)	18 (60)		
TT (8/13.3)	3 (10)	5 (16.7)		
SVR12 (N/%)				
Responders (35/58.3)	12 (40)	23 (76.7)	$X^2 = 8.5$	
Non responders (25/41.7)	18 (60)	7 (23.3)	p = 0.004	
SVR24 (N/%)		• 4 (0.0)	72 740	
Responders (38/63.3)	14 (46.7)	24 (80)	$X^2 = 7.18$	
Non responders (22/36.7)	16 (53.3)	6 (20)	p = 0.007	

*Note:* BMI: body mass index, (calculated as the square of height in meters divided by weight in kilograms), AST: aspartate aminotransferase, Alb: albumin, GGTP: gamma-glutamyl transpeptidase, TLC: total leukocytic count, Hb: hemoglobin, PT: prothrombin time, AFP: alpha fetoprotein, HDL: high-density lipoprotein, LDL: low-density lipoprotein, IR: insulin resistance, SVR12: sustained virologic response 12 weeks after the completion of HCV, SVR24: sustained virologic response 24 weeks after the completion of HCV therapy. Data are presented as mean ± SD unless otherwise indicated.

In the cirrhotic group of patients, GGTP, INR, PT, AFP showed a higher mean among non responders than responders and this was shown to be significant. The higher mean of HDL, negative schistosomal serology and CC IL28 B genotype showed a significant association with SVR24. Nine responders (81.8%) had the CC genotype, and 5 (58.3%) had the CT/TT genotype (p = 0.009) (Table 4).

As for the OR and the 95% CI of SVR24 after 6 months, positive schistosomal serology is the strongest predictor of non response, being more potent (p = 0.001) than the presence of cirrhosis (p = 0.020) and IL28B genotype (Table 5).

## 5. Discussion

This study was undertaken to assess the predictive power of rs12979860 IL-28B variations on the response to PEG-IFN-./ RBV therapy in a group of Egyptian patients infected with chronic HCV genotype 4 with and without compensated cirrhosis, in relation to other predictors of response.

There are many pretreatment factors such as age, sex, ethnicity, body mass index, insulin resistance, hepatic steatosis, and degree of liver fibrosis, HCV genotype, baseline viral load and viral kinetics during treatment which can influence the response to the therapy with Peg-IFN and ribavirin. <sup>12</sup>

Our findings showed that the higher mean age and the presence of negative schistosomal serology were statistically significant in the non-cirrhotic group (p=0.045, p=0.010 respectively), denoting a more frequent schistosoma infection among chronic HCV cirrhotic patients. Kamal et al. <sup>21</sup> reported that HCV/schistosomiasis coinfected patients have more rapid progression of hepatic fibrosis than those with HCV monoinfection.

Base line AST level was significantly lower in the non-cirrhotic group (p = 0.028), as reported by Al Ashgar et al.<sup>22</sup> who demonstrated that the AST reflects less severe histopathological parameters.

Recent genome-wide association studies (GWAS) have shown that human genetic variations (single-nucleotide polymorphisms, SNPs) around the gene for interleukin 28B

<b>Table 2</b> Relationship between SVR12 and demographic and biochemical variables among both studied group	Table 2	Relationship between	n SVR12 and demographic a	and biochemical variables amou	ng both studied groups
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Parameter	Chronic HCV with cirrhosis		<i>p</i> -Value	Chronic HCV without cirrhosis		<i>p</i> -Value
	Responders $(n = 12)$	Non-responders $(n = 18)$		Responders $(n = 23)$	Non-responders $(n = 7)$	
Age (years)	43.1 ± 9.8	46.8 ± 5.8	0.106	$48.4 \pm 7.5$	51.0 ± 8.5	0.446
BMI (kg/m <sup>2</sup> )	$27.2 \pm 2.3$	$28.6 \pm 3.6$	0.363	$27.7 \pm 2.8$	$30.7 \pm 1.9$	0.015
Alb (g/dl)	$4.2 \pm 0.3$	$3.8 \pm 0.4$	0.018	$4.1 \pm 0.5$	$3.9 \pm 0.7$	0.507
GGTP (IU/L)	$40.3 \pm 29.0$	$74.2 \pm 52.8$	0.026	$3.7 \pm 96.9$	$67.0 \pm 83.1$	0.624
AST (IU/L)	$51.9 \pm 25.3$	$53.7 \pm 23.3$	0.932	$51.2 \pm 43.1$	$34.4 \pm 11.6$	0.508
TLC ( $\times 10^3 / \text{mm}^3$ )	$4.3 \pm 1.1$	$4.6 \pm 1.6$	0.671	$5.1 \pm 1.5$	$4.2 \pm 2.0$	0.141
Hb (g/dl)	$12.9 \pm 1.5$	$12.5 \pm 1.2$	0.351	$12.8 \pm 1.7$	$12.3 \pm 1.0$	0.572
Platelets ( $\times 10^3 / \text{mm}^3$ )	$176.5 \pm 54.7$	$181.9 \pm 69.4$	0.816	$185.9 \pm 65.5$	$175.6 \pm 40.6$	0.961
INR	$1.0 \pm 0.1$	$1.1 \pm 0.1$	0.029	$1.0 \pm 0.0$	$1.0 \pm 0.0$	0.499
PT	$12.5 \pm 0.8$	$13.5 \pm 1.5$	0.020	$12.3 \pm 0.4$	$12.2 \pm 0.4$	0.587
PCR-HCV (×10 <sup>6</sup> IU/ml)	1722030.9 ± 1114642.4	$1089318.8 \pm 1480267.6$	0.057	$1112806.3 \pm 1067043.9$	$4184974.9 \pm 6605613.9$	0.249
AFP (ng/ml) (mean ± SD)	$4.1 \pm 5.1$	$12.4 \pm 28.1$	0.028	$8.0 \pm 12.3$	$19.2 \pm 41.4$	0.731
Cholesterol (mg/dl)	$152.3 \pm 32.5$	$12.4 \pm 20.1$ $147.6 \pm 43.4$	0.687	$154.3 \pm 29.6$	$19.2 \pm 41.4$ $142.4 \pm 34.4$	0.364
HDL (mg/dl)	$57.4 \pm 23.1$	$42.0 \pm 11.0$	0.087	$54.1 \pm 20.8$	$53.3 \pm 11.6$	0.864
LDL (mg/dl)	$73.8 \pm 16.8$	$82.3 \pm 32.1$	0.485	$79.2 \pm 30.2$	$64.9 \pm 28.0$	0.229
IR	$1.6 \pm 0.7$	$6.9 \pm 12.9$	0.034	$2.4 \pm 1.4$	$2.3 \pm 1.4$	0.922
Male $(N/\%)$	8 (36.4)	14 (63.6)	FETp = 0.678	15 (71.4)	6 (28.6)	FETp = 0.393
Female $(N/\%)$	4 (50.0)	4 (50.0)	1L1p 0.076	8 (88.9)	1 (11.1)	1E1p 0.55.
Schistosomal serology $(N/\%)$						
Negative	10 (58.8)	7 (41.2)	FET p = 0016.	20 (76.9)	6 (23.1)	FETp = 0.999
Positive	2 (15.4)	11 (84.6)	·	3 (75.0)	1 (25.0)	•
IL28B (N/%)						
CC	6 (54.5)	5 (45.5)	MCP = 0.267	5 (71.4)	2 (28.6)	MCP = 0.591
CT	6 (37.5)	10 (62.5)		13 (72.2)	5 (27.8)	
TT	0 (0.0)	3 (100)		5 (100)	0 (0.0)	

**Table 3** Odds ratio of SVR12 depending on the patient group, BMI and schistosomal serology.

Factors affecting SVR12	p value	OR	95% CI
Cirrhotic group BMI Positive schistosomal serology	0.016 0.001 0.003	6.39 19.07 12.88	1.41–28.87 3.15–115.66 2.34–70.83
Model $X^2 = 26.57 \text{ p} = 0.000$	0.003	12.00	2.51 70.05

OR: odds ratio, CI: confidence interval.

(IL-28B) may explain differences in the results of the treatment of adults chronically infected with HCV and that they can be useful as therapy response markers.<sup>17</sup>

The IL28B polymorphism rs12979860 has a marked differential distribution between racial groups, being least frequent in African Americans, most frequent in Asians, and with an intermediate frequency in Hispanics and Caucasians.<sup>23</sup>

This global difference of allele frequency might explain the ethnic variations seen in the treatment response among these populations.<sup>24</sup>

The frequency of IL28 genotype in our genotype 4 Egyptian patients showed that almost half of them were of the CT genotype (56.7%) followed by CC (30%) while TT had the least expression (13%). The CC genotype occurred more frequently in cirrhotic patients compared to non cirrhotic patients (11/30)

vs. 7/30), unlike CT and TT genotypes that were more frequent in chronic HCV patients compared to HCV with cirrhosis (18/30 and 5/30 vs. 16/30 and 3/30 respectively), but this was statistically non significant.

These results were matching with the study of Khairy et al.<sup>23</sup> who reported that the frequency of IL28 genotype in their 263 chronic HCV patients genotype 4 Egyptian patients was 56% for the CT genotype, followed by 25% for CC while TT had the least expression (19%). De Nicola group (2012)<sup>25</sup> which included 128 patients with genotype 4, 68% Egyptians, showed 63% CT, 14% TT, and 23% CC expression. Also, Asselah and colleagues<sup>26</sup> studied 164 patients with genotype 4 (43% Egyptians), and found the difference in distribution of IL 28 B genotype between Egyptians and Subsaharan Africans; in the Egyptian ethnicity the frequency was 55% CC, 11% TT and 34% CT, while in the in sub-Saharan group the TT genotype was the most predominant form (48%). El-Awady and colleagues during 2012 also in a study on genotype 4, found that the frequencies of genotypes were 48% CC, 14% TT, and 38% CT for their studied patients.<sup>27</sup>

This was inconsistent with previous results describing a higher incidence of TT or CT allele in cases of cirrhosis and faster fibrosis progression in HCV-infected liver transplant recipients and liver from donors with the TT genotype, <sup>28,29</sup> and that TT genotype occurred more frequently in patients with end stage liver disease, <sup>23</sup> as well as that of Ciesla et al. <sup>15</sup>, who studied 64 Caucasian chronic HCV patients on IFN and

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Table 4	Relationship	between SVR24 at	nd demographic and	biochemical variab	oles among both studied groups.

Parameter	Chronic HCV with cirrhosis		<i>p</i> -Value	Chronic HCV without cirrhosis		<i>p</i> -Value
	Responders $(n = 14)$	Non-responders $(n = 16)$		Responders $(n = 24)$	Non-responders $(n = 6)$	
Age (years)	44.1 ± 9.0	$46.4 \pm 6.5$	0.337	$49.3 \pm 7.6$	$48.0 \pm 8.7$	0.815
BMI $(kg/m^2)$	$28.2 \pm 2.7$	$27.9 \pm 3.6$	0.506	$28.8 \pm 2.7$	$26.7 \pm 3.3$	0.186
Alb (g/dl)	$4.1\pm0.4$	$3.8 \pm 0.4$	0.132	$4.1 \pm 0.5$	$4.0 \pm 0.6$	0.406
GGTP (IU/L)	$32.6 \pm 17.6$	$85.2 \pm 51.8$	0.001	$72.3 \pm 102.2$	$71.7 \pm 39.4$	0.169
AST (IU/L)	$53.1 \pm 28.2$	$52.8 \pm 19.9$	0.901	$50.0 \pm 42.7$	$36.5 \pm 8.2$	0.917
TLC ( $\times 10^3 / \text{mm}^3$ )	$4.6 \pm 1.8$	$4.4 \pm 1.1$	0.917	$4.7 \pm 1.7$	$5.6 \pm 1.4$	0.213
Hb (g/dl)	$12.4 \pm 1.2$	$12.9 \pm .4$	0.454	$12.8 \pm 1.7$	$12.4 \pm 1.3$	0.604
Platelets ( $\times 10^3 / \text{mm}^3$ )	$175.4 \pm 52.2$	$183.6 \pm 72.6$	0.835	$182.3 \pm 58.8$	$188.3 \pm 71.0$	0.795
PT	$12.4 \pm 0.8$	$13.7 \pm 1.5$	0.001	$12.2 \pm 0.4$	$12.6 \pm 0.5$	0.036
INR	$1.0 \pm 0.1$	$1.1 \pm 0.1$	0.000	$1.0 \pm 0.0$	$1.1 \pm 0.0$	0.058
PCR-HCV (×10 <sup>6</sup> IU/ml)	$1801612.1 \pm$	$940596.3 \pm$	0.212	$1977492.3 \pm$	$1238258.8 \pm$	0.717
i i i	1743753.6	763112.5		3780153.0	1115577.7	
AFP (ng/ml)	$3.3 \pm 2.7$	$14.2 \pm 29.7$	0.010	$5.6 \pm 7.3$	$30.9 \pm 44.7$	0.062
Cholesterol (mg/dl)	$158.1 \pm 38.0$	$141.8 \pm 39.2$	0.339	$158.1 \pm 29.7$	$125.5 \pm 18.7$	0.022
HDL (mg/dl)	$57.3 \pm 21.8$	$40.2 \pm 9.2$	0.019	$52.8 \pm 20.6$	$58.3 \pm 9.2$	0.222
LDL (mg/dl)	$75.9 \pm 24.8$	$1.5 \pm 29.4$	0.708	$81.2 \pm 29.9$	$54.5 \pm 19.3$	0.038
IR	$2.0 \pm 1.4$	$7.2 \pm 13.7$	0.096	$2.3 \pm 1.4$	$2.5 \pm 1.4$	0.678
Male $(N/\%)$	9 (40.9)	13 (59.1)	FETp = 0.417	16 (76.2)	5 (23.8)	FETp = 0.400
Female $(N/\%)$	5 (62.5)	3 (37.5)	Ť	8 (88.9)	1 (11.1)	Ť
Schistosomal serology (N/	%)					
Negative	13 (76.5)	4 (23.5)	FETp = 0.000	23 (88.5)	3 (11.5)	FETp = 0.018
Positive	1 (7.7)	12 (92.3)	Ť	1 (25.0)	3 (75.0)	Ť
IL28B (N/%)						
CC	9 (81.8)	2 (18.2)	MCP = 0.009	7 (100.0)	0 (0.0)	MCP = 0.372
CT	4 (25.0)	12 (75.0)		13 (72.2)	5 (27.8)	
TT	1 (33.3)	2 (66.7)		4 (80.0)	1 (20.0)	

Table 5 Odds ratio of SVR24 depending on the patient group, type of polymorphism of IL28B and schistosomal serology.

Factors affecting SVR24	p value	OR	95% CI		
Cirrhotic group	0.020	11.40	1.46-88.84		
CT	0.043				
CC	0.384	2.99	0.25-35.24		
TT	0.064	0.03	0.00-1.22		
Positive schistosomal serology	0.001	132.25	7.34-2383.52		
Model $X^2 = 46.43$	p = 0.00	0			
OR: odds ratio, CI: confidence interval.					

Ribavirin therapy, where patients with the TT genotype had a more active state of necroinflammation in the histological analysis, and borderline significance of greater severity of fibrosis. Taking into consideration that the distribution of IL28B genotypes among both studied groups in the present study was not statistically significant, it was not possible to confirm the impact of the genotype on the progress of CHC.

Endpoint of therapy is the SVR, defined by undetectable HCV RNA 24 weeks after the end of therapy, as assessed by a sensitive molecular method with a lower limit of detection < 15 IU/ml (SVR24). Long-term follow-up studies have shown that an SVR corresponds to a definitive cure of HCV infection in more than 99% of cases. The validity of using undetectable HCV RNA at 12 weeks after the end of therapy (SVR12) has been accepted by regulators in the US and Europe, given that the concordance with SVR24 is 99%.<sup>30</sup>

Various studies from European and Middle Eastern countries showed that the SVR in genotype 4 for combination therapy, pegylated interferon and ribavirin, ranges between 43% and 70%. 31,32

In the current study, SVR12 and SVR24 showed significant association with non cirrhotic patients (p = 0.004, p = 0.007respectively) denoting higher response among the same group.

This was close to Aghemo et al. results, 33 where 53% of 409 HCV cirrhotic patients achieving an SVR compared to 75% of non cirrhotics, and Thompson et al. study,<sup>34</sup> showing lower SVR rates in IL28B CC patients with bridging fibrosis and cirrhosis.

Little is known about predictors of response within populations infected with genotype 4. In previous studies on genotype 4; age, pretreatment viral load, and stage of fibrosis were considered as good predictive factors. 35,36

The gene expression and the role of IL28B gene SNP rs12979860 in response to treatment in genotype 4 were recently studied by limited research with CC genotype of higher response rate.<sup>25,26</sup>

Relationship between baseline parameters and virological response in the current study showed that, higher base line GGTP, INR, PT, AFP levels, higher IR index, lower albumin level and positive schistosomal serology antibodies are significantly associated with unfavorable outcomes after 3 months

(SVR12) in cirrhotic group, while lower BMI value was the only significant predictor of SVR12 in non cirrhotic group.

The role of insulin resistance in response to HCV therapy is controversial, <sup>20</sup> the correlation between insulin resistance and higher BMI and response to antiviral therapy observed in the present study is consistent with Ciesla et al. results. <sup>15</sup>

In this study, although CC patients are more prone than patients with the T allele to achieve an SVR12 in cirrhotic patients, this is not the case in HCV non cirrhotic patients, where CT/TT alleles are more prone to achieve SVR12, but this was statistically non significant.

Concerning rates of SVR24 in our study, cirrhotic patients (genotype CC) did achieve it more frequently as compared to genotype CT/TT, which was statistically significant (p = 0.009), while in the non cirrhotic group, this difference was statistically non significant.

This was in harmony with Khairy et al.<sup>23</sup> study, where the CC genotype was significantly correlated with SVR in comparison to CT and TT. The response rates were 50%, 47.4% and 25% for genotype CC, CT, and TT respectively. Absence of C allele (TT genotype) was associated with 75% failure of response; either early failure, e.g. non response (54.5%), or late failure, e.g. relapsers (20.5%). This is in agreement with previous studies reported on genotype 1<sup>15,37,38</sup> and studies conducted on genotype 4.<sup>26,25</sup>

On the other hand, in the study of Ciesla et al., <sup>15</sup> 60% of patients with SVR had TT (8%) or CT (52%) genotypes, which are not known factors associated with higher IFN responsiveness. These values are similar to those described in previous reports, where analysis of the differences in the course of therapy in patients with the CT allele and a SVR revealed higher baseline platelet and neutrophil levels. A low baseline platelet count was significantly associated with the need for IFN dose reduction in the group without a SVR. <sup>39</sup>

In the present study, negative serology for schistosoma antibodies was a common predictor of SVR24 in both groups of patients. Besides, we found 2 independent prognostic factors for SVR24 in cirrhotic patients: Lower INR and PT level and higher cholesterol and LDL levels. As regards the non cirrhotic group, higher HDL level was a predictor of SVR24, while higher GGT, INR, PT and AFP levels were associated with failure of response. Base line viral load and base line AST levels were not predictors of SVR24 in both groups.

In the study of Abdel-Rahman et al. <sup>40</sup> the EVR, virological response at week 24, and SVR were significantly higher in patients with negative schistosomal serology. This finding may be attributed to the fact that coinfected patients with a down-regulated immune response to HCV leading to reduced IFN., interleukin (IL)-4 and IL-10 secreted by HCV-specific T cells.

In contrast to our study, the analysis presented by McCarthy et al.<sup>38</sup> showed that lower viral load before treatment predicted higher SVR. This phenomenon was not confirmed in our study, possibly due to the small sample size. This result has been affirmed by Correia et al.<sup>41</sup> emphasizing that the baseline HCV RNA is trivial for SVR.

In the study of Khairy et al.<sup>23</sup>, lower base line AST but not ALT was an independent predictor of SVR in patients with chronic HCV genotype 4. This was in accordance with Al Ashgar et al. study<sup>22</sup> which demonstrated that the AST reflects less severe histopathological parameters in sustained responders.

Concerning the plasma levels of LDL, in vitro studies have shown that LDL may competitively inhibit the binding of HCV to the LDL receptor, which functions as one of the cellular receptors for HCV. This competitive blockade would hamper the infection of hepatocytes with HCV. Accordingly, higher levels of plasma LDL (as in the current results) have been shown to be an independent predictor of SVR, in studies specifically designed to appraise this issue.<sup>20</sup>

Higher serum AFP level was a strong negative predictor of SVR24 in the studied patients. Previous studies including HCV genotype 4<sup>42,43</sup> and genotype 1<sup>44,45</sup> highlighted the same findings. Abdoul et al.<sup>46</sup> examined the association between serum alpha-fetoprotein level and sustained SVR in 93 chronic hepatitis C patients and found that the SVR rate was much higher among patients with serum AFP levels below rather than above a median value of 5.7 ng/ml, denoting that serum AFP should be added to the list of factors predictive of treatment response in chronic hepatitis C.

The results of multivariate analysis in this study show that 3 factors only were considered to be significant negative predictors of response after 3 months (SVR12), which are presence of cirrhosis, high BMI and positive schistosomal serology. As regards SVR24 after 6 months, positive schistosomal serology is the strongest predictor of non response, being more potent than the presence of cirrhosis and IL28B genotype.

We can conclude that concomitant absence of cirrhosis and negative serology for bilharzial infections do seem to improve responses achieved with pegylated interferon plus ribavirin combination therapy for HCV genotype 4. In Egypt, where chronic HCV genotype 4 and schistosoma coinfection predominate, both schistosoma infection and cirrhosis are more potent than IL28B polymorphisms as strong baseline negative predictors of hepatitis C treatment response.

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### Conflict of interest

None declared.

# References

- European Association for the Study of the Liver. EASL Clinical Practice Guidelines: management of hepatitis C virus infection. J Hepatol 2014;60:392–420.
- El-Zanaty F, Egypt Way A. Demographic and health survey, 2009. Cairo: Ministry of Health, El-Zanaty and Associates, and Macro International; 2008.
- Ray SC, Arthur RR, Carella A, Bukh J, Thomas DL. Genetic epidemiology of hepatitis C virus throughout Egypt. J Infect Dis 2000:182(3):698–707.
- Hoofnagle JH. Course and outcome of hepatitis C. Hepatology 2002;36(5 Suppl. 1):S21–9.
- McHutchison JG, Lawitz EJ, Shiffman ML, Muir AJ, Galler GW, McCone J. Peginterferon alfa-2b or alfa-2a with ribavirin for treatment of hepatitis C infection. N Engl J Med 2009;361:580–93.
- Ge D, Fellay J, Thompson AJ, Simon JS, Shianna KV, Urban TJ, et al. Genetic variation in IL28B predicts hepatitis C treatmentinduced viral clearance. *Nature* 2009;461:399–401 PMID: 19684573 DOI: 10.1038/nature08309.

- Lyra AC, Ramrakhiani S, Bacon BR, Di Bisceglie AM. Infection with hepatitis C virus genotype 4 in the United States. J Clin Gastroenterol 2004;38(1):68-71.
- Van Asten L, Verhaest I, Lamzira S, Hernandez-Aguado I, Zangerle R, Boufassa F, et al. Spread of hepatitis C virus among European injection drug users infected with HIV: a phylogenetic analysis. *J Infect Dis* 2004;189(2):292–302.
- 9. Payan C, Roudot-Thoraval F, Marcellin P, Bled N, Duverlie G, Fouchard-Hubert I, et al. Changing of hepatitis C virus genotype patterns in France at the beginning of the third millennium: The GEMHEP GenoCII Study. *J Viral Hepatol* 2005;12(4):405–13.
- Degasperi E, Aghemo A. Clinical drivers in naïve patient eligibility for treatment of chronic hepatitis C. J Viral Hepat 2012;19(1):3–6.
- Bochud PY, Bibert S, Negro F, Haagmans B, Soulier A, Ferrari C, et al. IL28B polymorphisms predict reduction of HCV RNA from the first day of therapy in chronic hepatitis C. *J Hepatol* 2011;55:980–8 PMID: 21354446 DOI: 10.1016/j.jhep.2011.01.050.
- Jablonowska E, Piekarska A, Koslinska-Berkan E, Omulecka A, Szymanska B, Wojcik K. Sustained virologic response and IL28B single-nucleotide polymorphisms in patients with chronic hepatitis C treated with pegylated interferon alfa and ribavirin. *Acta Biochim Polon* 2012:59(3):333-7.
- Ordi-Ros J, Villarreal J, Monegal F, Sauleda S, Esteban I, Ge D, et al. Genetic variation in IL28B predicts hepatitis C treatmentinduced viral clearance. *Nature* 2009;461:399–401.
- McHutchison JG, Goldstein DB, Carrington M. Genetic variation in IL28B and spontaneous clearance of hepatitis C virus. *Nature* 2009;461:798–801 PMID: 19759533 DOI: 10.1038/nature08463.
- Ciesla A, Bociaga-Jasik M, Sobczyk-Krupiarz I, Glowacki MK, Owczarek D, Cibor D, et al. IL28B polymorphism as a predictor of antiviral response in chronic hepatitis C. World J Gastroenterol 2012;18(35):4892–7.
- Khattab MA, Ferenci P, Hadziyannis SJ, Colombo M, Manns PL, Almasio PL, et al. Management of hepatitis C virus genotype 4: recommendations of an international expert panel. *J Hepatol* 2011;54(6):1250–62.
- Domagalski K, Pawlowska M, Tretyn A, Halota W, Pilarczyk M, Smukalska E, et al. Impact of IL-28B polymorphisms on pegylated interferon plus ribavirin treatment response in children and adolescents infected with HCV genotypes 1 and 4. Eur J Clin Microbiol Infect Dis 2013;32:745–54. http://dx.doi.org/10.1007/ s10096-012-1799-z.
- 18. Bedossa P, Poynard T. An algorithm for the grading of activity in chronic hepatitis C. The METAVIR Cooperative Study Group. *Hepatology* 1996;**24**(2):289–93.
- Ghany MG, Strader DB, Thomas DL, Seeff LB. Diagnosis, management and treatment of hepatitis C: an update. *Hepatology* 2009;49(4):1335–74. <a href="http://dx.doi.org/10.1002/hep.22759">http://dx.doi.org/10.1002/hep.22759</a>.
- Pineda JA, Caruz A, Rivero A, Neukam K, Salas I, Camacho A, et al. Prediction of response to pegylated interferon plus ribavirin by IL28B gene variation in patients coinfected with HIV and hepatitis C virus. *Clin Infect Dis* 2010;51(7):788–95.
- Kamal SM, Graham CS, He Q, Bianchi L, Tawil AA, Rasenack JW, et al. Kinetics of intrahepatic hepatitis C virus (HCV)-specific CD4+ T cell responses in HCV and Schistosoma mansoni coinfection: relation to progression of liver fibrosis. *J Infect Dis* 2004;189:1140–50 PMID: 15031780 DOI: 10.1086/382278.
- 22. Al Ashgar H, Helmy A, Khan MQ, Al Kahtani K, Al Quaiz M, Rezeig M, et al. Predictors of sustained virological response to a 48-week course of pegylated interferon alfa-2a and ribavirin in patients infected with hepatitis C virus genotype 4. Ann Saudi Med 2009;29(1):4–14.
- 23. Khairy M, Fouad R, Mabrouk M, El-Akel W, Awad A, Salama R, et al. The impact of Interleukin 28b gene polymorphism on the virological response to combined pegylated interferon and ribavirin therapy in chronic HCV genotype 4 infected Egyptian patients using data mining analysis. *Hepat Mon* 2013;13(7):e10509. <a href="http://dx.doi.org/10.5812/hepatmon.10509">http://dx.doi.org/10.5812/hepatmon.10509</a>.

- 24. Jun-qiang X, Xiao-yan G, Xiao-hong Z, Bing-liang L, Dong-ying G, Zhi-liang G, et al. Relationship between the genetic variation in interleukin 28B and response to antiviral therapy in patients with chronic hepatitis C. *Chin Med J* 2012;**125**(13):2334–8.
- 25. De Nicola S, Aghemo A, Rumi MG, Galmozzi E, Valenti L, Soffredini R, et al. Interleukin 28B polymorphism predicts pegylated interferon plus ribavirin treatment outcome in chronic hepatitis C genotype 4. *Hepatology* 2012;55(2):336–42.
- Asselah T, De Muynck S, Broet P, Masliah-Planchon J, Blanluet I, Bieche I, et al. IL28B polymorphism is associated with treatment response in patients with genotype 4 chronic hepatitis C. J Hepatol 2012;56(3):527–32.
- 27. El-Awady MK, Mostafa L, Tabll AA, Abdelhafez TH, Bader El Din NG, Zayed N, et al. Association of IL28B SNP with progression of Egyptian HCV genotype 4 patients to end stage liver disease. *Hepat Mon* 2012;12(4):271–7.
- 28. Charlton MR, Thompson A, Veldt BJ, Watt K, Tillmann H, Poterucha JJ, et al. Interleukin-28B polymorphisms are associated with histological recurrence and treatment response following liver transplantation in patients with hepatitis C virus infection. *Hepatology* 2011;53:317–24.
- 29. Fabris C, Falleti E, Cussigh A, Bitetto D, Fontanini E, Bignulin S, et al. IL-28B rs12979860 C/T allele distribution in patients with liver cirrhosis: role in the course of chronic viral hepatitis and the development of HCC. *J Hepatol* 2011;54:716–22.
- 30. Martinot-Peignoux M, Stern C, Maylin S, Ripault MP, Boyer N, Leclere L, et al. Twelve weeks post treatment follow-up is as relevant as 24 weeks to determine the sustained virologic response in patients with hepatitis C virus receiving pegylated interferon and ribavirin. *J Hepatol* 2014;60:392–420.
- 31. Hasan F, Asker H, Al-Khaldi J, Siddique I, Al-Ajmi M, Owaid S, et al. Peginterferon alfa-2b plus ribavirin for the treatment of chronic hepatitis C genotype 4. *Am J Gastroenterol* 2004;99(9):1733–7.
- Moucari R, Ripault MP, Martinot-Peignoux M, Voitot H, Cardoso AC, Stern C, et al. Insulin resistance and geographical origin: major predictors of liver fibrosis and response to peginterferon and ribavirin in HCV-4. Gut 2009;58(12):1662–9.
- 33. Aghemo A, Degasperi E, Rumi MG, Galmozzi E, Valenti L, De Francesco R, et al. Cirrhosis and rapid virological response to peginterferon plus ribavirin determine treatment outcome in HCV-1 IL28B rs12979860 CC patients. BioMed Res Int 2013; Article ID 580796, 6 pages.
- 34. Thompson AJ, Muir AJ, Sulkowski MS, et al. Interleukin28B polymorphism improves viral kinetics and is the strongest pretreatment predictor of sustained virologic response in genotype 1 hepatitis C virus. *Gastroenterology* 2010;139(1):120–9.
- 35. Kamal SM, El Kamary SS, Shardell MD, Hashem M, Ahmed IN, Muhammadi M, et al. Pegylated interferon alpha-2b plus ribavirin in patients with genotype 4 chronic hepatitis C: the role of rapid and early virologic response. *Hepatology* 2007;46(6):1732–40.
- Gad RR, Males S, El Makhzangy H, Shouman S, Hasan A, Attala M, et al. Predictors of a sustained virological response in patients with genotype 4 chronic hepatitis C. Liver Int 2008;28(8):1112–9.
- 37. Rauch A, Kutalik Z, Descombes P, Cai T, Di Iulio J, Mueller T, et al. Genetic variation in IL28B is associated with chronic hepatitis C and treatment failure: a genome-wide association study. *Gastroenterology* 2010;138(4):1338–45.
- **38.** McCarthy JJ, Li JH, Thompson A, Suchindran S, Lao XQ, Patel K, et al. Replicated association between an IL28B gene variant and a sustained response to pegylated interferon and ribavirin. *Gastroenterology* 2010;**138**(7):2307–14.
- **39.** Romero-Gomez M, Eslam M, Ruiz A, Maraver M. Genes and hepatitis C: susceptibility, fibrosis progression and response to treatment. *Liver Int* 2011;**31**:443–60.
- **40.** Abdel-Rahman M, El-Sayed M, El Raziky M, Elsharkawy A, El-Akel W, Ghoneim H, et al. Coinfection with hepatitis C virus and

- schistosomiasis: fibrosis and treatment response. World J Gastro-enterol 2013;19(17):2691-6.
- 41. Correia MC, Domingues AL, Lacerda HR, Santos EM, Machado V, Hora V, et al. Platelet function and the von Willebrand factor antigen in the hepatosplenic form of schistosomiasis mansoni. *Trans R Soc Trop Med Hyg* 2009;103:1053–8.
- **42.** Gad HH, Dellgren C, Hamming OJ, Vends S, Paludan SR, Hartmann R. Interferon-lambda is functionally an interferon but structurally related to the interleukin-10 family. *J Biol Chem* 2009;**284**(31):20869–75.
- Males S, Gad RR, Esmat G, Abobakr H, Anwar M, Rekacewicz C, et al. Serum alpha-fetoprotein level predicts treatment outcome in chronic hepatitis C. *Antivir Ther* 2007;12(5):797–803.
- 44. Akuta N, Suzuki F, Kawamura Y, Yatsuji H, Sezaki H, Suzuki Y, et al. Predictors of viral kinetics to peginterferon plus ribavirin combination therapy in Japanese patients infected with hepatitis C virus genotype 1b. *J Med Virol* 2007;79(11):1685–6.
- 45. Chen TM, Huang PT, Tsai MH, Lin LF, Liu CC, Ho KS, et al. Predictors of alpha-fetoprotein elevation in patients with chronic hepatitis C, but not hepatocellular carcinoma, and its normalization after pegylated interferon alfa 2a-ribavirin combination therapy. J Gastroenterol Hepatol 2007;22(5):669–75.
- Abdoul H, Mallet V, Pol S, Fontanet A. Serum alpha-fetoprotein predicts treatment outcome in chronic hepatitis C patients regardless of HCV genotype. *PLoS One* 2008;3(6):e2391. <a href="http://dx.doi.org/10.1371/journal.pone.0002391">http://dx.doi.org/10.1371/journal.pone.0002391</a>.