

Research Paper

## Weight loss, leukopenia and thrombocytopenia associated with sustained virologic response to Hepatitis C treatment

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### Abstract

**OBJECTIVE:** To identify apparent adverse effects of treatment of chronic hepatitis C and their relationship to sustained virologic response (SVR).

**METHODS:** A retrospective study was conducted of all Hepatitis C virus (HCV)-infected patients treated with pegylated interferon and ribavirin in an academic ambulatory infectious disease practice. Clinical and laboratory characteristics were compared between patients with SVR and without SVR.

**RESULTS:** Fifty-four patients completed therapy with the overall SVR rate of 76%. SVR was associated with genotype non-1 ( $P=0.01$ ), weight loss more than 5 kilograms ( $P=0.04$ ), end of treatment leukopenia ( $P=0.02$ ) and thrombocytopenia ( $P=0.05$ ). In multivariate analysis, SVR was significant associated with HCV genotype non-1 (Adjusted Odd Ratio [AOR] 15.22; CI 1.55 to 149.72;  $P=0.02$ ), weight loss more than 5 kilograms, (AOR 5.74; CI 1.24 to 26.32;  $P=0.04$ ), and end of treatment white blood cell count level less than  $3 \times 10^3$  cells/ $\mu$ l (AOR 9.09; CI 1.59 to 52.63;  $P=0.02$ ). Thrombocytopenia was not significant after adjustment. Other factors including age, gender, ethnicity, injection drug use, viral load, anemia, alanine transaminase level, and liver histology did not reach statistical significance.

**CONCLUSION:** Besides non-1 genotype, SVR was found to be independently associated with weight loss during therapy, and leukopenia at the end of HCV treatment. These correlations suggest continuation of therapy despite adverse effects, may be of benefit.

Key words: Hepatitis C, pegylated interferon, ribavirin, weight loss, leukopenia, thrombocytopenia.

### INTRODUCTION

Hepatitis C virus (HCV) infection is a major cause of liver diseases and liver cancer (1-3). Among the six genotypes of HCV, the most common genotypes of HCV in the United States are genotype 1 (approximately 75%), genotype 2 (15%), and genotype 3 (7%) (2-5). At present, the standard treatment for chronic HCV genotype 1 infection is 48 weeks with a combination of pegylated (long-acting) inter-

feron alfa-2a or alfa-2b plus ribavirin. Outcomes are measured by sustained viral response (SVR), defined as an undetectable viral load 24 weeks after the end of therapy. HCV genotype 1 has been reported to have a 54-56% SVR (2, 6-9). Prior studies have shown better response rates with genotype 2 or 3 with a 24 week-course of therapy (2, 8,9). Response rates have been found higher in Caucasians (52%) compared to

African-Americans (28%) (2). Response rates are reported lower with initial levels of HCV RNA >600,000 IU/ml, male gender, high body weight, and advanced liver fibrosis (11-17). Recent studies have also shown a poorer response to treatment associated independently with HCV genotype 1 infection but a better response with weight loss (18). Limited knowledge exists regarding the influence of on-treatment factors during therapy. The purpose of our analysis was to identify clinical, biological, virological and histological predictive factors that may be helpful in guiding decisions during therapy, and for counseling patients about outcomes.

## MATERIALS AND METHODS

**Study population.** We conducted a retrospective cohort study of 54 adults (age  $\geq$  18 years) patients who were diagnosed with HCV infection and completed treatment with pegylated interferon plus ribavirin through an academic ambulatory infectious disease practice from January 2004 to December 2007. The study was approved by the University of Hawaii Committee on Human Studies (CHS # 1541). Patients infected with HCV genotype 1 completed a 48 week-course with once weekly injections of pegylated interferon alfa-2a or alfa 2b (180  $\mu$ g) plus ribavirin (1000 or 1200 mg/day in divided dose) for 48 weeks. Patients infected with HCV genotypes 2 or 3 completed a 24 week-course with once weekly injections of pegylated interferon alfa-2a (180  $\mu$ g) plus ribavirin (800 mg/day in divided doses). SVR rates were measured at 24 weeks after the end of therapy. Interferon dosing was not adjusted but ribavirin occasionally was. Hematology growth factors were occasionally used but not consistently.

**Study design and definitions.** The study patients were identified from the clinic's medical records using the International Statistical Classification of Diseases and Related Health Problems (ICD) code of 070.44 and 070.54. During the period of study, 78 patients started on therapy. Twenty four patients were not included in the study analysis due to fail to complete treatment and evaluation. Of these, 13 failed to return for follow-up, 6 stopped because of adverse effects, 3 for poor response, and 2 because of death. Finally, a total of 54 charts were reviewed in the patients who completed therapy and returned for follow-up at 24 weeks after treatment. Information on demographic characteristics, co-morbid conditions, genotypes of HCV, laboratory data, treatment, and follow-up data were collected by using a study data collection form. SVR was indicated by undetectable HCV RNA at 24 weeks after therapy with combined pegylated interferon alfa-2a or alfa-2b plus ribavirin.

Genotypes of HCV were defined using the Bayer TRUGENE HCV Genotyping Test. Serum concentration of HCV RNA was determined shortly before study and at 24 weeks after complete course of treatment (treatment duration: 48 weeks in genotype 1 and 24 weeks in genotype 2 or 3) by the COBAST (TaqMan) test, which has a limit of detection of 28 IU/mL. Body weight measurement and laboratory data were collected and compared at the beginning and at the end of therapy. The Knodell scoring system for liver biopsy (obtained at baseline) result was used as the indicator of histologic activity.

**Statistical analysis.** Descriptive statistics were produced comparing SVR versus non-SVR group. Categorical variables were compared using the Pearson's  $\chi^2$  or Fisher's exact test, as appropriate. The distribution of continuous data was evaluated by the O'Brien test for homogeneity of variances. When necessary, an appropriate normalizing or variance stabilizing transformation was applied. Analysis of variance was used to compare groups. Cut-offs of continuous data was determined by evaluating the median results of the groups and deciding among the investigators clinical importance. From the data, clinically relevant cut-offs determined by the investigators were as follows: for weight loss (more than 5 kilograms.), leukopenia (WBC less than  $3 \times 10^3$  cells/ $\mu$ l) and thrombocytopenia (platelets less than  $1 \times 10^5$  cells/ $\mu$ l). Logistic regression and multivariate analysis were performed to identify independent predictors for SVR. Statistical significance was considered with *P* value of 0.05. Odds ratio (OR) with 95% confidence interval (CI) was reported in categorical data. All statistical analyses were conducted using SPSS for Window software, version 15.0 (SPSS Inc, Chicago, IL).

## RESULTS

A total of 54 medical records of patients who completed HCV therapy with follow-up viral load result at 24 weeks were reviewed. The majority of the study population was male (67%) and Caucasian (57%) with the mean age of 52.5 years (Table 1). Fifty six percent were infected with genotype 1, 16% with genotype 2 and 28% with genotype 3. Clinical characteristics, relevant laboratory data and liver biopsy results of all patients are shown in Table 1. The major risk factors for acquiring HCV infection were related to injection drug use and tattooing. Mild and moderate liver inflammation determined by Knodell score was presented in the majority of patients while cirrhosis was rarely observed. A total of 22 patients had liver fibrosis: portal fibrosis was the most common (n=11, 50%), followed by septal fibrosis (n=4, 18%),

bridging fibrosis (n=4, 18%), periportal fibrosis, and only one patient with cirrhosis pattern (n=1, 5%). Forty-one percent of the patient population had a history of alcohol abuse and 35% had underlying psychiatric problems. Concomitant psychiatric disorders among this patient population included depression (38%), anxiety (18%), schizophrenia (6%) and bipolar disorder (3%). Underlying medical problems included hypertension (15%) and diabetes mellitus (DM) (14%).

Demographics, clinical and laboratory data of patients (percent and means) were compared between patients with SVR and without SVR as shown in Table 2. All continuous variables were normally distributed and did not require log transformation. SVR was significantly affected by genotype non-1 ( $P=0.01$ ) as well as low white blood cell (WBC) count (leukopenia) at the end of treatment ( $P=0.02$ ) compared to patients without SVR. Moreover, low platelet count (thrombocytopenia) at the end of treatment was associated with SVR ( $P=0.05$ ). Age, ethnicity,

gender, drug abuse, and histology were not significantly associated with SVR. To allow for clinical applicability, logistic regression with clinical cut-offs in the significant relevant covariants, found in the linear regression models, were performed. Laboratory characteristics and categorical variables for the predictors of SVR were compared by using logistic regression and multivariate analysis (categorical data) as displayed in Table 3. In multivariate analysis, SVR was significant associated with HCV genotype non-1 (Adjusted Odd Ratio [AOR] 15.22; CI 1.55 to 149.72;  $P=0.02$ ), weight loss more than 5 kilograms, (AOR 5.74; CI 1.24 to 26.32;  $P=0.04$ ), and end of treatment white blood cell count level less than  $3 \times 10^3$  cells/ $\mu$ l (AOR 9.09; CI 1.59 to 52.63;  $P=0.02$ ). Thrombocytopenia was not significant after adjustment. Other factors including age, gender, ethnicity, injection drug use, viral load, anemia, alanine transaminase level, and liver histology and did not reach statistical significance.

**Table 1:** Baseline demographic, clinical characteristics and laboratory data of the study population

Characteristics	Total, n = 54
Male gender, n (%)	36 (67)
Ethnicity, n (%)	
Caucasian	31 (57)
Asian	10 (19)
Hispanic	7 (13)
Hawaiian	6 (11)
Others	3 (6)
Age, years, mean $\pm$ SD	52.5 $\pm$ 5.8
HCV genotypes, n (%)	
Genotype 1	30 (56)
Genotype 2	9 (16)
Genotype 3	15 (28)
Weight, kilograms, mean $\pm$ SD	85.0 $\pm$ 19.1
BMI, kilograms/m <sup>2</sup> , mean $\pm$ SD	28.2 $\pm$ 5.5
Baseline HCV RNA ( $\times 10^6$ IU/mL $\pm$ SD)	6.45 $\pm$ 13.4
HCV RNA > 600,000 IU/mL, n (%)	47 (87)
ALT >90 U/L, n (%)	45 (83)
Total bilirubin, mg/dL, mean $\pm$ SD	1.0 $\pm$ 0.5
Serum albumin, g/dl, mean $\pm$ SD	3.9 $\pm$ 0.3
Liver biopsy results;	n = 38
Knodell score, mean $\pm$ SD	8.0 $\pm$ 3.2
Minimal inflammation ( $\leq 4$ ), n (%)	4 (11)
Mild inflammation (5-8), n (%)	18 (47)
Moderate inflammation (9-12), n (%)	14 (37)
Marked inflammation ( $\geq 13$ ), n (%)	2 (5)
Risk factors for acquiring HCV infection, n (%)	
Injection drug use	38 (70)
Tattooing	41 (76)
Sexual partners with HCV infection	10 (19)
Blood transfusion	5 (9)

History of cocaine use, n (%)	22 (41)
History of alcohol abuse, n (%)	22 (41)
History of methamphetamine use, n (%)	2 (4)
HIV co infection, n (%)	3 (6)
Underlying psychiatric disorders (depression, bipolar disorder, schizophrenia and anxiety), n (%)	19 (35)
Underlying medical conditions (diabetes mellitus, hypertension, others), n (%)	28 (52)

**Abbreviations:** ALT = alanine transaminase; BMI = body mass index; HCV = hepatitis C virus; HIV = human immunodeficiency virus; RNA = ribonucleic acid; SD = standard deviation.

**Table 2:** Comparison of demographics, clinical characteristics and laboratory data between patients with and without sustained virological response (SVR).

Characteristics	Patients with SVR (n=41)	Patients without SVR (n=13)	P-Value
Male gender, n, (%)	30 (73)	6 (46)	0.11
Caucasians, n (%)	23 (56)	8 (62)	0.94
Age <50 years, n (%)	25 (60)	10 (77)	0.24
Genotype 1, n (%)	18 (44)	12 (92)	<b>0.01*</b>
Risk factors for acquiring HCV infection, n (%)			
Injection drug use	26 (63)	12 (92)	0.28
Blood transfusion	4 (10)	1 (8)	0.21
<b>Baseline characteristic and risk factors</b>			
Alcohol abuse, n (%)	14 (47)	8 (62)	0.22
Cocaine use, n (%)	13 (32)	7 (54)	0.15
Methamphetamine use, n (%)	0 (0)	2 (15)	0.15
HIV co-infection, n (%)	1 (3)	2 (15)	0.47
Psychiatric disorders, n (%)	14 (34)	5 (38)	0.75
Knodell score < 11, n (%)	16 (64) <sup>a</sup>	10 (77) <sup>b</sup>	0.07
HCV RNA (x10 <sup>6</sup> IU/mL, mean ± SD)	4.6±5.6	8.3±15.6	0.22
Weight (kilograms, mean ± SD)	92.1±22.2	77.9±16.0	0.10
<b>Baseline laboratory data</b>			
WBC (X 10 <sup>3</sup> cells/μl, mean ± SD)	6.6±3.1	6.8±2.8	0.97
Hemoglobin (g/dl, mean ± SD)	15.3±0.9	14.8±1.2	0.91
Platelet (X 10 <sup>5</sup> cells/μl, mean ± SD)	184.0±93.8	239.7±99.7	0.14
ALT (U/L, mean ± SD)	78.5±70.5	63.6±28.8	0.31
Total bilirubin (g/dl, mean ± SD)	1.1±0.7	0.8±0.2	0.47
Albumin (g/dl, mean ± SD)	3.8±0.4	4.0±0.2	0.10
<b>End of treatment laboratory data</b>			
WBC (X 10 <sup>3</sup> cells/μl, mean ± SD)	2.9 ±1.1	3.6±0.9	<b>0.02*</b>
Hemoglobin (g/dl, mean ± SD)	11.8 ± 1.8	11.7±1.5	0.42
Platelet (X 10 <sup>6</sup> cells/μl, mean ± SD)	122.4 ± 96.8	207.2±86.5	<b>0.05*</b>
ALT (U/L, mean ± SD)	35.3 ± 26.8	36.6±28.1	0.60
Total bilirubin (g/dl, ) mean ± SD)	1.4 ± 1.4	0.8± 0.3	0.13
Albumin (g/dl, mean ± SD)	3.8 ±0.3	3.8±0.5	0.95
<b>Clinically significance relevantce characteristic and laboratory data</b>			
Weight loss > 5 Kilograms, n (%)	29 (71)	5 (38)	<b>0.04*</b>
End of treatment WBC count less than 3 X 10 <sup>3</sup> cells/μl	34 (83)	3 (23)	<b>0.01*</b>
End of treatment platelet count less than 1X 10 <sup>5</sup> cells/μl	36 (88)	7(54)	0.06

**Note:** \* Statistical significant. Biopsy results; <sup>a</sup> total n for SVR = 25; <sup>b</sup> total n for non SVR =13.

**Abbreviations:** ALT = alanine transaminase; HCV = hepatitis C virus; RNA = ribonucleic acid; SD = standard deviation; SVR = sustained virological response; WBC = white blood cell count.

**Table 3:** Logistic regression and multivariate analysis for the predictors of SVR (Clinical significance relevance characteristic and laboratory data)

Variables	OR (95% CI)	P-value	Adjusted# OR (95% CI)	P-value
Age < 50 year	2.5 (0.57-10)	0.24	1.38 (0.23-8.33)	0.73
Genotype Non-1	15.33 (1.82-129.18)	0.01*	15.22 (1.55-149.72)	0.02*
Weight loss more than 5 kilograms	3.86 (1.05-14.82)	0.04*	5.74 (1.24-26.32)	0.04*
End of treatment WBC count less than $3 \times 10^3$ cells/ $\mu$ l	9.52 (1.86-50.00)	0.01*	9.09 (1.59-52.63)	0.02*
End of treatment platelet count less than $1 \times 10^5$ cells/ $\mu$ l	6.9 (0.8-5.9)	0.06	2.43 (0.13-45.45)	0.51

Note: \* Statistical significant. # Model adjusted with all table variables.

Abbreviation; CI = confidence interval; OR = odds ratio; SVR = sustained virological response; WBC = white blood cell.

## DISCUSSION

The overall rate of response was 76% comparable with those reported in the previous US studies of 55%. Responses were better for both genotype 1 (56% vs 52%) and others (96% vs 81-84%).<sup>2</sup> Our study strongly indicated that being infected with HCV genotype 1 was associated with treatment failure. The higher treatment response in our study compared to the US literatures may have occurred from differences in our demographic population. Our study had less African Americans (only 1 patient), and a lower prevalence of HCV genotype 1 and HIV co-infection. Higher response rates among Asians have been noted before. This may be related to the *IL28B* gene polymorphism as it is apparently associated with higher response to treatment among Asian Americans patients compared to Caucasians, Hispanics and African Americans, respectively (19).

In our study, the patients with SVR had significant weight loss compared to those without SVR. The association between body weight, weight loss and SVR is controversial. Some previous studies showed that initial body weight is an independent risk factor for HCV treatment failure, and weight reduction was associated with treatment success (10, 20). However, other studies demonstrated that weight loss did not improve the treatment outcomes (15, 21). It is known that interferon therapy is associated with weight loss, but the mechanism and pathophysiology are still unclear. The reasons for weight loss among the study population are unclear. Many patients complained of loss of appetite or ate less related to the fatigue. Studies suggest that interferon decreases appetite via induction of tumor necrosis factors (TNF), the level of leptin, insulin and cytokines but findings are not consistent (21-24). Alternatively, several studies showed that TNF levels were not increased during interferon

therapy and postulated other mechanisms of weight loss including changes in level of leptin and insulin which affect glucose metabolism (21, 22). In our studies, weight loss usually began within the first few dosages of interferon treatment. It was also associated with fatigue, nausea, vomiting. The weight loss was usually sustained during therapy or gradually progressed. On the follow-up after complete treatment, patients usually gained weight but did not achieve their baseline bodyweight.

In our study, patients with SVR had significantly lower WBC and platelet count at the end of treatment compared to those without SVR. These findings suggested that patients who developed leukopenia and/or thrombocytopenia during the interferon treatment responded well to the therapy and these side effects, if not severe, may not be indications for withholding or reducing the dose of the treatment. We hypothesized that the greater cytopenia is a marker for greater TNF activity in a specific treatment recipient which translates into greater SVR. Common side effects of interferon treatment that need to be monitored during the therapy are anemia, thrombocytopenia and leukopenia (25-29). These side effects may prevent physicians from continuing their patients on interferon therapy or reducing dosage (28, 29). A comparison of outcome by the approach of reducing interferon and/or ribavirin versus use of growth factors to stimulate promotion of white blood cell or red blood cell or platelet counts has not been done but is clearly needed. We believe that the differences in magnitude of interferon effects on individual pharmacokinetics, pharmacodynamics, and genetic determination of interferon susceptibility may play an important role in each patient's treatment response and side effect experience.

Our patients had a high rate of mental illnesses and social problems because of our open referral sys-

tem and the population at risk. However, the majority of patients tolerated HCV treatment which can be attributed to our supportive staff and follow-up system. The potential interaction between psychiatric illness and its treatment with HCV therapy could not be assessed with our limited study.

This study is limited in that it is retrospective and examined only subjects who completed HCV therapy. The association of SVR with factors reported by others, such as age, histological (fibrosis) score, viral load and alanine transaminase level was not significant in our study. This may be related to the small sample size. Other factors such as adherence and insulin resistance were not assessed. The majority of patient with genotype 2 and 3 did not have liver biopsy reports. Further studies with larger sample size and molecular testing may enhance our understanding of the relationship between interferon and antiviral therapy in chronic hepatitis C infection.

In conclusion, the virological outcomes of the current HCV infection therapy in our study were comparable with the outcomes of the previous studies. The treatment failure was significantly associated with genotype 1. Side effects of interferon therapy including leukopenia, thrombocytopenia and weight loss were predictors of good treatment response. Giving these findings, a careful assessment is needed in regards to continuing therapy with interferon and ribavirin despite opposed adverse effects by laboratory parameters and weight loss.

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## CONFLICT OF INTEREST

Alan Tice has recently been a consultant or speaker or investigator for Roche, Schering, 3 Rivers, Human Genome Science, and Novartis.

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