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***Prospective Study***

**Importance of virological response in the early stage of telaprevir-based triple therapy for hepatitis C**

Hiramine S *et al*. Viral response to telaprevir-based triple therapy

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**Abstract**

**AIM:** To investigate the efficacy of virological response (VR) to telaprevir　(TVR)-based triple therapy in predicting the treatment outcome.

**METHODS:** This prospective, multicenter study consisted of 253 Japanese patients infected with hepatitis C virus (HCV) genotype 1b. All received 12 wk of TVR in combination with 24 weeks of pegylated-interferon-α and ribavirin. Serum HCV RNA was tested at weeks 1, 2, 3, 4, 6, 8, 12, 16, 20, and 24. VR was defined as undetectable serum HCV RNA. Sustained virological response (SVR) is VR at 24 wk after the end of treatment and is regarded as a successful outcome.

**RESULTS:** Of 253 patients, 207 (81.8%) achieved SVR. The positive predictive value of VR for SVR was 100% at week 2, after which it gradually decreased, however it remained over 85% to week 12. The negative predictive value (NPV) gradually increased, reaching 100% at week 12. The upslope of the NPV showed a large increase from week 4 (40.6%) to week 6 (82.4%). There was a moderate concordance between the SVR and VR at week 6 (kappa coefficient = 0.44) although other VRs had poor concordance to SVR. Multiple logistic regression analysis extracted VR at week 6 (*P* < 0.0001, OR = 63.8) as an independent factor contributing to SVR, in addition to the IL28B SNPs and response to previous pegylated-interferon-α and ribavirin therapy.

**CONCLUSION:** VR at week 6, but not at week 4, is an efficient predictor of both SVR and non-SVR to TVR-based triple therapy.

**Key words:** Chronic hepatitis C; Direct-acting antiviral agent; Rapid virological response; Early virological response; Response-guided treatment

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**Core tip:** An undetectable viral level at week 4 or 12 was a good predictor of the outcome of conventional interferon therapy without direct-acting antiviral agents (DAAs); however, the transition of the viral level during DAA therapy has not been well documented. In this prospective multicenter study, the authors did frequent testing of 253 patients to investigate viral activity during triple therapy containing telaprevir, the first approved DAA, and an undetectable viral level at week 6 was the most effective predictor. It is suggested that the most predictive time point in DAA therapy is different from the traditional markers.

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**INTRODUCTION**

Since the approval of Interferon (IFN)-α for the treatment of hepatitis C virus (HCV) infected patients in 1991, the treatment regimens used have seen great change and improvements. The rate of sustained virological response (SVR) to dual therapy with ribavirin (RBV) and pegylated IFN (PegIFN) of patients with HCV genotype 1 has remained approximately 50%[[1-3](#_ENREF_1)], but with telaprevir (TVR), the first direct-acting antiviral agent (DAA) approved in the United States, Canada, the European Union, and Japan, the rate of SVR to a triple therapy of PegIFN-α, RBV and TVR against HCV genotype 1 has reached over 70%[[4-6](#_ENREF_4)]. New DAAs have since been developed and approved, and it has become common for patients to be treated with IFN therapy that contains a DAA or a DAA based IFN-free oral therapy. Unfortunately, the cost of DAAs can be prohibitive and some have serious side effects. If patients destined not to achieve SVR can be identified before or in the early stage of treatment, they can avoid starting or continuing a treatment that has no possibility of success. Studies of factors that can be used to predict the outcome of DAA based therapies are needed.

For dual therapy with PegIFN-α/RBV, it has been consistently reported that virological response (VR: undetectable serum HCV RNA) at week 4 or 12 of therapy is strongly associated with the outcome[[7-10](#_ENREF_7)]. Rapid virological response (RVR), VR at week 4, and early virological response (EVR), VR at week 12, are terms coined before the approval of DAAs, and this criterion is still used for determining the best form of antiviral treatment management, as recommended by international consensus conferences such as the American Association for the Study of Liver Diseases (AASLD)[[11](#_ENREF_11)] and the European Association for the Study of the Liver (EASL)[[12](#_ENREF_12)]. However, the viral kinetics during DAA therapy are unclear and it is possible that the time point most predictive of success might be different than under the older regimens.

To clarify the timing of VR most predictive of SVR during DAA based treatment, we measured serum HCV RNA at seven time points during the early stage of TVR-based triple therapy for Japanese patients to determine their relation to SVR.

**MATERIALS AND METHODS**

***Patients***

Since 2004, the Kyushu University Liver Disease Study (KULDS) Group has conducted prospective, multicenter studies to investigate the efficacy and safety of antiviral treatment for chronic hepatitis C patients[[3](#_ENREF_3),[6](#_ENREF_6)]. For this study, we recruited 253 chronic hepatitis C patients infected with HCV genotype 1b who started TVR-based triple therapy between December 2011 and December 2012 and whose 24-wk post-therapy follow-up ended by June 2013. Exclusion criteria were as reported previously[[6](#_ENREF_6)]. The study was conducted in accordance with the ethical principles of the Declaration of Helsinki and was approved by the Ethics Committee of our hospital. Informed consent was obtained from all patients before enrollment. The study was registered as a clinical trial on the University Hospital Medical Information Network (ID 000009711).

***Treatment response***

VR is defined as undetectable serum HCV RNA. Successful treatment is SVR at 24 weeks after the end of treatment. Relapse is defined as VR during the treatment, but non-SVR. Patients with HCV RNA detectable throughout treatment are classified as non-responders. Patients who had not been previously treated with PegIFN-α/RBV therapy are classified as treatment naïve.

***Clinical and laboratory assessment***

Clinical parameters included hemoglobin, platelet count, serum albumin, aspartate aminotransferase (AST), alanine aminotransferase (ALT), γ-glutamyl-transpeptidase (γGTP), low-density lipoprotein (LDL) cholesterol, ferritin, and estimated glomerular filtration rate (eGFR). HCV RNA was tested at baseline, weeks 1, 2, 3, 4, 6, 8, 12, 16, 20, and 24 during the treatment and at weeks 4, 8, 12, and 24 after the end of treatment. We defined the early stage of treatment as the period between day 1 and week 12. The timing of VR in the early stage of treatment was evaluated for candidate predictors of SVR. Liver biopsy was done for 154 (60.9%) patients before the induction of therapy. For each specimen, the stage of fibrosis (F0-4) and grade of activity (A0-3) were established according to the Metavir score[[13](#_ENREF_13)].

***Determination of HCV markers***

The baseline and follow-up tests for HCV viremia were done by real-time polymerase chain reaction (PCR) assay (COBAS TaqMan HCV test, Roche Diagnostics, Tokyo, Japan), with a detectability of ≥ 15 IU/mL and a linear dynamic range of 1.2-7.8 log IU/mL. HCV genotype and the core amino acid substitution at position 70 of the HCV genome were determined before treatment for all patients. HCV genotype was determined by sequence determination in the 5’ non-structural region of the HCV genome, followed by phylogenetic analysis [[14](#_ENREF_14)].

***Interleukin 28B and inosine triphosphate pyrophosphatase polymorphism genotyping***

Human genomic DNA was extracted from peripheral blood. Genotyping by the single-nucleotide polymorphism (SNP) of the interleukin 28B (IL28B) (rs8099917) gene was done using the TaqMan Allelic Discrimination Demonstration Kit (7500 Real-Time PCR System; Applied Biosystems, Foster City, CA). Patients were genotyped as TT, TG, or GG at the polymorphic site. Similarly, genotyping by the SNP of the inosine triphosphate pyrophosphatase (ITPA) (rs1127354) gene was done using the TaqMan Allelic Discrimination Demonstration Kit. Patients were genotyped as CC, CA, or AA at the polymorphic site. IL28B and ITPA SNPs were not available for only two patients (1.2%).　Although rs12979860, another IL28B SNP that is also strongly correlated to the therapeutic outcome, has been reported[[15](#_ENREF_15)], we determined only rs8099917 because it was previously reported that rs8099917 and rs12979860 represent 98.6% of the Japanese population[[16](#_ENREF_16)].

***Therapeutic protocol***

All patients received 12-wk triple therapy that included TVR (2,250 mg/day) (Telavic; Mitsubishi Tanabe Pharma, Osaka, Japan), PegIFN-α-2b (60-150 μg/wk) (PEG-Intron; MSD, Tokyo, Japan) and RBV (600-1000 mg/d) (Rebetol; MSD), followed by a 12-wk dual therapy that included PegIFN-α-2b and RBV. TVR (750 mg) was administered orally three times a day at an 8-h interval after each meal. PegIFN-α-2b was injected subcutaneously once weekly at a dose of 1.5 μg/kg. RBV was given orally at a daily dose of 600-1000 mg based on body weight (600 mg for patients weighing < 60 kg, 800 mg for those weighing 60-80 kg, and 1000 mg for those weighing > 80 kg). The above durations and dosages are those approved by the Japanese Ministry of Health, Labor, and Welfare. If marked anorexia, an elevation of serum creatinine, or severe anemia developed, the TVR dose could be reduced to 1500 mg/d (750 mg at a 12-h interval, after meals). The method of RBV/TVR dose reduction in the case of anemia was as reported[[17](#_ENREF_17)]. The completed assigned total cumulative dosages of each drug were calculated by reviewing the patients’ medical records and by counting the pills not consumed by each patient. The actual dosage of TVR given is calculated as the percentage of target TVR (2250 mg/d). The dosages of PegIFN-α-2b and RBV were calculated individually as averages on the basis of body weight at baseline.

***Definition of positive predictive value and negative predictive value***

To evaluate the precision rate of on-treatment VR for predicting the outcome, we calculated the positive predictive value (PPV) and the negative predictive value (NPV). PPV is defined as the probability that a patient with a given on-treatment VR will achieve SVR. In contrast, NPV is defined as the probability that a patient without a given on-treatment VR will not achieve SVR.

***Statistical analysis***

Statistical analyses were performed using the SAS system, version 9.1.3 (SAS Institute, Cary, NC). Continuous data are expressed as median with interquartile range. Univariate analyses were performed using the χ2 test, Fisher’s exact test, paired *t*-test, or Mann-Whitney *U* test, as appropriate, with SVR as the outcome. Kappa coefficient was used for the analysis of the concordance between SVR and VR at the seven time points. To identify independent factors predictive of SVR, variables that reached the *P* < 0.1 level in univariate tests were used as candidates in the multiple logistic regression analysis. Continuous parameters that were significant in univariate analysis were converted into categorical variables by dichotomizing at the round number closest to their median for analysis in the multiple logistic regression model. Because liver histology data was missing for 99 (39.1%) patients, it was excluded from the multiple logistic regression model. A *P* value less than 0.05 was regarded as statistically significant in all analyses.

**RESULTS**

***Transition of VR rate during telaprevir-based triple therapy and follow-up***

Of the 253 patients, 207 (81.8%) achieved SVR, 37 (14.6%) relapsed and 9 (3.6%) were non-responders. The VR rate increased dramatically to week 6 (5.9%, 22.0%, 53.4%, 74.0%, and 93.1% at weeks 1, 2, 3, 4, and 6, respectively). Two hundred and forty-four patients (96.4%) had achieved VR by week 12. The rate gradually decreased to 81.8% after the end of treatment. A graph of the VR rates classified by SVR status is shown in Figure 1. Comparison of the VR rates of the SVR and non-SVR groups in the early stage of treatment shows that although there was no statistical difference at weeks 1 or 12 (7.3% *vs* 0.0% and 100% *vs* 97.3%, respectively), the rates were significantly higher for the SVR than for the non-SVR group for weeks 2 to 8 (26.9% *vs* 0.0%, 59.8% *vs* 25.6%, 81.6% *vs* 35.0%, 98.5% *vs* 65.0%, and 98.5% *vs* 86.5% at weeks 2, 3, 4, 6, and 8, respectively. *P* < 0.0001 at weeks 2, 3, 4, and 6. *P* = 0.0027 at week 8).

***Demographic and clinical features of patients, by SVR status***

The patient characteristics are summarized by the clinical outcome in Table 1. Sex, age, genotype of IL28B SNP (rs8099917), hemoglobin level, platelet count, serum albumin, AST, and LDL-cholesterol at baseline were significantly correlated with SVR in the univariate analysis (all *P* < 0.05). The rate of non-responders to previous PegIFN-α/RBV therapy was significantly higher in the non-SVR group than in the SVR group (44.8% *vs* 12.5%, *P* < 0.0001). The SVR rate significantly decreased as the stage of fibrosis progressed, but was not related to the grade of activity.

***Concordance between SVR and VR during the early stage of treatment***

The PPV and NPV, calculated on the basis of VR in the early stage of treatment, are shown in Table 2. The PPV of VR for SVR was 100% at week 2, after which it gradually decreased, however it remained over 85% to week 12. The NPV gradually increased, reaching 100% at week 12. The upslope of the NPV showed a large increase from week 4 (40.6%) to week 6 (82.4%). Kappa coefficients were calculated to evaluate the concordance between SVR and VRs (Table 2). There was a moderate concordance between the SVR and VR at week 6 (kappa coefficient = 0.44, 95%CI: 0.24-0.76), although the other VRs had poor concordance to SVR.

***Multivariate Analysis for factors predictive of SVR***

Multiple logistic regression analysis was done to determine factors predictive of SVR. VR at week 6, which had the highest kappa coefficient, was included as a candidate in order to compare its predictive power. IL28B SNP (rs8099917) genotype [*P* < 0.0001, odds ratio (OR) = 8.24, 95%CI: 2.81-26.8], response to previous PegIFN-α/RBV therapy (*P*=0.0281, OR = 3.29, 95%CI: 1.14-9.46), and VR at week 6 (*P <* 0.0001, OR = 63.8, 95%CI: 10.8-563) were extracted as factors contributing to SVR. VR at week 6 had a high statistical correlation with SVR (Table 3).

**DISCUSSION**

VR in the early stage of treatment has in the past been used to manage the treatment of patients with HCV. Since the advent of DAAs, no research has been published that describes the detailed transition of serum HCV RNA during DAA therapy. Although the guidelines of AASLD and EASL recommend checking VR at weeks 4 (RVR) and 12 (EVR) for the assessment of initial response to therapy and adherence, other time points such as weeks 1, 2, 3, 6, and 10 were not mentioned in these guidelines[[11](#_ENREF_11),[12](#_ENREF_12)]. It is possible that RVR and EVR were chosen because they are traditional conventional markers that have been used in PegIFN-α/RBV therapy and because the efficacy of other time points in DAA-containing therapy have not yet been fully investigated. By the testing done at frequent intervals in this prospective multicenter study of 253 patients infected with HCV genotype 1b, we were able to show that the transition during treatment with a DAA is different than what was seen in the past. VR at week 6 had a high PPV (88.6%), NPV (82.4%), and kappa coefficient (0.44), which indicate its usefulness as a single time point for predicting both SVR and non-SVR during TVR-based triple therapy. In addition, multiple logistic regression analysis that included pretreatment factors such as the patient’s genotype, laboratory parameters at baseline and response to previous therapy extracted VR at week 6 as an independent factor contributing to SVR.

For dual therapy with PegIFN-α and RBV, RVR and EVR were correlated with the outcome and have traditionally been utilized as predictors. It has consistently been reported that RVR has a high PPV, around 90%[[7-9](#_ENREF_7)], making it a useful marker for the prediction of SVR. In contrast, EVR has a high NPV, over 90%, making it a useful predictor of non-SVR[[10](#_ENREF_10)]. In our study, the rates of EVR were not significantly different between the SVR and the non-SVR group. Although RVR had a high PPV, the NPV showed a sharp rise, from 45.7% at week 4 to 87.0% at week 6. This suggests that in DAA therapy, which has a direct mechanism and much stronger power to eliminate HCV than does dual therapy, the most useful and meaningful time points for predicting the outcome may be different than in PegIFN-α/RBV therapy.

Although the DAAs strongly eliminate HCV, they are costly and some have serious side effects, such as the rash and anemia that often accompany TVR. To avoid unproductive expenditures and side effects, attempts have been made to establish response-guided treatment regimens that include early stopping rules for unproductive DAA therapy[[5](#_ENREF_5),[18](#_ENREF_18)]. It has been suggested that patients who have a rapid decline in their viral level can be treated with a shorter treatment duration while preserving the high rate of SVR and that treatment can be discontinued earlier than in the past for patients who are unlikely to respond the treatment. Our results showed that checking VR at week 6 would contribute to shortening the duration of TVR-based triple therapy. Furthermore, because at week 6 both SVR and non-SVR can be predicted at a single time point, unnecessary testing can be eliminated, which will contribute to patient comfort and economic efficiency.

One of the limitations of our study is that TVR is no longer the standard of care in many countries. It is not recommended for the treatment of patients with decompensated cirrhosis or in a post-liver transplantation setting, and it should not be administered as co-medication. In addition, TVR can cause serious rash and anemia. In Japan, IFN-based therapy with RBV and Simeprevir, a new NS3/4A inhibitor, has become the standard of care against HCV genotype 1[[19](#_ENREF_19)]. More recently, a number of novel DAAs, such as NS5A and NS5B inhibitors, have been developed and approved, and the current standard of care in the US is an IFN-free DAA regimen[[11](#_ENREF_11)]. Although our results might seem late to the game, TVR-containing treatment will continue to be an option in regions of the world where the newly approved DAAs are not available or as an option for patients with no alternative. Another limitation of our study is that the patients were all Japanese and infected with HCV genotype 1b. The rate of SVR significantly differs by the race of the patient and the genotype of HCV[[20](#_ENREF_20)]. Hence, our results may not be able to be simply extended to the up-to-date IFN-free DAA regimens or to every patient with chronic hepatitis C; however, the results give useful information because 253 patients were enrolled and frequent HCV RNA testing during DAA-containing therapy was analyzed that included numerous variables, including the genotype and laboratory parameters of each patient in this study. We believe that our study is sufficiently reliable to show that the most efficient time point for checking VR in DAA therapy might be different than the RVR and EVR that was developed for earlier therapies. Our results will need to be validated for the current DAA regimens, and further studies of patients with other HCV genotypes and of other racial cohorts will be necessary.

It is also a limitation of our study that we did not test for mutations of various HCV strains. Many studies have revealed that the variations in the amino acid sequences of HCV affect the antiviral activity of DAAs. Bartels *et al*[[21](#_ENREF_21)] , using a direct-sequencing technique, reported that the mutant strain resistant to NS3/4A protease inhibitors was detected in 2% of treatment naïve patients and that it was the pre-existing dominant strain in some of the patients. Nasu *et al*[[22](#_ENREF_22)], using an ultra-deep sequencing technique, found some resistant mutations in a surprisingly high percentage of treatment naïve patients. In the coming era of IFN-free regimens, it will be essential to determine the mutations of the patients' HCV strains before treatment.

In conclusion, VR at week 6 is the time point most predictive of both SVR and non-SVR in the early stage of TVR-based triple therapy. This shows the possibility that the most efficient time point for checking VR in DAA therapy might be different than the conventional RVR and EVR. Our results will need to be validated in light of the newly developed DAA regimens.

**COMMENTS**

***Background***

Since direct-acting antiviral agents (DAAs) were approved for the treatment of chronic hepatitis C, the treatment success rate has greatly improved. However, DAAs are costly and some of have serious side effects. To avoid unproductive expenditures and side effects, attempts have been made to predict the treatment response by checking the serum viral load of patients during the early stage of treatment. It has been suggested that patients who have a rapid decline in viral level can be treated with a shorter treatment duration while preserving the high rate of success and that patients who are unlikely to respond treatment should discontinue it early.

***Research frontiers***

An undetectable viral level at week 4 or 12 has consistently been correlated to the outcome of conventional interferon therapy without DAAs, with RVR and early virological response commonly used as predictors of treatment success. The transition of the viral level during DAA therapy has not been well documented. In this prospective multicenter study, the authors did frequent testing of 253 patients to investigate viral activity during triple therapy containing telaprevir, the first approved DAA.

***Innovations and breakthroughs***

This is the first study to report the detailed transition of the viral level during the early stage of DAA therapy for chronic hepatitis C. The important finding is that an undetectable serum viral level at week 6, not at week 4 or 12, is the most efficient predictor of the outcome.

***Applications***

Checking the serum viral level at week 6 would be useful for establishing a response-guided treatment regimen for patients treated with DAAs, which would help reduce the total duration of treatment.

***Terminology***

Virological response (VR) is defined as undetectable serum HCV RNA. Sustained virological resoponse (SVR) is VR at 24 wk after the end of treatment and is regarded as successful treatment. We evaluated the ability of the VR between weeks 1 and 12 during the treatment to predict SVR or non-SVR.

***Peer-review***

Hiramine *et al* in this article describe in detail the factors which can be used for the prediction of therapeutic outcome. The limitation of the study is as mentioned by the authors too that all the results are only for HCV genotype 1 Japanese patients. IL28b polymorphism are now a known factor influencing the treatment response, author has taken this as a predictive factor too. But the polymorphism at IL28b rs8099917 is studied while another very important polymorphism rs12979860 which is well documented to influence the therapeutic outcome is not studied, it would be great if authors have studied that too on these patients. Overall the study is well-designed and well written.

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**Figure 1 Transition of the virological response rate by sustained virological response status.** The virological response (VR) rates were significantly higher in the SVR than the non-SVR group between weeks 2 and 8 (26.9% *vs* 0.0%, 59.8% *vs* 25.6%, 81.6% *vs* 35.0%, 98.5% *vs* 65.0% and 98.5% *vs* 86.5% at weeks 2, 3, 4, 6 and 8, respectively. *P* < 0.0001 at weeks 2, 3, 4 and 6. *P* = 0.0027 at week 8), although there was no statistical difference at week 1 or 12 (7.3% *vs* 0.0% and 100% *vs* 97.3%, respectively).

**Table 1 Patient characteristics**

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  | **All (*n =* 253)** | | **SVR (*n =* 207)** | | **non-SVR (*n =* 46)** | | ***P* value** |
| Sex, male (%) | 123 | (48.6) | 108 | (52.2) | 15 | (32.6) | 0.0153 |
| Age (yr) | 61 | (12.5) | 60 | (12) | 63.5 | (11.25) | 0.0340 |
| Body mass index (kg/m2) | 23.4 | (3.9) | 23.4 | (3.9) | 23.9 | (3.8) | 0.2198 |
| Baseline HCV RNA (log10 IU/mL) | 6.5 | (0.9) | 6.5 | (0.9) | 6.4 | (0.7) | 0.4468 |
| IL28B SNP (rs8099917), TT/TG or GG (%)1 | 186/65 | (74.1/25.9) | 166/40 | (80.6/19.4) | 20/25 | (44.4/55.6) | < 0.0001 |
| ITPA SNP (rs1127354), CC/CA or AA (%)1 | 193/58 | (76.9/23.1) | 157/49 | (76.2/23.8) | 36/9 | (80.0/20.0) | 0.5802 |
| Hemoglobin level (g/L) | 138 | (22) | 140 | (21) | 134 | (20) | 0.0031 |
| Platelet count (× 109/L) | 157 | (69) | 159 | (65) | 129 | (69) | 0.0006 |
| Serum albumin (g/L) | 40 | (6.0) | 40 | (6.0) | 39 | (5.0) | 0.0143 |
| Aspartate aminotransferase (U/L) | 48 | (42) | 46 | (43) | 59 | (34.5) | 0.0350 |
| Alanine aminotransferase (U/L) | 54 | (58) | 53 | (64) | 58 | (44) | 0.4955 |
| γ-glutamyl-transpeptidase (U/L) | 40 | (51) | 39 | (47) | 46 | (59) | 0.1270 |
| LDL-cholesterol (mg/dL) | 95 | (38) | 98 | (36) | 75 | (35) | < 0.0001 |
| Ferritin (μg/L) | 164.6 | (232.3) | 160.5 | (223.2) | 181.7 | (253.9) | 0.3583 |
| Estimated glomerular filtration rate (mL/min/1.73 m2) | 79.4 | (19.1) | 79.7 | (18.9) | 77.7 | (19.4) | 0.6210 |
| Response to previous PegIFN-α/RBV therapy |  |  |  |  |  |  | < 0.0001 |
| Treatment naïve, *n* (%) | 92 | (36.4) | 81 | (39.4) | 11 | (23.9) |  |
| Prior relapse, *n* (%) | 113 | (44.7) | 100 | (48.1) | 13 | (28.3) |  |
| Prior non-response, *n* (%) | 48 | (19.0) | 26 | (12.5) | 22 | (44.8) |  |
| Liver histology |  |  |  |  |  |  |  |
| Stage, F0-2/F3-4 (%) | 96/58 | (62.3/37.7) | 87/38 | (69.6/30.4) | 9/20 | (31.0/69.0) | < 0.0001 |
| Grade, A0-1/A2-3 (%) | 54/100 | (35.1/64.9) | 45/80 | (36.0/64.0) | 9/20 | (31.0/69.0) | 0.6136 |
| Not determined, *n* | 99 |  | 82 |  | 17 |  |  |

1IL28B and ITPA SNPs were not available for only two patients (1.2%). Continuous variables are expressed as median (interquartile range). *P* value draws a comparison between SVR and non-SVR patients. SVR: Sustained virological response; HCV: Hepatitis C virus; IL28B: Interleukin 28B; SNP: Single-nucleotide polymorphism; ITPA: Inosine triphosphate pyrophosphatase; LDL: Low-density lipoprotein.

**Table 2 Precision rate for the prediction of sustained virological response and non-sustained virological response in the early stage of telaprevir-based triple therapy**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | **Patients who achieved SVR/ Patients with VR, *n*** | **PPV (%)** | **Patients who did not achieve SVR/Patients without VR, *n*** | **NPV (%)** | **Kappa coefficient**  **(95%CI)** |
| Week 1 | 14/14 | 100 | 45/222 | 20.3 | 0.03 (0.01-0.05) |
| Week 2 | 52/52 | 100 | 44/185 | 23.8 | 0.12 (0.08-0.16) |
| Week 3 | 113/124 | 91.1 | 33/109 | 30.3 | 0.22 (0.12-0.33) |
| Week 4 | 168/182 | 92.3 | 26/64 | 40.6 | 0.38 (0.24-0.51) |
| Week 6 | 202/228 | 88.6 | 14/17 | 82.4 | 0.44 (0.27-0.61) |
| Week 8 | 200/232 | 86.2 | 5/8 | 62.5 | 0.18 (0.02-0.34) |
| Week 12 | 198/234 | 84.6 | 1/1 | 100 | - |

PPV: Positive predictive value, the probability that a patient with a given on-treatment virological response (VR) will achieve sustained virological response (SVR); NPV: Negative predictive value, the probability that a patient without a given on-treatment VR will not achieve SVR.

**Table 3 Factors contributing to sustained virological response**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | **Univariate analysis** | | **Multivariable analysis** | | |
| OR | *P* value | OR (95%CI) | | *P* value |
| Sex (male to female) | 2.25 | 0.0153 |  |  |  |
| Age (< 60 yr to ≥ 60 yr) | 1.79 | 0.0822 |  |  |  |
| IL28B SNPs (rs8099917) (TT to TG/GG) | 5.19 | < 0.0001 | 8.24 | (2.81-26.8) | < 0.0001 |
| Hemoglobin level (≥ 140 g/L to < 140 g/L) | 2.13 | 0.0245 |  |  |  |
| Platelet count (≥ 150 × 109/L to < 150 × 109/L) | 3.21 | 0.0005 |  |  |  |
| Serum albumin (> 35 g/L to ≤ 35 g/L) | 2.51 | 0.0308 |  |  |  |
| Aspartate aminotransferase (< 50 U/L to ≥ 50 U/L) | 2.30 | 0.0123 |  |  |  |
| LDL-cholesterol (≥ 95 mg/dL to < 95 mg/dL) | 4.39 | < 0.0001 |  |  |  |
| Response to previous PegIFN-α/RBV therapy (naïve/relapse to non-response) | 6.38 | < 0.0001 | 3.29 | (1.14-9.46) | 0.0281 |
| VR at week 6 | 31.1 | < 0.0001 | 63.8 | (10.8-563) | < 0.0001 |

*P* value draws a comparison between SVR and non-SVR patients. SVR: Sustained virological response; HCV: Hepatitis C virus; IL28B: Interleukin 28B; SNP: Single-nucleotide polymorphism; LDL: Low-density lipoprotein; PegIFN: Pegylated interferon; RBV: Ribavirin; VR: Virological response.