Influence of Host and Viral Factors on Patients with Chronic Hepatitis C Virus Genotype 6 Treated with Pegylated Interferon and Ribavirin: A Systematic Review and Meta-Analysis

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Objectives: We conducted a systematic review and meta-analysis of the influence of host and viral factors on the sustained virologic response (SVR) in hepatitis C virus genotype 6 (HCV-6) patients treated with pegylated interferon (PEG-IFN) and ribavirin (RBV).

Methods: Data were retrieved from Medline, Embase, PubMed and the Cochrane Library for ‘genotype 6’ studies published up to December 2014 and for abstracts from international scientific meetings. Inclusion criteria were efficacy of PEG-IFN+RBV based on SVR, 24- or 48-week therapy and treatment-naïve patients. Patients with hepatitis B, D and E and HIV coinfection or another concurrent liver disease were excluded. Pooled standard difference, odds ratio and confidence intervals (CIs) were calculated using a random-effect model with STATA 11.

Results: Fourteen studies were included in the meta-analysis. The pooled SVR rate was 80% (95% CI: 0.78–0.83, p < 0.0001; I² = 71.2%). SVR of the PEG-IFN+RBV-treated HCV-6 patients was markedly higher than that of HCV-1 patients (80.1 vs. 55.3%). The SVR rate was significantly higher for the 48- week treatment, but not different among HCV-infected patients with rs12979860 and ss469415590 polymorphisms of the ILFN4 gene (80.6% CC vs. 66.7% non-CC, p = 0.593; 81.1% TT/TT vs. 60% non-TT/TT, p = 0.288). Gender and type of PEG-IFN did not affect SVR rates. Conclusions: Treatment outcomes for HCV-6 patients are superior to those for HCV-1 patients and comparable to those of HCV-2 and HCV-3 patients, especially at 48 weeks. The level of fibrosis affects treatment outcome, but SVR rates are not significantly different between genders. IL28B and IFNL4 polymorphisms are not significantly associated with HCV-6 treatment outcome.

Introduction

Hepatitis C virus (HCV) is an important public health problem with more than 160 million cases of chronic infection worldwide [1]. HCV infection commonly progresses to liver fibrosis, cirrhosis and hepatocellular car-
cinoma, and related mortality is predicted to increase over the next two decades [2]. Genotype 1 (HCV-1) is found globally, especially in developed regions, such as America and Europe. HCV-3 and -6 are predominantly detected in Asian countries, which have a high prevalence of HCV [3].

Various factors can be effectively used to identify treatment responses in HCV patients. The standard treatment for HCV is a combination of pegylated interferon (PEG-IFN) and ribavirin (RBV) with a 24- to 48-week regimen, depending on the viral genotype of infected individuals [4–6]. Previous studies have reported that treatment outcomes of HCV-6 are closer to those of HCV-2 and HCV-3 [7–9], but patients infected with HCV-6 respond better to therapy than those with HCV-1 [3], defined as sustained virologic response (SVR). Several host factors, including age, sex, race and the level of fibrosis, additionally influence treatment outcome. Single nucleotide polymorphisms on chromosome 19 within or near the *interleukin-28B* gene (*IL28B* encoding interferon λ-3) represent one of the strongest baseline predictors of SVR in HCV treatment. Recent studies have additionally shown that the *IFN-λ-4* (*IFNL4*) gene polymorphisms are in high linkage disequilibrium with those near *IL28B* and more strongly associated with spontaneous or treatment-induced HCV clearance than *IL28B* genotypes.

While advances in HCV therapy continue to evolve rapidly with the development of potent direct-acting antiviral agents [10], limited data availability has prevented general recommendations for HCV-6 therapy. PEG-IFN+RBV combination therapy continues to be a predominant option in resource-limited settings due to the high costs associated with new agents. To date, only one study reported on the influence of host and viral factors in HCV-4 [11]. Moreover, this treatment option remains elusive for patients in developing countries, such as Asia, where HCV-6 infection is high and funding for medications is inadequate. The current study aims to provide a systematic review and meta-analysis of the influence of host and viral factors on virologic response in HCV-6 patients treated with PEG-IFN+RBV.

**Materials and Methods**

**Data Sources and Search Strategy**

This systematic review was conducted according to PRISMA guidelines [12]. PubMed, Embase, Medline and the Cochrane Library were used for comprehensive literature searches. We searched the databases for the following term: genotype 6. The search was conducted for studies published from inception to December 2014 and also included a manual search of abstracts using the term genotype 6 from annual international scientific meetings in the areas of liver disease. To ensure maximum sensitivity of our search strategy, we kept the search string as simple as possible, using only key words without filters. Search and study selection were performed without language limitations. We reviewed the bibliographies of relevant published articles for inclusion in our study. References concerning previous meta-analyses were additionally explored for eligibility.

**Inclusion and Exclusion Criteria**

Data were included based on the following criteria: (i) treatment-naïve patients, (ii) assessment of the efficacy of PEG-IFN+RBV therapy based on SVR, which was defined as undetectable HCV-RNA at least 24 weeks after the end of treatment, (iii) course of treatment (24 or/and 48 weeks), and (iv) reports in English. We excluded studies comprising patients with hepatitis B, D and E or HIV coinfection and those with other concurrent liver diseases. Studies were additionally excluded if insufficient data were available for pooling. Two of the authors independently reviewed the articles, and discrepancies were reviewed by another author and resolved by consensus.

**Data Extraction**

Information extracted from published material included study data (author, publication year and status, country of origin, continent, study design, total sample size, HCV genotype and type of PEG-IFN used), demographic data on study participants (age, sex and ethnicity) and treatment outcomes (number of patients who achieved or failed to achieve SVR). Data on treatment characteristics were additionally collected, including *IL28B* and *IFNL4* polymorphisms, duration of treatment (48 vs. 24 weeks), baseline HCV-RNA levels and stage of liver fibrosis/cirrhosis.

**Statistical Analysis**

Pooled standard differences in means (overall SVR rates) and 95% confidence intervals (CIs) were calculated for each group using a random-effect model and inverse variance method [13]. Heterogeneity was tested with *I*² test-based Cochran’s Q statistics with *p* values up to 0.05, and the degree of heterogeneity quantified using the *I*² statistic representing the percentage of total variability across studies due to heterogeneity. *I*² values of 25, 50 and 75% corresponded to low, moderate and high degrees of heterogeneity, respectively. Univariate and multivariate random-effect meta-regression on study level characteristics was performed to explain any observed heterogeneity in primary outcomes and identify associated patient level factors [13, 14]. We quantified publication bias using Egger’s regression model. All analyses were performed with STATA 11 (Stata Corporation, College Station, Tex., USA) [15, 16].

**Results**

**Study Selection**

In total, 207 articles relevant to the HCV-6 were initially identified from our searches. The modified process employed for study search and selection is summarized in figure 1. After the titles were screened and abstracts...
Influence of Host and Viral Factors on HCV-6

Clinical Study and Patient Characteristics

Characteristics of the included studies are presented in table 1. The pooled number of HCV-6 patients was 1,176 and study sizes ranged from 12 to 70, except for one study, which included 242 patients. Six studies evaluated patients with HCV-1 and -6, and 3 evaluated patients with HCV-1, -2, -3 and -6. Seven of the studies evaluated HCV-6 only. The majority of studies (n = 11) were nonrandomized clinical trials (non-RCT; 2 RCT) with predominantly male patients. Mean age ranged from 19 to 50 years. In all studies, patients underwent combination treatment with PEG-IFN+RBV for 48 weeks; only 5 studies evaluated the effects of therapy for 24 weeks. The funnel plot was symmetrical and Egger’s regression for publication bias was not statistically significant (p = 0.912; fig. 2).

Overall SVR Rates

Thirteen trials reported SVR data from a total of 891 HCV-6 patients. SVR to PEG-IFN plus weight-based RBV therapy in HCV patients ranged from 39 to 92.8%. In influence analysis, the pooled SVR rate for all studies was 80% (95% CI: 0.78–0.83; p < 0.0001; I² = 71.2%; fig. 3).

Upon comparing RCT versus non-RCT, we observed a pooled SVR rate of 70% (95% CI: 0.64–0.77) in 2 RCTs [20, 27], compared with 83% (95% CI: 0.79–0.86) in 11 non-RCTs. This finding was statistically significant (p = 0.032). Heterogeneity was observed in both subgroups (I² = 72.1% for non-RCT; I² = 67.1% for RCT).

SVR Rates in Relation to Host and Viral Factors

Male versus Female Patients

In 11 of the 14 studies with 891 HCV-6 patients, gender was reported for 537 patients, 320 (60%) males and 217 (40%) females, which included information on the influence of sex on SVR. The SVR rate was 72% in males (95% CI: 0.62–0.87) and 73% in females (95% CI: 0.61–0.89). Notably, no significant differences in SVR were evident between males and females with an odds ratio (OR) of 0.81 (95% CI: 0.71–1.12, p = 0.71). Weak heterogeneity was observed in our model (I² = 2%, p = 0.42).

IL28B and IFNL4 Polymorphisms

Two studies involving 102 patients assessed SVR in relation to IL28B polymorphisms while 1 study focused on IFNL4 polymorphisms [23, 29]. Among the 2 studies on IL28B polymorphisms, 1 study investigating 60 HCV-6 patients [52 (90%) TT and 8 (10%) GT/GG] as-

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Fig. 1. Flow diagram of study inclusion criteria for meta-analysis.
sessed SVR in rs8099917 TT compared with GT/GG. One study evaluated SVR in rs12979860 CC versus CT/TT and ss469415590 TT/TT versus non- TT/TT (table 2). With IL28B rs8099917, the SVR rate in TT patients was 96.2% (50 of 52) compared to 62.5% (5 of 8) in patients with the IL28B TG genotype (p = 0.014, OR: 15.0, 95% CI: 2.0–112.1). Upon analysis of the allelic frequency of major allele T versus minor allele G, the same significant association was found (p = 0.001, OR: 10.3, 95% CI: 1.7–45.6). In contrast, for rs12979860 and ss469415590, no such differences in SVR were observed among patients infected with HCV-6 (80.6% in CC vs. 66.7% in non-CC, p = 0.593; 81.1% in TT/TT vs. 60% in non-TT/TT, p = 0.288).

Table 1. Characteristics of studies included in the meta-analysis of HCV-6 patients treated with PEG-IFN+RBV

<table>
<thead>
<tr>
<th>Study</th>
<th>Study design/publication type</th>
<th>Country (population)</th>
<th>Treatment regime (PEG-IFN+RBV)</th>
<th>Geno-type</th>
<th>Treatment, weeks</th>
<th>HCV-6 patients</th>
<th>Males, n (%)</th>
<th>Age, years</th>
<th>SVR, n (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Nguyen et al. [24], 2008</td>
<td>Non-RCT US (Asian American)</td>
<td>PEG-IFN α-2a  6  24  23 16 (69.6) 49±10 9 (39)</td>
<td>PEG-IFN α-2b  6  48  12 7 (58.3) 50±10 9 (75)</td>
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<tr>
<td>Fung et al. [27], 2008</td>
<td>Non-RCT Hong Kong (Chinese)</td>
<td>PEG-IFN α-2a  1  48  21 12 (57) 52 (30–63) 11 (52)</td>
<td>PEG-IFN α-2b  6  48  21 11 (52) 49.6 (14–64) 18 (86)</td>
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<tr>
<td>Lam et al. [26], 2010</td>
<td>RCT US (Asian American)</td>
<td>PEG-IFN α-2a  6  24  27 13 (48) 49.6±10 9 (70)</td>
<td>PEG-IFN α-2b  6  48  33 15 (46) 52.8±8.0 26 (79)</td>
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<tr>
<td>Nguyen et al. [30], 2010</td>
<td>Non-RCT US (Asian American)</td>
<td>PEG-IFN α-2a  1  48  70 51 (73) 49.6±10 9 (39)</td>
<td>PEG-IFN α-2b  6  48  27 19 (56) 49.4±10 9 (70)</td>
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<tr>
<td>Tsang et al. [19], 2010</td>
<td>Non-RCT Hong Kong (Chinese)</td>
<td>PEG-IFN α-2a  1  48  70 44 (63) 48 (18–64) 11 (52)</td>
<td>PEG-IFN α-2b  6  48  34 19 (56) 49.4±10.8 25 (49)</td>
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<tr>
<td>Zhou et al. [18], 2011</td>
<td>Non-RCT full China (Chinese)</td>
<td>PEG-IFN α-2a  1  48  39 22 (56.4) 15 (38) 23 (59.0)</td>
<td>PEG-IFN α-2b  6  48  22 14 (63.6) 18 (86.4) 26 (79.8)</td>
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<tr>
<td>Qing-Xian et al. [28], 2011</td>
<td>Non-RCT full China (Chinese)</td>
<td>PEG-IFN α-2a  6  48  84 NR NR 74 (88.1)</td>
<td>PEG-IFN α-2b  6  48  84 NR NR 74 (88.1)</td>
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<tr>
<td>Tangkijvanich et al. [21], 2012</td>
<td>Non-RCT full Thailand (Thai)</td>
<td>PEG-IFN α-2a  1  48  16 9 (56.3) 46.4±12.5 21 (60)</td>
<td>PEG-IFN α-2b  6  48  34 23 (67.6) 41.2±8.4 26 (76.5)</td>
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</tr>
<tr>
<td>Shao et al. [22], 2012</td>
<td>Non-RCT abstract China (Chinese)</td>
<td>PEG-IFN α-2a  6  48  28 NR NR 26 (92.8)</td>
<td>PEG-IFN α-2b  6  48  28 NR NR 26 (92.8)</td>
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</tr>
<tr>
<td>Mauss et al. [25], 2012</td>
<td>Non-RCT full Germany (Caucasian, African, Asian, Hispanic)</td>
<td>PEG-IFN α-2a  6  48  27 17 (63) 47 (37–52) 16 (59)</td>
<td>PEG-IFN α-2b  6  48  27 17 (63) 47 (37–52) 16 (59)</td>
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</tr>
<tr>
<td>Thu Thuy et al. [20], 2012</td>
<td>RCT full Vietnam (Vietnamese)</td>
<td>PEG-IFN α-2a  6  24  35 22 (62.9) 46.8±7.2 21 (60)</td>
<td>PEG-IFN α-2b  6  48  70 43 (61.4) 48.5±8.4 50 (71)</td>
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<td></td>
<td></td>
</tr>
<tr>
<td>Seto et al. [23], 2013</td>
<td>Non-RCT full Hong Kong (Chinese)</td>
<td>PEG-IFN α-2a  1  48  60 41 (68.3) 49 (14–71) 55 (91.7)</td>
<td>PEG-IFN α-2b  6  48  60 41 (68.3) 49 (14–71) 55 (91.7)</td>
<td></td>
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<td></td>
<td></td>
</tr>
<tr>
<td>Qing-Xian et al. [17], 2013</td>
<td>RCT abstract China (Chinese)</td>
<td>PEG-IFN α-2a  6  48  242 (total) NR NR 47 (68.1)</td>
<td>PEG-IFN α-2b  6  48  242 (total) NR NR 47 (68.1)</td>
<td></td>
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<td></td>
<td></td>
</tr>
<tr>
<td>Akkarathamrongsin et al. [29], 2014</td>
<td>Non-RCT full Thailand (Thai)</td>
<td>PEG-IFN α-2a  1  48  69 43 (62.3) 49.0±10.6 47 (68.1)</td>
<td>PEG-IFN α-2b  6  48  69 43 (62.3) 49.0±10.6 47 (68.1)</td>
<td></td>
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</tr>
</tbody>
</table>

NR = Not reported. Age is shown as mean ± SD or median (range). * Including HCV-2, 3 and HCV-6.

Fig. 2. Funnel plot of the included studies.

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Mild versus Advanced/Severe Hepatic Fibrosis

A total of 7 studies involving patients with mild and advanced fibrosis were included to evaluate the influence of fibrosis on treatment response \[23–27, 29, 30\]. Mild and advanced fibrosis were defined as F0–F2 and F3–F4 based on the METAVIR scoring system, respectively. In influence analysis, the SVR rate was 76% (95% CI: 0.67–0.84) for patients with mild fibrosis and 67% (95% CI: 0.61–0.77) for those with severe fibrosis. Our data indicate that the SVR rate in patients with mild fibrosis is significantly higher than in patients with advanced fibrosis (p < 0.001).

HCV-6 versus Other Genotypes

Six trials assessed treatment response in patients infected with HCV-6 compared with those infected with other genotypes. The overall SVR rate in HCV-6 patients treated with PEG-IFN+RBV is shown in Fig. 3.

<table>
<thead>
<tr>
<th>Study name</th>
<th>Year</th>
<th>Study design</th>
<th>Duration, weeks</th>
<th>Total</th>
<th>Proportion (95% CI)</th>
<th>Weight, %</th>
</tr>
</thead>
<tbody>
<tr>
<td>Seto et al. [23], 2013</td>
<td>2013</td>
<td>Non-RCT</td>
<td>24</td>
<td>50/52</td>
<td>0.39 (0.20, 0.61)</td>
<td>2.31</td>
</tr>
<tr>
<td>Nguyen et al. [24], 2010</td>
<td>2010</td>
<td>Non-RCT</td>
<td>48</td>
<td>5/8</td>
<td>15.0 (2.0, 112.1)</td>
<td>0.014</td>
</tr>
<tr>
<td>Lam et al. [26], 2010</td>
<td>2010</td>
<td>RCT</td>
<td>24</td>
<td>9/23</td>
<td>0.75 (0.43, 0.95)</td>
<td>1.44</td>
</tr>
<tr>
<td>Lam et al. [26], 2010</td>
<td>2010</td>
<td>RCT</td>
<td>48</td>
<td>19/27</td>
<td>0.70 (0.50, 0.81)</td>
<td>4.04</td>
</tr>
<tr>
<td>Fung et al. [27], 2008</td>
<td>2008</td>
<td>Non-RCT</td>
<td>48</td>
<td>26/33</td>
<td>0.79 (0.61, 0.91)</td>
<td>4.31</td>
</tr>
<tr>
<td>Nguyen et al. [30], 2010</td>
<td>2010</td>
<td>Non-RCT</td>
<td>48</td>
<td>17/27</td>
<td>0.63 (0.42, 0.81)</td>
<td>2.55</td>
</tr>
<tr>
<td>Nguyen et al. [30], 2010</td>
<td>2010</td>
<td>Non-RCT</td>
<td>48</td>
<td>34/46</td>
<td>0.74 (0.59, 0.86)</td>
<td>5.33</td>
</tr>
<tr>
<td>Zhou et al. [18], 2011</td>
<td>2011</td>
<td>Non-RCT</td>
<td>24</td>
<td>18/22</td>
<td>0.82 (0.60, 0.95)</td>
<td>3.17</td>
</tr>
<tr>
<td>Tsang et al. [19], 2010</td>
<td>2010</td>
<td>Non-RCT</td>
<td>48</td>
<td>53/70</td>
<td>0.76 (0.64, 0.85)</td>
<td>8.80</td>
</tr>
<tr>
<td>Qing-Xian et al. [28], 2011</td>
<td>2011</td>
<td>Non-RCT</td>
<td>48</td>
<td>74/84</td>
<td>0.88 (0.79, 0.94)</td>
<td>17.26</td>
</tr>
<tr>
<td>Shao et al. [22], 2012</td>
<td>2012</td>
<td>Non-RCT</td>
<td>48</td>
<td>26/28</td>
<td>0.93 (0.76, 0.99)</td>
<td>7.34</td>
</tr>
<tr>
<td>Mauss et al. [25], 2012</td>
<td>2012</td>
<td>Non-RCT</td>
<td>48</td>
<td>16/27</td>
<td>0.59 (0.39, 0.78)</td>
<td>2.55</td>
</tr>
<tr>
<td>Tangkijvanich et al. [21], 2012</td>
<td>2012</td>
<td>Non-RCT</td>
<td>24</td>
<td>26/34</td>
<td>0.76 (0.59, 0.91)</td>
<td>4.31</td>
</tr>
<tr>
<td>Thu Thuy et al. [20], 2012</td>
<td>2012</td>
<td>RCT</td>
<td>24</td>
<td>21/35</td>
<td>0.65 (0.53, 0.76)</td>
<td>7.34</td>
</tr>
<tr>
<td>Thu Thuy et al. [20], 2012</td>
<td>2012</td>
<td>RCT</td>
<td>48</td>
<td>50/70</td>
<td>0.71 (0.59, 0.82)</td>
<td>7.34</td>
</tr>
<tr>
<td>Seto et al. [23], 2013</td>
<td>2013</td>
<td>Non-RCT</td>
<td>48</td>
<td>55/60</td>
<td>0.92 (0.82, 0.97)</td>
<td>17.26</td>
</tr>
<tr>
<td>Akkarathamrongsin et al. [29], 2014</td>
<td>2014</td>
<td>Non-RCT</td>
<td>48</td>
<td>33/42</td>
<td>0.95 (0.67, 1.27)</td>
<td>1.08</td>
</tr>
<tr>
<td>Overall ((I^2 = 71.2%), p &lt; 0.0001)</td>
<td></td>
<td></td>
<td>48</td>
<td>504/661</td>
<td>0.80 (0.77, 0.83)</td>
<td>100</td>
</tr>
</tbody>
</table>

Fig. 3. Overall SVR rate in HCV-6 patients treated with PEG-IFN+RBV.

Table 2. Overall SVR rates in \(IL28B\) (rs8099917 and rs12979860) and \(IFNL4\) (ss4694 15590) patient groups

<table>
<thead>
<tr>
<th>Study name</th>
<th>SVR, n/total, n</th>
<th>OR</th>
<th>Lower limit</th>
<th>Upper limit</th>
<th>p value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Seto et al. [23], 2013</td>
<td>50/52</td>
<td>15.0</td>
<td>2.0</td>
<td>112.1</td>
<td>0.014</td>
</tr>
<tr>
<td>Akkarathamrongsin et al. [29], 2014</td>
<td>29/36</td>
<td>2.07</td>
<td>0.31</td>
<td>13.68</td>
<td>0.450</td>
</tr>
<tr>
<td>Akkarathamrongsin et al. [29], 2014</td>
<td>30/37</td>
<td>NR</td>
<td>NR</td>
<td>NR</td>
<td>NR</td>
</tr>
</tbody>
</table>
HCV-1 [18, 19, 21, 27, 29, 30]. Among these, 3 were direct comparisons of response to antiviral therapy by HCV-6 versus HCV-3 and/or HCV-2 patients. Patients with HCV-6 displayed significantly better SVR to PEG-IFN+RBV combination therapy than those with HCV-1 (80.1 vs. 55.3%), with a relative risk of 1.33 (95% CI: 1.14–1.53, p = 0.904; fig. 4). Our results are consistent with the majority of previous studies reporting that HCV-6 behaves in a more similar manner to HCV-2 and HCV-3 (SVR rates of 76–80%) [7–9].

**Discussion**

Although direct-acting antiviral agents have recently been approved for the treatment of chronic HCV infection, PEG-IFN+RBV combination therapy remains appropriate for HCV-6, since the relevant data obtained are insufficient to make general recommendations [32, 33] and not available in most countries in Southeast Asia [3]. Except for a previous study about HCV-4 [11], this is the first systematic review and meta-analysis to evaluate the influence of host and viral factors in patients with chronic HCV-6 treated with PEG-IFN+RBV. In our analysis, which included 14 studies comprising a total of 819 HCV-6 patients treated with PEG-IFN+RBV, the pooled SVR rate was 79.8% (95% CI: 0.77–0.83). No publication bias was found. Only 1 earlier study performed a meta-analysis of HCV-6 patients treated with PEG-IFN+RBV [31]. The pooled SVR rate in this report (75%) was slightly lower than in our results. However, their study did not include subgroup analysis of host and viral factors on SVR, while our investigation incorporated recent studies with a large number of patients and evaluated the influence of host and viral factors on SVR rates in HCV-6 patients. Differences in SVR rates between non-RCT and RCT were evident, with a significantly higher response in non-RCTs (p = 0.032).

The optimal duration of PEG-IFN+RBV therapy in HCV-6 patients depends on several factors and is yet to
be determined. Recent studies reported no significant differences in SVR rates in patients subjected to 48- and 24-week regimens. The abbreviated regimen has many advantages, including reducing unnecessary medication exposure, increasing affordability of treatment and maximizing the cost-effectiveness of therapy. Interestingly, in our subgroup analysis of 4 studies [20, 27, 32, 33], the SVR rate at 24 weeks of treatment was significantly lower than that at 48 weeks. Specifically, pooled SVR rates were 67% for 24 weeks (95% CI: 0.61–0.74; p = 0.038; $I^2 = 57.6\%$) and 84% for 48 weeks (95% CI: 0.80–0.87; p = 0.006; $I^2 = 59.7\%$). While 1 retrospective study demonstrated that SVR in HCV-1 and HCV-6 patients is not significantly different [19], the majority of investigations have shown that HCV-6 is closer to HCV-2 and -3 in that patients in these groups respond better to therapy than those with HCV-1 infection. With regard to the treatment regimen, we did not observe significant differences in SVR rates between patients treated with PEG-IFN α-2a and PEG-IFN α-2b (p = 0.352).

Several host factors (such as age, sex and fibrosis level) play an important role in response to HCV therapy [4, 5, 19]. In our analysis, no significant differences in SVR were found between males and females (OR: 0.81, 95% CI: 0.71–1.12, p = 0.71). Notably, the SVR rate in patients with mild fibrosis was markedly higher than that for advanced fibrosis (76% for mild fibrosis and 67% for severe fibrosis; p < 0.001). Baseline viral load has been established as the determinant of treatment response. Lower starting HCV-RNA levels are reported to be associated with higher SVR rates [34]. Consistent with these results, our data showed that lower baseline viral load is signifi-
cantly related to increased SVR than the higher baseline viral load (OR: 4.01, 95% CI: 1.62–5.43, p < 0.001). However, larger-scale studies are required to further validate our findings on the effects of gender, fibrosis level and baseline viral load on treatment outcome.

A number of studies have reported that rs8099917 and rs12979860 polymorphisms in IL28B are significantly associated with treatment outcome, especially in HCV-1 patients. Also, a previous meta-analysis on HCV-4 reported an association with SVR with favorable IL28B polymorphisms (rs12979860 CC and rs8099917 TT genotypes) [11]. The 2 studies on IL28B in our subgroup analyses showed a favorable association between SVR and rs8099917 TT genotype versus GT/GG (p = 0.014, OR: 4.01, 95% CI: 1.62–5.43, p < 0.001). How- ever, based on the small numbers and limited patient data, these findings should be interpreted with caution.

In our subgroup analysis, IL28B polymorphism was significantly associated with treatment outcome, especially in HCV-6 patients. Our findings on the effects of gender, fibrosis level and baseline viral load on treatment outcome are in line with other studies [29]. The efficacy of PEG-IFN+RBV varies depending on the type of treatment regimen and dose schedule, so we were not able to perform sensitivity analyses on these parameters owing to limited data. Despite the above drawbacks, this study provides solid evidence that PEG-IFN+RBV treatment significantly improves the SVR rate in HCV-6 patients and validates the effects of specific host and virus factors on treatment outcome.

In conclusion, the PEG-IFN+RBV combination is effective in HCV-6 patients, with a pooled SVR rate of 79.8% in our study. Moreover, treatment outcomes of HCV-6 patients are superior to HCV-1 and comparable to those of HCV-2 and -3. As HCV-1, the optimal treatment duration of HCV-6 should be 48 weeks, although a shortened treatment duration of 24 weeks could be sufficient. Considering the level of fibrosis in HCV-6 patients may significantly benefit from the recently FDA-approved triple therapies, where available. Gender and type of PEG-IFN did not affect SVR rates. In addition, UL28B and IFNL4 polymorphisms are not significantly associated with treatment outcome in HCV-6 patients.

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