

Table III. Comparison of characteristics between low insulin group and high insulin group.

| Variable                                       | Number or mean (SD)      |                           | P      |
|--|--------------------------|---------------------------|--------|
|  | Low insulin group (n=70) | High insulin group (n=70) |        |
| Onset age, y.o.                                | 64.7 (9.5)               | 65.5 (9.5)                | 0.620  |
| Gender   |                          |                           | 0.999  |
| Male   | 55                       | 55                        |        |
| Female   | 15                       | 15                        |        |
| BMI, kg/m <sup>2</sup>                         | 22.5 (3.0)               | 23.7 (3.0)                | 0.019  |
| Alcohol intake                                 |                          |                           | 0.687  |
| <80 g/day                                      | 53                       | 55                        |        |
| ≥80 g/day                                      | 17                       | 15                        |        |
| Etiology                                       |                          |                           | 0.818  |
| HBsAg(+)                                       | 16                       | 13                        |        |
| HCVAb(+)                                       | 48                       | 51                        |        |
| Non-B, non-C                                   | 6                        | 6                         |        |
| Underlying liver diseases and Child-Pugh grade |                          |                           | 0.539  |
| CH   | 17                       | 13                        |        |
| LC grade A                                     | 37                       | 35                        |        |
| LC grade B                                     | 14                       | 17                        |        |
| LC grade C                                     | 2                        | 5                         |        |
| Total bilirubin, ng/ml                         | 1.5 (2.9)                | 1.3 (1.3)                 | 0.634  |
| Ferritin, ng/ml                                | 305.2 (351.7)            | 303.2 (341.2)             | 0.974  |
| Serum iron, μg/ml                              | 155 (77)                 | 148 (74)                  | 0.571  |
| Fasting insulin, μIU/ml                        | 4.3 (2.2)                | 15.8 (10.6)               | <0.001 |
| Fasting blood glucose, mg/dl                   | 94.0 (16.6) (n=31)       | 111.4 (70.5) (n=31)       | 0.185  |
| HOMA-R, %                                      | 0.9 (0.6) (n=13)         | 4.8 (2.4) (n=15)          | <0.001 |
| Total adiponectin, μg/ml                       | 7.9 (4.3)                | 8.3 (5.3)                 | 0.611  |
| HMW, μg/ml                                     | 3.7 (2.7)                | 4.1 (3.3)                 | 0.473  |
| MMW, μg/ml                                     | 1.9 (1.2)                | 1.8 (1.1)                 | 0.570  |
| LMW, μg/ml                                     | 2.3 (1.1)                | 2.5 (1.4)                 | 0.600  |

calculated in 28 patients, and the level of HOMA-R in the high insulin group was significantly higher than that in the low insulin group. Table IV shows the comparison of characteristics of HCC between the two groups. Patients with more than three HCC lesions and diffuse HCC were more prevalent in the low insulin group than in the high insulin group. The other characteristics of HCC did not differ substantially between the two groups.

Fig. 2A indicates the cumulative survival rates of all stage HCC patients between the low insulin group (70 of 140) and the high insulin group (70 of 140). There was no significant difference between the two groups ( $P=0.235$ ). Next, to evaluate the relationship of the fasting insulin level with the prognosis of early stage HCC patients, we analyzed the cumulative survival rates in HCC patients with TNM stage I and II disease ( $n=92$ ). As shown in Fig. 2B, the high insulin group (49 of 92) exhibited a poor prognosis with a significant difference in comparison to the low insulin group (43 of 92) ( $P=0.018$ ).

*Association of fasting total adiponectin level with prognosis of HCC.* Similarly, we evaluated the association of the total adiponectin level with the prognosis of HCC. One hundred and forty patients were divided into 2 groups in terms of the 50th percentile of the value of total adiponectin (6.95 μIU/ml). The mean level of total adiponectin in the low adiponectin group (<6.95 μIU/ml,  $n=70$ ) was 4.5 μg/ml. That in the high adiponectin group (≥6.95 μIU/ml,  $n=70$ ) was 11.8 μg/ml. We estimated the cumulative survival rates of all stages of HCC and early stage HCC between the low adiponectin group and the high adiponectin group. Fig. 3A and B show each result. No significant differences were found in all stages of HCC, or in the early stage of HCC (all stage HCC: low group vs. high group;  $P=0.886$ , early stage HCC: low group vs. high group;  $P=0.804$ ).

*Univariate and multivariate analyses of the factors associated with HCC prognosis.* Univariate and multivariate analyses

Table IV. Comparison of HCC characteristics between low insulin group and high insulin group.

| Variable               | Number or mean (SD)      |                           | P     |
|------------------------|--------------------------|---------------------------|-------|
|                        | Low insulin group (n=70) | High insulin group (n=70) |       |
| Tumor size, cm         | 3.6 (3.2)                | 3.2 (2.5)                 | 0.479 |
| Number of tumor lesion |                          |                           | 0.032 |
| 1                      | 36                       | 39                        |       |
| 2                      | 10                       | 19                        |       |
| 3 - and diffuse        | 24                       | 12                        |       |
| TNM stage              |                          |                           | 0.300 |
| I                      | 22                       | 17                        |       |
| II                     | 21                       | 32                        |       |
| III                    | 21                       | 16                        |       |
| IV                     | 6                        | 5                         |       |
| AFP, ng/ml             | 5136.8 (30566.9)         | 10487.6 (83045.4)         | 0.614 |
| Therapy                |                          |                           | 0.492 |
| Surgical resection     | 5                        | 2                         |       |
| PEIT and/or RFA        | 27                       | 26                        |       |
| TACE or TAI            | 35                       | 38                        |       |
| Others                 | 3                        | 4                         |       |

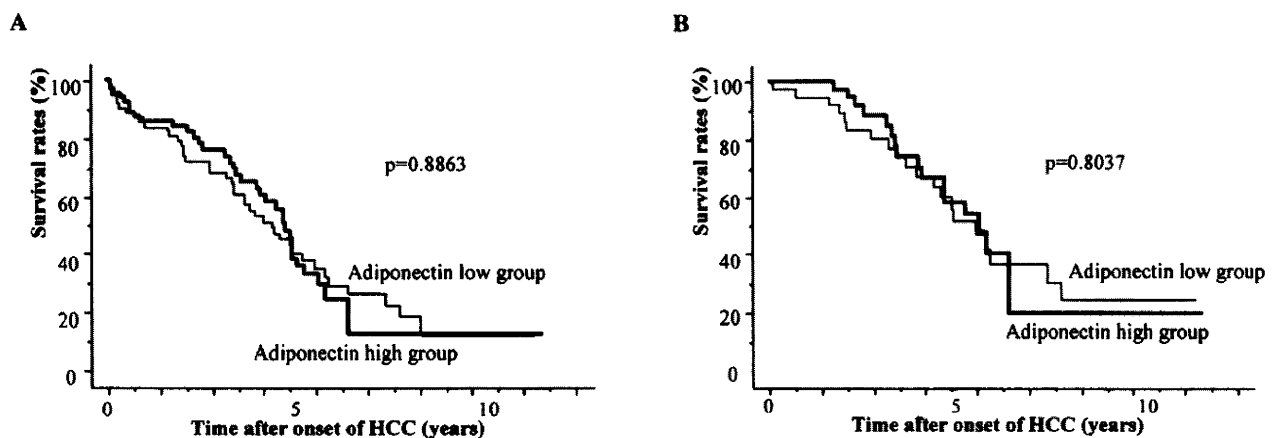


Figure 3. Kaplan-Meier curves for survival between low adiponectin group (thin line) and high adiponectin group (heavy line) in all stages of HCC patients (n=140) (A) and in TNM stage I + II HCC patients (n=92) (B).

using the Cox proportional hazards model in 140 patients diagnosed with HCC were performed to identify the relevant independent prognostic factors in all stages of HCC. In a univariate analysis, the following three factors significantly influenced the prognosis: alcohol intake (excessive drinker, RR 2.033, 95% CI 1.206-3.425,  $P=0.008$ ), Child-Pugh grade (grade C, RR 9.906, 95% CI 3.547-27.666,  $P<0.001$ ), and therapy for HCC (TACE or TAI, RR 1.856, 95% CI 1.143-3.015,  $P=0.012$ ). However, a multivariate analysis revealed that only two factors influenced the HCC prognosis significantly: Child-Pugh grade (grade C, RR 9.807, 95% CI 2.710-30.471,  $P<0.001$ ) and therapy for HCC (TACE or TAI, RR 1.803, 95% CI 1.104-2.943,  $P=0.018$ ).

Next, univariate and multivariate analyses in 92 patients diagnosed with HCC, all TNM stage I or II, were performed to identify the independent prognostic factors of early stage HCC. In the univariate analysis, the following three factors significantly influenced prognosis: alcohol intake (excessive drinker, RR 2.488, 95% CI 1.160-5.319,  $P=0.019$ ), Child-Pugh grade (grade B, RR 4.582, 95% CI 1.370-15.323,  $P=0.014$ , grade C, RR 41.104, 95% CI 6.403-263.831,  $P<0.001$ ), and the value of insulin ( $>7.73 \mu\text{IU/ml}$ , RR 2.196, 95% CI 1.126-4.292,  $P=0.021$ ) (Table V).

Similarly, when we performed multivariate analysis, only two factors, the Child-Pugh grade and the level of fasting insulin influenced the prognosis of early stage HCC with a

Table V. Univariate analyses of prognosis factors for HCC of TNM stage I and II.

| Variable                                       | Relative risk (95% CI) | P      |
|--|------------------------|--------|
| Onset age, >60 y.o.                            | 1.248 (0.536-2.907)    | 0.606  |
| Gender, male                                   | 1.637 (0.715-3.745)    | 0.244  |
| BMI, >25.0 kg/m <sup>2</sup>                   | 1.488 (0.746-2.967)    | 0.260  |
| Alcohol intake, ≥80g/day                       | 2.488 (1.160-5.319)    | 0.019  |
| Background                                     |                        | 0.647  |
| non-B, non-C                                   | -                      | -      |
| HBsAg(+)                                       | 1.111 (0.230-5.369)    | 0.896  |
| HCVAb(+)                                       | 1.566 (0.366-6.700)    | 0.545  |
| Underlying liver diseases and Child-Pugh grade |                        | <0.001 |
| CH   | -                      | -      |
| LC Child-Pugh grade A                          | 2.531 (0.866-7.395)    | 0.090  |
| LC Child-Pugh grade B                          | 4.582 (1.370-15.323)   | 0.014  |
| LC Child-Pugh grade C                          | 41.104 (6.403-263.831) | <0.001 |
| Serum ferritin, <185 ng/ml                     | 1.193 (0.621-2.295)    | 0.596  |
| Serum iron, <141 μg/ml                         | 1.222 (0.641-2.331)    | 0.542  |
| Fasting insulin, >7.73 μIU/ml                  | 2.196 (1.126-4.292)    | 0.021  |
| Fasting blood glucose, >110 mg/dl              | 0.949 (0.118-7.634)    | 0.961  |
| HOMA-R, >2.0%                                  | 4.762 (0.475-47.619)   | 0.184  |
| Total adiponectin, >6.95 μg/ml                 | 0.921 (0.479-1.767)    | 0.804  |
| HMW, 3.0 μg/ml                                 | 0.799 (0.418-1.529)    | 0.498  |
| MMW, >1.6 μg/ml                                | 1.171 (0.613-2.232)    | 0.633  |
| LMW, >2.1 μg/ml                                | 1.038 (0.544-1.984)    | 0.908  |
| Therapy, TACE or TAI                           | 1.429 (0.743-2.748)    | 0.285  |

Table VI. Multivariate analyses of prognosis factors for HCC of TNM stage I and II.

| Variable                                       | Relative risk (95% CI) | P     |
|--|------------------------|-------|
| Alcohol intake, ≥80 g/day                      | 2.217 (0.933-5.263)    | 0.071 |
| Underlying liver diseases and Child-Pugh grade |                        | 0.022 |
| CH   | -                      | -     |
| LC Child-Pugh grade A                          | 2.884 (0.975-8.531)    | 0.056 |
| LC Child-Pugh grade B                          | 3.771 (1.099-12.529)   | 0.035 |
| LC Child-Pugh grade C                          | 19.039 (2.782-130.298) | 0.003 |
| Fasting insulin, >7.73 μIU/ml                  | 2.033 (1.019-4.049)    | 0.044 |

significant difference: Child-Pugh grade (grade B, RR 3.771, 95% CI 1.099-12.529, P=0.035, grade C, RR 19.039, 95% CI 2.782-130.298, P=0.003), and the level of fasting insulin (>7.73 μIU/ml, RR 2.033, 95% CI 1.019-4.049, P=0.044) (Table VI).

*Association of fasting insulin and total adiponectin level with recurrence-free survival.* To evaluate the association of fasting insulin level with the recurrence-free survival time, 59 patients who underwent curative therapy, defined as a condition characterized by the no findings of recurrence over six months after the initial therapy, were extracted from 140 patients and subjected to analysis. Of 59 patients, the mean level of insulin in the low insulin group (<7.73 μIU/ml, n=32) or that in the high insulin group (>7.73 μIU/ml, n=27) was 3.8 μIU/ml or 14.4 μIU/ml, respectively. Fig. 4A indicates the cumulative recurrence-free survival rates of 59 patients who underwent curative therapy. The high insulin group exhibited a lower recurrence-free survival with a significant difference in comparison to the low insulin group (P=0.017).

Similarly, we evaluated the association of the total adiponectin level with the recurrence-free survival time. The mean level of total adiponectin in the low adiponectin group (<6.95 μIU/ml, n=28) or that in the high adiponectin group (≥6.95 μIU/ml, n=31) was 4.5 μIU/ml or 11.7 μIU/ml, respectively. We compared the cumulative recurrence-free survival rates of HCC between the low adiponectin group and the high adiponectin group, but no significant difference was found (Fig. 4B).

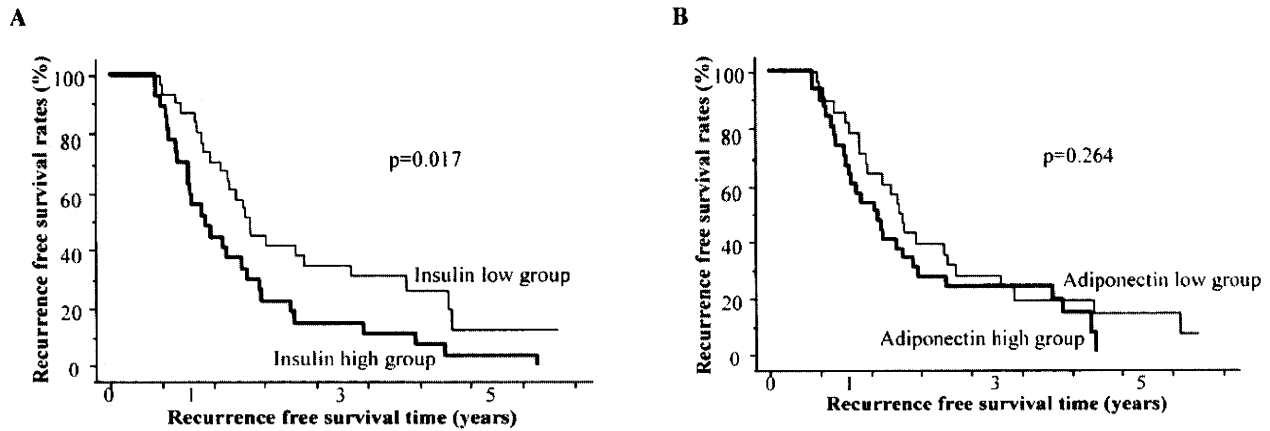


Figure 4. Kaplan-Meier curves for recurrence-free survival in HCC patients who underwent curative therapy (n=59) between low insulin group (thin line) and high insulin group (heavy line) (A) and between low adiponectin group (thin line) and high adiponectin group (heavy line) (B).

Table VII. Univariate analyses of the factor that contribute to recurrence-free survival.

| Variable                                       | Relative risk (95% CI) | P     |
|--|------------------------|-------|
| Onset age, >60 y.o.                            | 1.905 (0.951-3.871)    | 0.069 |
| Gender, male                                   | 1.984 (0.355-1.243)    | 0.201 |
| BMI, >25.0 kg/m <sup>2</sup>                   | 2.268 (0.805-4.367)    | 0.014 |
| Alcohol intake, ≥80 g/day                      | 1.289 (0.653-2.538)    | 0.465 |
| Background                                     |                        | 0.671 |
| Non-B, non-C                                   | -                      | -     |
| HBsAg(+)                                       | 1.302 (0.289-5.855)    | 0.731 |
| HCVAb(+)                                       | 1.566 (0.390-6.773)    | 0.505 |
| Underlying liver diseases and Child-Pugh grade |                        | 0.093 |
| CH   | -                      | -     |
| LC Child-Pugh grade A                          | 2.300 (1.095-4.831)    | 0.028 |
| LC Child-Pugh grade B                          | 2.883 (1.086-7.650)    | 0.034 |
| LC Child-Pugh grade C                          | 3.655 (0.774-17.263)   | 0.102 |
| Serum ferritin, <185 ng/ml                     | 1.157 (0.663-2.019)    | 0.607 |
| Serum iron, <141 μg/ml                         | 1.379 (0.772-2.464)    | 0.278 |
| Fasting insulin, >7.73 μIU/ml                  | 1.946 (1.117-3.378)    | 0.019 |
| Fasting blood glucose, >110 mg/dl              | 4.975 (0.903-27.778)   | 0.065 |
| HOMA-R, >2.0%                                  | 4.255 (0.816-22.222)   | 0.086 |
| Total adiponectin, >6.95 μg/ml                 | 1.376 (0.784-2.410)    | 0.266 |
| HMW, >3.0 μg/ml                                | 1.076 (0.611-1.893)    | 0.799 |
| MMW, >1.6 μg/ml                                | 1.258 (0.711-2.222)    | 0.430 |
| LMW, >2.1 μg/ml                                | 1.012 (0.572-1.792)    | 0.967 |
| Therapy, TACE or TAI                           | 1.165 (0.646-2.101)    | 0.610 |

Univariate and multivariate analyses of the factors associated with recurrence-free survival. To clarify the factors that contribute to recurrence-free survival except for tumoral

factors, univariate and multivariate analyses were performed using the Cox proportional hazards model in 59 patients who underwent curative therapy. In a univariate analysis, only two

Table VIII. Multivariate analyses of the factors that contribute to recurrence-free survival.

| Variable                           | Relative risk (95% CI) | P     |
|------------------------------------|------------------------|-------|
| BMI, >25.0 kg/m <sup>2</sup>       | 1.992 (1.026-3.861)    | 0.042 |
| Fasting insulin, >7.73 $\mu$ IU/ml | 1.767 (1.004-3.117)    | 0.049 |

factors significantly influenced the recurrence-free survival: BMI (>25.0 kg/m<sup>2</sup>, RR 2.268, 95% CI 0.805-4.367, P=0.014) and the value of insulin (>7.73  $\mu$ IU/ml, RR 1.946, 95% CI 1.117-3.378, P=0.019) (Table VII). Multivariate analysis showed that both factors influenced the recurrence-free survival with a significant difference: BMI (>25.0 kg/m<sup>2</sup>, RR 1.992, 95% CI 1.026-3.861, P=0.042) and the value of insulin (>7.73  $\mu$ IU/ml, RR 1.767, 95% CI 1.004-3.117, P=0.049) (Table VIII).

## Discussion

Several prior studies have reported that the coexistence of DM influences the prognosis of HCC patients (10,11,28,29). However, the mechanism responsible for this finding remains unclear. Since the glucose tolerance of an individual is defined by the potential insulin secretion from  $\beta$ -cells and by the insulin sensitivity of target tissues including the liver, serum levels of fasting and postprandial insulin could differ in each HCC patient. In addition, advanced liver fibrosis is directly linked to an increase in the insulin resistance in HCV-infected patients (13,30).

In the present study, we therefore focused on the serum level of insulin rather than on the glucose tolerance in the HCC patients. Our study indicates that a high value of fasting insulin heralds not only a poor prognosis in the early stage of HCC but also a high recurrence rate in the curative HCC. There are a few studies on the prognostic value of hyperinsulinemia on patients with HCC. Saito *et al* have demonstrated that the area under the plasma insulin curve for the oral glucose tolerance test can serve as a significant prognostic tool, and can assist in forecasting the doubling time of HCC (16), and that continuous infusion of octreotide in five patients inhibited insulin secretion resulting in a decrease in the HCC growth rate.

Komura *et al* reported that insulin therapy for coexisting DM is an independent risk factor for HCC recurrence after a curative resection (10). Taken together, it is possible that hyperinsulinemia promotes the progression and development of HCC. This is consistent with the results from the following *in vitro* studies, that insulin has the potential to accelerate the growth of hepatoma cells and inhibits apoptosis through the upregulation of Bcl-xl (14), and that insulin stimulates the motility and invasiveness of hepatoma cells (31). In addition, there have been several clinical studies supporting the association between hyperinsulinemia and the advancement of cancers. A high level of fasting insulin is associated with distant recurrence and death in early stage breast cancer (32). High insulin levels are associated with a poorer prognosis in prostate cancer and endometrial cancer, and malignant

degeneration of adenomatous polyps (33-36). These findings suggest that, in addition to an effect on glucose metabolism, insulin functions to promote the proliferation and metastasis of various types of cancer cells.

Hyperinsulinemia is inextricably linked to insulin resistance of the peripheral tissues including the liver. In our study, HOMA-R, a good indicator of insulin resistance, was not associated with a poor prognosis in early stage HCC (univariate analysis, P=0.184) and a recurrence-free survival in curative HCC (univariate analysis, P=0.086) although HOMA-R was significantly higher in the high insulin group than in the low insulin group (Table III). It is probably due to the small number of cases used to determine the HOMA-R (28 of 140 subjects or 11 of 59 subjects).

Since adiponectin has a potent insulin-sensitizing effect, we determined its value in HCC patients. In contrast to fasting serum insulin, the mean value of total adiponectin apparently increased with the decline of liver function. The HCC stage did not affect the values of total adiponectin. A similar observation has been reported by Tacke *et al* (23), in which they suggest that the elevation of adiponectin in chronic liver disease is due to the decrease of clearance from the serum, and possibly decreased biliary excretion of adiponectin, and that portal hypertension and the development of HCC do not affect the values of adiponectin. In addition to total adiponectin, we measured the levels of HMW, MMW, and LMW adiponectins. These adiponectins increased in direct relation to the decline in the liver function (data not shown), thus suggesting that higher molecular weight adiponectin is also metabolized by the liver. It is surprising that the values of total, HMW, MMW, and LMW adiponectins showed no significant differences between the high insulin group and the low group (Table III). However, Tacke *et al* have already reported a similar observation that the elevated adiponectin in LC patients is not directly involved in insulin sensitivity. Recently, adiponectin is known to possess antitumoral activity. The circulating adiponectin level is inversely associated with an increased risk of breast cancer, endometrial, prostate, gastric, and colorectal cancer (37-41). Furthermore, Miyazaki *et al* reported that adiponectin shows an antitumor effect against HepG2 hepatoma cells through JNK activation and suppression of STAT3 function (42). However, our study showed that total adiponectin has no impact on the prognosis of any stage of HCC. It is unclear why there is a discrepancy between these literature findings and our own. We are now speculating that certain cirrhotic environments such as advanced liver fibrosis, decreased liver function and portal systemic shunting may diminish the anti-tumoral activity of adiponectin against HCC. Further studies are thus needed to clarify this.

Although the present study is retrospective and involves a limited number of participants, this is a first study indicating that fasting hyperinsulinemia is a risk factor associated with a poor prognosis in the early stage of HCC and a high recurrence rate in the curative HCC. We have to validate our findings with a prospective study and also clarify the mechanism by which insulin impacts the clinical course of HCC. However, our study suggests that treatment modalities which lower the level of fasting insulin could improve the prognosis of the early stage of HCC and reduce the recurrence of HCC.

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## Original Article

# Hepatitis C virus kinetics during the first phase of pegylated interferon- $\alpha$ -2b with ribavirin therapy in patients with living donor liver transplantation

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**Aim:** To identify the problems of pegylated interferon (PEG IFN) with ribavirin therapy against hepatitis C virus (HCV) reinfection in living donor liver transplantation (LDLT) patients. HCV kinetics during the PEG IFN with ribavirin therapy were analyzed in LDLT patients, as well as in chronic hepatitis C (CHC) patients.

**Methods:** The study included 80 consecutive HCV infected patients undergoing PEG IFN with ribavirin therapy (64 CHC and 16 LDLT patients) who attended the Nagasaki University Hospital for an initial visit between January 2005 and December 2007.

**Results:** The sustained viral response (VR) rate of the CHC group (80%) was superior to the LDLT group (22%). The viral

disappearance rate of the CHC group was also superior to the LDLT group, regardless of the HCV serotype. The HCV core antigen (cAg) titer under treatment in the LDLT group was more than that of the CHC group from day 0 to week 12. The HCV cAg decrease rate of the LDLT group on the first day of treatment was less than that of the CHC group.

**Conclusion:** The HCV infection of a transplanted liver is more refractory to treatment than a non-transplanted liver. The low reduction HCV cAg rate on day 1 is one of the problems of the combination therapy.

**Key words:** chronic hepatitis C, first phase, hepatitis C virus, interferon, living donor liver transplantation

## INTRODUCTION

HEPATITIS C VIRUS (HCV) infection is widespread throughout the world. Chronic HCV infection leads to cirrhosis and hepatocellular carcinoma. Liver transplantation for HCV-related liver disease has been an option worldwide.<sup>1</sup> Recently, it has been shown that the prognosis for liver transplanted (LT) patients with HCV-related disease deteriorates over time,<sup>2</sup> thus resulting in a poorer outcome than in the non-HCV course.<sup>3</sup> The transplanted liver for HCV-related disease undergoes a rapidly progressive fibrosis and acute graft

failure.<sup>3,4</sup> Consequently, anti-HCV treatment after LT is important for the prognosis. Interferon (IFN) has been recognized as the only treatment method for HCV infection. For the transplanted liver, it is known that IFN treatment improves liver fibrosis or halts the progression.<sup>5</sup> Recently, the combination of pegylated IFN (PEG IFN) with ribavirin was used and produced an excellent result for non-transplanted patients with HCV.<sup>6</sup> However, that was not the case for the HCV re-infected transplanted liver.<sup>7</sup> It is important that the cause of refractory HCV infection in the transplanted liver be more fully clarified. Immunosuppressant therapy, especially with glucocorticoid, has been speculated to be the cause of the refractory nature of the transplanted liver to IFN.<sup>8,9</sup> The cause of this is considered to be that glucocorticoid downregulated the IFN signal transduction in the hepatocytes.<sup>8</sup> The authors recently found that calcineurin inhibitors also inhibited IFN induced STAT-1 phosphorylation and antiviral activity in the HCV

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replicon system.<sup>10</sup> Therefore, the problem of IFN signaling in the hepatocyte induced an IFN refractory condition<sup>11</sup> and decreased the first phase of HCV decline, which was IFN induced HCV decay during the first day of IFN treatment.<sup>12</sup>

In the present study, we attempted to better understand PEG IFN and ribavirin therapy by comparing patients with chronic hepatitis from HCV infection (CHC) with living donor LT (LDLT) patients. When the non-transplanted CHC patients were used as a reference against the HCV reinfected LDLT patients, we expected that the differences in the clinical data in the two groups would help to clarify the problem of IFN refractory HCV infection, and shed light on the analysis of HCV kinetics under IFN and ribavirin treatment, and to elucidate the damaged segment of the IFN induced antiviral mechanism in the LDLT condition.

## PATIENTS AND METHODS

### Patients

THE PRESENT RESEARCH is a prospective study. The study included 80 consecutive HCV-infected patients undergoing PEG IFN with ribavirin combination therapy (64 CHC and 16 LDLT patients) who attended the Nagasaki University Hospital for an initial visit between January 2005 and December 2007. All patients received the targeted dose of 1.5 µg/kg PEG IFN-α-2b (Pegintron; Schering-Pough K.K., Osaka, Japan) once weekly with daily ribavirin (Rebetol; Schering-Pough K.K., Osaka, Japan) for a total dose of 600 mg (bodyweight < 60 kg), 800 mg (60 kg < bodyweight < 80 kg) or 1000 mg (bodyweight > 80 kg) according to bodyweight (BW). The number of patients who were judged to have obtained a curative effect from IFN therapy was 42 in total, and 12 were LDLT patients. If the HCV-RNA had been negative in the patient serum until 12 weeks after the initiation of treatment or positive at 24 weeks, PEG IFN with ribavirin therapy was stopped at week 48. If the HCV-RNA had been negative from weeks 12 to 24, PEG IFN with ribavirin therapy was continued for 24 weeks to a predetermined 48 weeks. CHC patients were diagnosed on the basis of a persistently raised alanine aminotransferase (ALT) level and biopsy proven disease. All LDLT patients, who had undergone liver transplantation for HCV related cirrhosis at Nagasaki University Hospital from June 2002 to May 2007, had the HCV-RNA in their serum at the commencement of PEG IFN with ribavirin treatment. To prevent HCV related hepatitis after liver trans-

plantation, pre-emptive therapy using IFN is the strategy used at the Nagasaki University Hospital. After the recovery of the general condition without ascites and icterus after transplantation, and establishment of the diagnosis using the liver biopsy, PEG IFN with ribavirin therapy was started. The interval between LDLT and IFN treatment was a mean of 281 days (range 16–989 days). Tacrolimus (Astellas, Tokyo, Japan), an immunosuppressive agent, was used together with steroids for all LDLT patients as the induction therapy. When IFN therapy was commenced, tacrolimus was switched to cyclosporin (Novartis, Tokyo, Japan) in 12/16 cases. A percutaneous liver biopsy assisted by ultrasonography was carried out in all cases. Liver histology was evaluated according to the degree of fibrosis and necroinflammatory activity.<sup>13</sup> The extent of fibrosis (staging) was classified as follows: F1 (periportal expansion), F2 (portoportal septa), F3 (portocentral linkage or bridging fibrosis) and F4 (cirrhosis). The necroinflammatory activity (grading) was classified as follows: A1 (mild), A2 (moderate) and A3 (severe). Liver biopsy specimens were fixed in 10% formalin, embedded in paraffin, cut to a thickness of 4 µm, and subjected to hematoxylin-eosin and Azan–Mallory staining.

### Hepatitis C virus kinetics assessment

We compared the HCV viral load in both groups, determined by the HCV core antigen (cAg), at baseline (D0), day 1 (D1), week 1 (W1), week 2 (W2), week 4 (W4), week 8 (W8), week 12 (W12), week 24 (W24) and week 48 (W48). The HCV viral serotype (ST) and HCV cAg were determined using available kits. In this assay, HCV serotypes 1 and 2 correspond to genotypes 1 and 2 of Simmonds' classification,<sup>14</sup> respectively. The HCV cAg correlates with HCV-RNA by quantitative PCR.<sup>15</sup> HCV cAg was measured at the indicated times and HCV-RNA qualitative PCR, the amplicor monitor method, was used after the level was under the detection range of HCV cAg in every month. In the present study, we proposed the calculation of the decreased HCV viral load during PEG IFN with ribavirin treatment and set as follows: a negative HCV cAg was 20 fmol/L and a negative HCV-RNA qualitative PCR was 1 fmol/L.

### Clinical and laboratory measurements

The body mass index (BMI) was calculated as weight (kg) divided by the square of height (m). Subjects fasted overnight before blood samples were obtained. Venous plasma glucose was measured with an automated analyzer, and basal serum insulin was measured using a standard radioimmunoassay. The index of insulin



resistance and  $\beta$ -cell function was calculated using the fasting value of plasma glucose (we excluded the patients with greater than 130 mg/dL), and the serum insulin level according to the homeostasis model assessment (HOMA) method. HOMA-IR, an insulin resistance marker, is calculated as follows: fasting plasma glucose  $\times$  fasting insulin/405. HOMA- $\beta$ , a  $\beta$ -cell function marker, was calculated as follows:  $360 \times$  fasting insulin/(fasting plasma glucose-63).<sup>16</sup> White blood cell, red blood cell, platelet, hemoglobin A1c, ALT, aspartate aminotransferase (AST),  $\gamma$ -GTP, total cholesterol (TC), triglyceride (TG), low density lipoprotein (LDL), high density lipoprotein (HDL), free fatty acid (FFA), and ferritin were determined by standard hematometry and laboratory techniques.

### Statistical analysis

The data were processed on a personal computer and analyzed using StatView 5.0 (SAS Institute, Cary, NC, USA). Differences between groups were analyzed by Mann-Whitney *U*-test and Pearson  $\chi^2$ -test. All data in the text and tables are shown as means, unless otherwise indicated. The statistical analysis of the HCV-RNA disappearance rate was by the Kaplan-Meier method with Wilcoxon assay. Values of  $P < 0.05$  were considered to be statistically significant.

## RESULTS

### Differences of patient characteristics

FIRST, THE PRETREATMENT clinical and laboratory characteristics were compared with All-CHC and All-LDLT patients (Table 1). The BW and BMI in the All-CHC group were higher than that of the All-LDLT group. Therefore, the levels of PEG IFN dose per BW and ribavirin dose per BW were even, but the levels of PEG IFN dose and ribavirin dose in the all LDLT group were lower than in the All-CHC group. The HCV viral load in the all LDLT group was greater than that in the All-CHC group and serotype 1 was the majority in the All-LDLT group. In hematology and laboratory data, the red blood cell count and hemoglobin in the All-LDLT group was lower than that of the All-CHC group, and the FFA level was higher in the All-LDLT group. In the histological examination, fibrosis is more advanced in the All-CHC group than in the All-LDLT group. There was the tendency toward higher levels of fasting plasma glucose and lower levels of HOMA- $\beta$  in the All-LDLT group than in the All-CHC group. Next, we targeted the serotype 1 and a high HCV titer (ST1H group) above 100 KIU/L by

the qualitative PCR method or 300 fmol/L of the cAg assay. These were examined in the same way (Table 2). The ST1H group might have shown the same result as the All group, except the levels of fasting plasma glucose and HOMA- $\beta$  did not differ with ST1H-CHC and ST1H-LDLT. The mean value of fasting plasma glucose (FPG) was higher than the normal range in the LDLT group. The discontinuance rates of treatment were almost equal, 19 cases (29.7%) and 4 cases (25%) in All-CHC and All-LDLT, respectively. The reasons for discontinuance were adverse effects in All-LDLT patients and the refractory nature of viral response in two All-CHC patients.

### The HCV infection in the LDLT group is more obstinate than in the CHC group

The response rate and cure rate of PEG IFN with ribavirin therapy were compared with both groups (Table 2A, All group and B, ST1H group). The HCV response rate to treatment, viral response (VR), was determined by the disappearance of HCV-RNA or by the decline of HCV cAg to less than 1/100 before treatment. The cure rate, sustained viral response rate (SVR), was determined by a negative HCV-RNA by qualitative PCR method at 6 months post-termination of treatment. The VR rate at 8 and 12 weeks, but not at 4 weeks, and the PP-SVR in the LDLT group (Table 3A,B) was worse than that in the CHC group. Non-viral responders, who did not achieve HCV-RNA negativity during the treatment, did not show statistical significance in either SG1H group (Table 3B). As a result, we calculated the prediction of the lack of SVR by non-viral response in the LDLT group. The sensitivity, specificity, positive predictive values and negative predictive value were 1, 0, 0.917 and the acalculia for null viral responders at 24 h, 0.7, 1, 1 and 0.25 at 4 weeks, 0.6, 1, 1 and 0.2 at 8 weeks and 0.6, 1, 1 and 0.2 at 12 weeks, respectively.

The disappearance rate of HCV-RNA was evaluated by the Kaplan-Meier method (Fig. 1 ST1H group). The disappearance rate in the LDLT group was statistically lower than the CHC group. Before 14 weeks after the initiation of treatment, the HCV-RNA disappearance case was not apparent in the ST1H group (Fig. 1).

### The decline of HCV load, especially early phase, is blocked in the LDLT group

For the analysis of viral kinetics, we evaluated the decline of the HCV load and the decline rate after treatment with particular emphasis of the early phase of treatment, including D1-W12. In the ST1H group (Fig. 2), the decreased rate on D1 in the LDLT group was

**Table 1** Difference of characteristics between all chronic hepatitis C cases and all living donor liver transplantation cases

| Characteristics                | All-CHC (n = 64) | All-LDLT (n = 16) | P-value  |
|--------------------------------|------------------|-------------------|----------|
| Age (years)                    | 58 ± 10.8        | 58.8 ± 4.62       | NS       |
| Sex (male : female)            | 36:28            | 7:9               | NS       |
| Height (m)                     | 1.60 ± 0.098     | 1.583 ± 0.010     | NS       |
| Bodyweight (kg)                | 61.0 ± 11.0      | 54.8 ± 8.52       | 0.025    |
| Body mass index                | 23.6 ± 2.94      | 21.8 ± 2.30       | 0.022    |
| PEG IFN dose (µg)              | 80.1 ± 18.7      | 71.9 ± 33.5       | 0.035    |
| PEG IFN/BW                     | 1.31 ± 0.304     | 1.35 ± 0.708      | NS       |
| Ribavirin dose (mg)            | 621.9 ± 151.7    | 525 ± 100         | 0.030    |
| Ribavirin/BW                   | 10.2 ± 2.23      | 9.72 ± 2.04       | NS       |
| Serotype (1:2)                 | 45:17            | 15:1              | 0.081    |
| HCV cAg (fmol/L)               | 5773 ± 5609      | 23144 ± 21059     | 0.001    |
| WBC (/µL)                      | 5006.3 ± 1335    | 5918.8 ± 2439     | NS       |
| RBC (10 <sup>4</sup> /µL)      | 445 ± 41.1       | 350 ± 56.7        | < 0.0001 |
| Hemoglobin (g/dL)              | 13.8 ± 1.06      | 10.9 ± 1.85       | < 0.0001 |
| Platelet (10 <sup>4</sup> /µL) | 16.4 ± 4.48      | 18.5 ± 10.6       | NS       |
| AST (U/L)                      | 62.9 ± 35        | 64.3 ± 37.2       | NS       |
| ALT (U/L)                      | 85 ± 53.0        | 89.9 ± 57.1       | NS       |
| γ-GTP (U/L)                    | 62.1 ± 56.5      | 138.9 ± 129.1     | 0.013    |
| Ferritin (ng/dL)               | 218 ± 216        | 254 ± 259         | NS       |
| TC (mg/dL)                     | 169.8 ± 26.6     | 167.3 ± 38.8      | NS       |
| TG (mg/dL)                     | 105.3 ± 46.8     | 122.8 ± 44.8      | 0.069    |
| HDL (mg/dL)                    | 45.2 ± 11.9      | 46.6 ± 14.9       | NS       |
| LDL (mg/dL)                    | 97.3 ± 24.3      | 88.8 ± 26.7       | NS       |
| FFA (mEq/L)                    | 0.492 ± 0.261    | 0.686 ± 0.299     | 0.019    |
| FPG (mg/dL)                    | 91.9 ± 15.4      | 125.1 ± 56.9      | 0.090    |
| Insulin (mIU/L)                | 9.16 ± 5.1       | 8.34 ± 5.16       | NS       |
| HOMA-IR                        | 2.08 ± 1.22      | 1.75 ± 1.42       | NS       |
| HOMA-β                         | 135.4 ± 86.2     | 89.7 ± 86.9       | 0.075    |
| Fibrosis                       | 1.86 ± 1.18      | 0.875 ± 0.806     | 0.004    |
| Activity                       | 1.03 ± 0.48      | 1.31 ± 0.48       | 0.067    |

Data are shown as the means ± standard deviation and values, with statistical analysis calculated by Mann–Whitney *U*-test for means and Pearson's  $\chi^2$ -test for values.

Normal values in laboratory tests: ALT (IU/L), 5–40; AST (IU/L), 10–40; γ-GTP (IU/L), < 70 in males, < 30 in females; TC (mg/dL), 150–219; TG (mg/dL), 50–149; FFA (mEq/L), 0.14–0.85; LDL (mg/dL), 70–139; HDL (mg/dL), 40–86 in male, 40–96 in female; hemoglobin (g/dL), 13.5–17.6 in male, 11.3–15.2 in female; WBC (/µL), 3900–9800 in males, 3500–9100 in females; RBC (10<sup>4</sup>/µL), 427–570 in males, 376–500 in females; ferritin (mg/dL), 27–320 in males, 3.4–89 in females; platelet (10<sup>4</sup>/µL), 13.1–36.2 in males, 13–36.9 in females; insulin (IU/L), 3.06–16.9; FPG (mg/L), 70–109. HOMA-IR, HOMA-β, and BMI are described in the text.

ALT, alanine aminotransferase; AST, aspartate aminotransferase; CHC, chronic hepatitis C; FFA, free fatty acid; FPG, fasting plasma glucose; HCV cAg, hepatitis C virus core antigen; HDL, high density lipoprotein; HOMA, homeostasis model assessment; LDL, low density lipoprotein; LDLT, living donor liver transplantation; PEG IFN, pegylated interferon; RBC, red blood cell count; TC, total cholesterol; TG, triglyceride; WBC, white blood cell count.

statistically lower than CHC (Fig. 2b) and the viral load of the LDLT group was larger than that in CHC from D0 to W12 (Fig. 2a). The decreased rate at the indicated time without D1 and W12 was not the difference between CHC and LDLT (Fig. 2b). We next analyzed the SG1H-group that matched the pre-treatment HCV cAg titer (Fig. 3). In a similar fashion to Figure 2, the viral load of the matched LDLT group was larger than that of the matched CHC from D1 to W12 (Fig. 3a) and the

decreased rate of the matched LDLT group was lower than that of the matched CHC at D1, W2 and W4 (Fig. 3b).

## DISCUSSION

IN THE PRESENT prospective study, we compared CHC and LDLT patients treated with PEG IFN and ribavirin for HCV infection. BMI, HCV cAg, red blood

**Table 2** Difference of characteristics of serotype 1 and high virus titer between chronic hepatitis C patients and living donor liver transplantation patients

| Characteristics                | ST1H-CHC (n = 42) | ST1H-LDLT (n = 15) | P-value  |
|--------------------------------|-------------------|--------------------|----------|
| Age (years)                    | 58.5 ± 10.8       | 58.8 ± 4.78        | NS       |
| Sex (male : female)            | 22:20             | 6:9                | NS       |
| Height (m)                     | 1.60 ± 0.10       | 1.566 ± 0.081      | NS       |
| Bodyweight (kg)                | 61.8 ± 12.1       | 53.8 ± 7.69        | 0.02     |
| Body mass index                | 24.0 ± 2.78       | 21.9 ± 2.37        | 0.012    |
| PEG IFN dose (µg)              | 81.4 ± 19.5       | 73.3 ± 34.2        | 0.052    |
| PEG IFN/BW                     | 1.33 ± 0.269      | 1.39 ± 0.711       | NS       |
| Ribavirin dose (mg)            | 642.8 ± 150.0     | 520 ± 101.4        | 0.011    |
| Ribavirin/BW                   | 10.5 ± 2.13       | 9.80 ± 2.08        | NS       |
| HCV cAg (fmol/L)               | 6969 ± 5281       | 24674 ± 20856      | 0.003    |
| WBC (/µL)                      | 5019.0 ± 1294     | 6033.8 ± 2479      | NS       |
| RBC (10 <sup>4</sup> /µL)      | 444 ± 40.1        | 351 ± 58.6         | < 0.0001 |
| Hemoglobin (g/dL)              | 13.9 ± 1.10       | 10.8 ± 1.88        | < 0.0001 |
| Platelet (10 <sup>4</sup> /µL) | 16.7 ± 4.68       | 18.9 ± 10.8        | NS       |
| AST (U/L)                      | 62.1 ± 31.6       | 64.2 ± 38.5        | NS       |
| ALT (U/L)                      | 84.5 ± 51.8       | 88.0 ± 58.6        | NS       |
| γ-GTP (U/L)                    | 64.0 ± 61.7       | 113.6 ± 83.1       | 0.036    |
| Ferritin (ng/dL)               | 206 ± 164.8       | 204.5 ± 188.4      | NS       |
| TC (mg/dL)                     | 172.6 ± 25.7      | 165.3 ± 39.2       | NS       |
| TG (mg/dL)                     | 108.2 ± 52.2      | 122.9 ± 46.4       | NS       |
| HDL (mg/dL)                    | 46.5 ± 11.9       | 45.4 ± 14.8        | NS       |
| LDL (mg/dL)                    | 97.7 ± 25.4       | 88.6 ± 27.8        | NS       |
| FFA (mEq/L)                    | 0.514 ± 0.251     | 0.693 ± 0.310      | 0.049    |
| FPG (mg/dL)                    | 92.4 ± 16.4       | 123.7 ± 58.6       | NS       |
| Insulin (mIU/L)                | 9.06 ± 5.5        | 8.34 ± 5.16        | NS       |
| HOMA-IR                        | 2.07 ± 1.31       | 1.86 ± 1.38        | NS       |
| HOMA-b                         | 128.0 ± 76.2      | 95.7 ± 86.5        | NS       |
| Fibrosis                       | 1.92 ± 1.19       | 0.933 ± 0.799      | 0.008    |
| Activity                       | 1.08 ± 0.474      | 1.33 ± 0.488       | 0.098    |

Data are shown as the means ± standard deviation and values, with statistical analysis calculated by Mann–Whitney *U*-test for means and Pearson's  $\chi^2$ -test for values.

Normal values in laboratory tests are same as in Table 1.

ALT, alanine aminotransferase; AST, aspartate aminotransferase; CHC, chronic hepatitis C; FFA, free fatty acid; FPG, fasting plasma glucose; HCV cAg, hepatitis C virus core antigen; HDL, high density lipoprotein; HOMA, homeostasis model assessment; LDL, low density lipoprotein; LDLT, living donor liver transplantation; PEG IFN, pegylated interferon; RBC, red blood cell count; TC, total cholesterol; TG, triglyceride; WBC, white blood cell count.

cell,  $\gamma$ -GTP, FFA and liver fibrosis in the pretreatment clinical characteristics were different in both groups (Tables 1,2). The VR rate of the CHC group was superior to that of the LDLT group, and the SVR by per-protocol analysis was also similar in result to the VR (Table 3). The viral disappearance rate of the CHC group was superior to the LDLT group, regardless of the HCV serotype (Fig. 1). The HCV cAg titer under the treatment in the LDLT group was more than that of the CHC group from D0 to W12 (Figs 2a,3a) and the HCV cAg decrease rate of the LDLT group at the D1 was less than that of the CHC group (Figs 2b,3b). We showed that the reinfected

HCV to the graft liver was more refractory than the non-transplanted CHC. The PEG IFN and ribavirin dose per BW was an equal dose in both groups. However, it was difficult to determine the pretreatment predictive factors for the LDLT cases, because only one case showed SVR in the LDLT group. Thus, we considered that the difference of the pretreatment clinical characteristics in both groups might be related to the refractory HCV infection.

The pretreated HCV cAg titer is known to be the principal factor for IFN resistance. For CHC and LDLT patients, a high HCV-RNA titer in the pretreatment sera

**Table 3** Result of pegylated interferon- $\alpha$ -2b plus ribavirin therapy

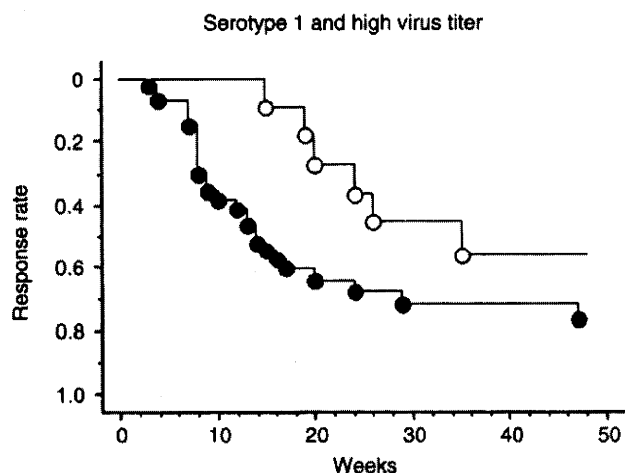
| A. All cases                             |             |            |         |
|--|-------------|------------|---------|
| Term                                     | All-CHC     | All-LDLT   | P-value |
| Viral response 4 weeks                   | 40/60 (67%) | 5/12 (42%) | NS      |
| Viral response 8 weeks                   | 47/55 (85%) | 6/12 (50%) | 0.011   |
| Viral response 12 weeks                  | 43/48 (90%) | 6/12 (50%) | 0.003   |
| Sustained viral response: ITT            | 20/42 (45%) | 2/12 (20%) | 0.054   |
| Sustained viral response: PP             | 20/28 (80%) | 2/9 (22%)  | 0.008   |
| B. Serotype 1 and high virus titer cases |             |            |         |
| Term                                     | ST1H-CHC    | ST1H-LDLT  | P-value |
| Viral response 4 weeks                   | 24/40 (67%) | 5/11 (45%) | NS      |
| Viral response 8 weeks                   | 30/36 (83%) | 5/11 (45%) | 0.012   |
| Viral response 12 weeks                  | 25/29 (86%) | 5/11 (45%) | 0.008   |
| Sustained viral response: ITT            | 8/27 (30%)  | 1/11 (8%)  | NS      |
| Sustained viral response: PP             | 8/15 (53%)  | 1/9 (11%)  | 0.029   |
| Non-virological response: ITT            | 11/27 (41%) | 5/11 (45%) | NS      |
| Non-virological response: PP             | 4/15 (27%)  | 4/9 (44%)  | NS      |

Data are shown as relevant numbers/target case numbers (percentage of relevant numbers) with statistical analysis using Pearson's  $\chi^2$ -test for numbers.

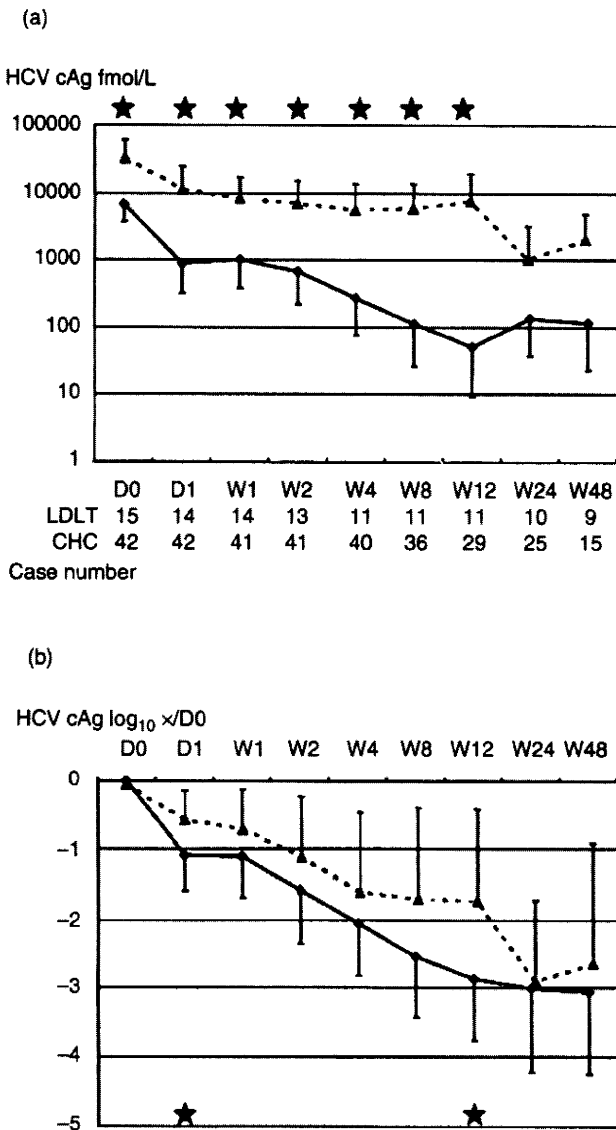
CHC, chronic hepatitis C; ITT, intention to treatment analysis; LDLT, living donor liver transplantation; PP, per-protocol analysis.

is associated with non-responder status for IFN treatment.<sup>7,17</sup> In the LDLT condition, the HCV-RNA titer was rapidly increased after immediately decreasing at transplant and the viral load after several weeks post-LDLT exceeded the value of pre-LDLT.<sup>18</sup> The HCV-RNA titer increased rapidly in patients receiving corticosteroids as part of the immunosuppressant regimen.<sup>18,19</sup> We have speculated that the massive amount of HCV, caused by immunosuppressant therapy after the LDLT, was part of the reason for the IFN refractory status. However, comparisons with the pretreated HCV cAg matched groups (Fig. 3) showed the existence of an important factor other than the pretreatment viral load. It will, therefore, be necessary to analyze this problem by evaluating many factors, for example immunosuppressants<sup>10</sup> and regeneration, in the future.

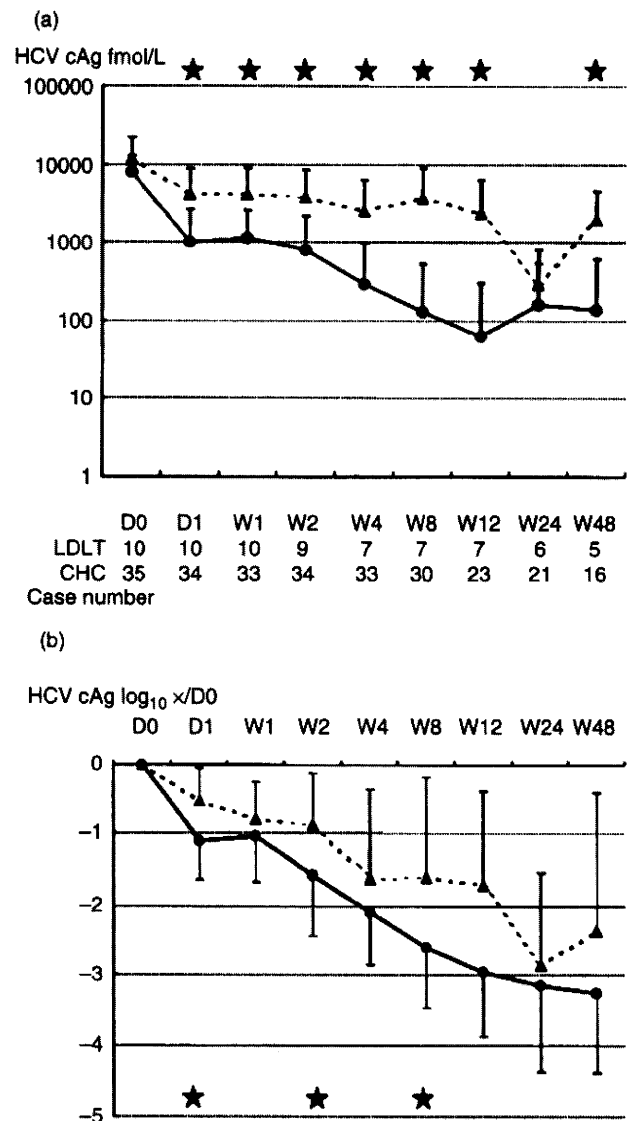
A high level of  $\gamma$ -GTP was also known to be an important factor for IFN treatment.<sup>7,17</sup> Usually, high levels of  $\gamma$ -GTP and FFA have been linked to insulin resistance.<sup>20,21</sup> Therefore, insulin resistance in the liver is assumed in the condition of IFN resistance. However, the LDLT group had the normal range of HOMA-IR,<sup>16</sup> which was lower than that of the CHC group (Tables 1,2). The HCV infection after liver transplantation is associated with insulin resistance.<sup>22</sup> Immunosuppressants, especially corticosteroids, induced insulin resistance.<sup>23</sup> In the present study, the LDLT group had a disturbance of insulin secretion



**Figure 1** The difference in the hepatitis C ribonucleic acid (HCV-RNA) disappearance rate between the chronic hepatitis C (CHC) group and living donated liver transplantation (LDLT) group during 48 weeks of treatment. HCV-RNA was evaluated by the qualitative PCR method. The disappearance rate was calculated as follows: serum HCV-RNA disappearance case number/all cases in indicated time. The statistical analysis was carried out using the Kaplan–Meier method with the Wilcoxon assay. ST1H group was plotted as the HCV-RNA disappearance line between the white circle of the LDLT group and the black circle of the CHC group. In all cases and the ST1H group, the disappearance rate was statistically significant between the CHC group and the LDLT group ( $P < 0.05$ ).



**Figure 2** Comparison of viral kinetics between the SG1H-chronic hepatitis C (CHC) group and the SG1H-living donor liver transplantation (LDLT) group during the 48 weeks of treatment. (a) The hepatitis C virus core antigen (HCV cAg) load and (b) reduction rates were plotted by a straight line (SG1H-CHC group), and dotted line (SG1H-LDLT group). The error bar represented the standard deviation. On the y-axis, D0 is pretreatment, D1 and WX is time post-treatment day 1 and week X, respectively. The reduction rate was calculated as follows:  $\log_{10}$ HCV cAg load in indicated time/in D0. HCV cAg titer at the indicated time between SG1H-CHC and SG1H-LDLT were compared. The asterisk mark indicates a significant difference,  $P < 0.05$ , calculated by Mann-Whitney *U*-test.



**Figure 3** Comparison of viral kinetics between matched pre-treatment hepatitis C virus core antigen (HCV cAg) ST1H-chronic hepatitis C (CHC) group and ST1H- living donor liver transplantation (LDLT) group during 48 weeks of treatment. (a) HCV cAg load and (b) reduction rate were plotted by a straight line (matched SG1H-CHC group) and dotted line (matched SG1H-LDLT group). The error bar represents the standard deviation. The asterisk mark is the significant difference,  $P < 0.05$ , calculated by Mann-Whitney *U*-test.

rather than insulin resistance and high levels of FPG might be caused by the disturbance of insulin secretion. Therefore, further study is necessary to clarify the relationship between the glucose metabolism and the IFN resistance in LDLT patients. The levels of  $\gamma$ -GTP rise at cholestatic conditions. It was reported that the presence

of a cholestatic profile is associated with an adverse response to IFN treatment in LT.<sup>7</sup> A cholestatic profile provoked the TH2-like lymphocyte response.<sup>19</sup> The authors have previously reported that IL-10, representative of TH2 cytokine, inhibits IFN signaling through an inducible suppressor of cytokine signaling.<sup>24</sup> The high levels of FFA were induced by a catabolic state, such as cirrhosis, and were not fully recovered after LDLT. As a result, the levels of FFA reflected a continuous catabolic state at the beginning of IFN treatment. FFA can induce oxidative stress in various cells,<sup>25,26</sup> and inhibit the IFN induced antiviral gene induction through the inactivation of Jak-1 and Tyk-2.<sup>27</sup> Therefore, we are speculating that high levels of  $\gamma$ -GTP and FFA in the LDLT group have the ability to inhibit IFN signaling as much as in the CHC patients.

We are paying attention to the viral decline of D1/D0 (Figs 2b,3b). The decreased rate of D1 is named as the first phase of HCV decline and is the predictor of SVR.<sup>28,29</sup> The first phase influenced the second phase, which is the decline of HCV after D2.<sup>28</sup> The IFN induced antiviral gene products were considered to be very important for antiviral activity.<sup>11</sup> The expressions of the IFN stimulating genes (ISG) were associated with the early phase of the decline<sup>11</sup> and it was reported that the lack of ISG caused early liver fibrosis in the LT patients with HCV.<sup>30</sup> In the LDLT group, the reduced HCV cAg decreased the rate of D1 and this might be part of the cause of being refractory to IFN. We speculate that an IFN signaling disturbance, related to high levels of  $\gamma$ -GTP and FFA, might have triggered the adverse effect to the HCV cAg decreased rate of D1.

In summary, it became clear that the viral response and SVR is worse in the LDLT group. The first phase of viral decay, the decreased rate of D1/D0, also declined in the LDLT group. High levels of  $\gamma$ -GTP and FFA in the pretreatment sera might also be related to IFN-signaling damage in hepatocytes. At the initiation of pre-emptive therapy, HCV had also been increasing in the graft liver and the catabolic status of energy did not recover for the relatively small size of the graft liver. When beginning treatment for an HCV infection after LT, we should carefully take into account the timing of IFN initiation, in addition to the types of immunosuppressants used.

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## CLINICAL STUDIES

**Study of liver-targeted regulatory T cells in hepatitis B and C virus in chronically infected patients**Hisamitsu Miyaaki<sup>1\*</sup>, Huijuan Zhou<sup>1,2\*</sup>, Tatsuki Ichikawa<sup>1</sup>, Kazuhiko Nakao<sup>1</sup>, Hidetaka Shibata<sup>1</sup>, Shigeyuki Takeshita<sup>1</sup>, Motohisa Akiyama<sup>1</sup>, Eisuke Ozawa<sup>1</sup>, Satoshi Miuma<sup>1</sup> and Katsumi Eguchi<sup>1</sup>

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**Keywords**

Foxp3 – hepatitis B virus – hepatitis C virus – regulatory T cell

**Abbreviations**

HBV, hepatitis B virus; HCV, hepatitis C virus; CHB, chronic hepatitis B; CHC, chronic hepatitis C; GITR, glucocorticoid-induced TNF receptor-related protein; Foxp3, forkhead box P3

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Hepatitis B virus (HBV) and hepatitis C virus (HCV) are global health problems and both cause chronic hepatitis, cirrhosis and hepatocellular carcinoma (1, 2). In some cases, these viruses lead to a chronic infection in the liver, but the mechanisms by which these hepatitis viruses are able to evade the immune system are still only poorly understood.

Recent studies have focused on regulatory T cells (Tregs). Tregs are engaged in the maintenance of self-tolerance by suppressing the activation and expansion of self-reactive lymphocytes (3–5). This suppressing function may lead to chronic inflammation and/or autoimmunity (6–12). Most of these Treg markers, including CD25, CTLA-4 and glucocorticoid-induced TNF receptor-related protein (GITR), do not accurately represent CD4<sup>+</sup> T cells with regulatory activity, and they overlap with activated T cells, which do not necessarily possess a regulatory activity (13, 14). Currently, the best indicator of the Tregs function is thought to be the intracellular expression of forkhead box P3 (Foxp3), which is also crucial for Treg development (15). Previous reports have indicated that Tregs play a role in viral persistence by suppressing the virus-specific T cell responses in a cell-to-cell contact manner (16, 17).

\*Co-first authors.

**Abstract**

**Introduction:** Regulatory T cells (Tregs) play a critical role in chronic viral infections. The role of Tregs in chronic hepatitis B (CHB) and chronic hepatitis C (CHC) is unknown. This study examined the distribution and frequency of forkhead box p3<sup>+</sup> (Foxp3<sup>+</sup>) Tregs in the liver tissue and compared the clinicopathological characteristics of CHB and CHC patients. **Methods:** Liver needle biopsies were obtained from 26 patients who were hepatitis B surface antigen positive and 27 patients who were hepatitis C virus antibody positive. **Results:** The ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells was similar in HBV and in HCV cases. In HBV cases, the variables that were positively associated with the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells included the serum alanine aminotransferase level ( $R = 0.402$ ,  $P = 0.025$ ) and the ratio of CD8<sup>+</sup> T cell plus CD56<sup>+</sup> NK cell against CD4<sup>+</sup> T cell ( $R = 0.53$ ,  $P = 0.005$ ). The ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells increased more in the severe activity group than in the mild activity group ( $P = 0.04$ ). In HCV cases, the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells increased significantly in terms of the genotype2 ( $P = 0.0002$ ) and male gender ( $P = 0.04$ ). In addition, the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells showed a negative correlation with the ratio of CD8<sup>+</sup> T cell plus CD56<sup>+</sup> NK cell against CD4<sup>+</sup> T cell ( $R = -0.508$ ,  $P = 0.005$ ) and HCV viral load ( $R = -0.482$ ,  $P = 0.001$ ). **Conclusions:** Liver-targeted regulatory T cells present similarly in CHB and CHC, but their relationship with the effector cell population, the inflammation grade or the viral load is different between CHB and CHC.

A more detailed analysis of this cell reaction is necessary not only in the peripheral blood but also in the target organ tissue. The liver is the target organ of both HBV and HCV. This study examined the distribution and frequency of Tregs in the liver and compared the clinicopathological characteristics of chronic hepatitis B (CHB) and chronic hepatitis C (CHC) patients.

**Patients and methods**

Liver needle biopsies were obtained from 26 patients who were hepatitis B surface antigen positive (mean age:  $33.3 \pm 10.4$ , male: female = 23:3) and 27 patients who were hepatitis C positive virus antibody positive (mean age:  $57.8 \pm 10.6$ , male: female = 15:12) at Nagasaki Universities and associated hospitals. The clinical and biological parameters, including HBV viral load, of all the patients were assessed using the HBV transcription-mediated amplification method or HCV virus load by the quantitative RT-PCR method. None of the HBV cases had been administered any antiviral treatment, while 10 of 27 HCV cases received antiviral treatment.

The clinical data of the patients are summarized in Table 1. Liver biopsy tissue specimens were taken by a needle puncture for diagnostic purposes. The diagnosis of each case was



**Table 1.** Clinical data of the patients

|  | HBV 26 cases | HCV 27 cases  |
|--|--------------|---------------|
| Mean age (years old)                         | 33.6 ± 10.4  | 57.8 ± 10.6   |
| Male:female                                  | 23:3         | 15:12         |
| ALT (IU/L)                                   | 310 ± 472    | 91 ± 54       |
| AST (IU/L)                                   | 208 ± 333    | 67 ± 36       |
| Plt (× 10 <sup>4</sup> /μl)                  | 16.6 ± 5.6   | 16.9 ± 4.7    |
| Inflammation grade (mild:severe)             | 19:7         | 22:5          |
| Fibrosis stage (mild:severe)                 | 15:11        | 14:13         |
| HBV viral load LEG/ml (TMA)                  | 6.8 ± 1.8    |               |
| HCV genotype (1:2)                           |              | 21:6          |
| Antiviral treatment (naïve:former treatment) |              | 17:10         |
| HCV viral load KIU/ml (RT-PCR)               |              | 1637 (5–5000) |

ALT, alanine aminotransferase; AST, aspartate aminotransferase; HBV, hepatitis B virus; HCV, hepatitis C virus; RT-PCR, reverse transcriptase polymerase chain reaction; TMA, transcription-mediated amplification.

independently confirmed histologically by liver pathologists according to the Japanese chronic hepatitis classification (New Inuyama classification). Mild activity was defined as A0 or A1, and severe activity was defined as A2 or A3 by the Inuyama classification. Mild fibrosis was defined as F0 or F1, and severe fibrosis was defined as F2, F3 or F4 by the Inuyama classification. All tissues were fixed in 10% neutral-buffered formalin and were then embedded in paraffin, and 4-μm-thick serial sections were cut from each paraffin block.

T cells were examined immunohistochemically with the anti-CD3 antibody (Novocastra, Newcastle, UK), regulatory T cells were examined with the anti-Foxp3 antibody (eBioscience, San Diego, CA, USA), CD4<sup>+</sup> T cells were examined with the anti-CD4 antibody (Novocastra), CD8<sup>+</sup> T cells were examined with the anti-CD8 antibody (Novocastra) and NK cells were examined with the anti-CD56 antibody (Novocastra). The number of Foxp3<sup>+</sup>, CD4<sup>+</sup>, CD8<sup>+</sup>, CD56<sup>+</sup> and CD3<sup>+</sup> cells contained within three portal tracts selected in each specimen were counted at a magnification of × 200. To correct for differences in the size of the portal tracts, the average proportion of Foxp3<sup>+</sup> Tregs among the total number of CD3<sup>+</sup> T cells and the average ratios of CD8<sup>+</sup> T cell plus CD56<sup>+</sup> NK cells against CD4<sup>+</sup> T cells were determined.

#### Statistical analysis

The SPSS statistical software program (SPSS, Chicago, IL, USA) was used to assess any correlations among multiple variables. Differences between the groups were analysed using the Mann–Whitney *U* test. Correlations between the groups were determined by the Spearman analysis. A *P* value of < 0.05 was considered to be statistically significant.

#### Results

##### The population of Foxp3<sup>+</sup> regulatory T cells in the liver

In HBV and HCV cases, both CD3<sup>+</sup> and Foxp3<sup>+</sup> lymphocytes were mainly seen in the portal areas (Fig. 1). In HBV cases, the average ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells in the portal tract was 0.09 ± 0.04, and the average ratio of CD8<sup>+</sup> T cell plus CD56<sup>+</sup> NK cell against CD4<sup>+</sup> T cell was 0.93 ± 0.63, while in

HCV cases, the average ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells was 0.09 ± 0.05, and the average ratio of CD8<sup>+</sup> T cell plus CD56<sup>+</sup> NK cell against CD4<sup>+</sup> T cell was 0.95 ± 0.51.

##### Association between the laboratory data and the frequency of Foxp3<sup>+</sup> regulatory T cells in the liver

In HBV and HCV cases, there was no significant correlation between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells and age. In HBV cases, the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells showed no significant difference in either gender (Table 2). On the other hand, in HCV cases, the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells was significantly increased in males in comparison with females (male:female = 0.11 ± 0.05:0.07 ± 0.03, *P* = 0.04; Table 3). In HBV cases, there was a significant correlation between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells and the serum ALT level (*R* = 0.481, *P* = 0.026; Fig. 2, Table 2). Similarly, there was a significant correlation between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cell and AST level (*R* = 0.402, *P* = 0.025; Table 2). On the other hand, in HCV cases, there was no significant correlation between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells and either serum AST or ALT level (*R* = −0.177, *P* = 0.38, *R* = −0.127, *P* = 0.53; Fig. 2, Table 3).

Both in HBV and in HCV cases, there was no significant correlation between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells and the platelet count (*R* = −0.10, *P* = 0.91, *R* = −0.18, *P* = 0.37, respectively; Tables 2 and 3).

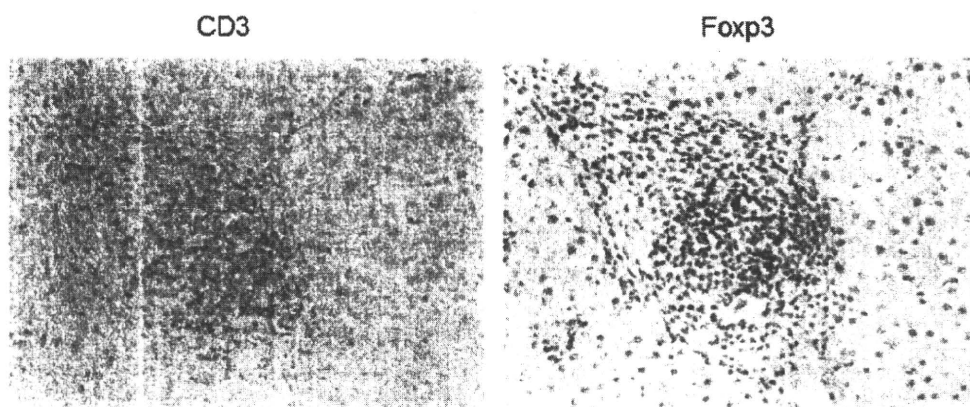
In HBV cases, there was no significant correlation between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells and HBV DNA viral load (*R* = −0.314, *P* = 0.19; Table 2). In HCV cases, there was a significant inverse correlation between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells and HCV viral load (*R* = −0.487, *P* = 0.01; Fig. 3a, Table 3). In addition, the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells was significantly increased in the genotype2 group in comparison with the genotype1 group (genotype1:genotype2 = 0.07 ± 0.03:0.11 ± 0.05, *P* = 0.0002; Fig. 3b, Table 3). In HCV cases, the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells showed no significant difference between the naïve group and the former antiviral treatment group (0.08 ± 0.05:0.10 ± 0.03, *P* = 0.40; Table 3).

##### Association between the histological findings and the frequency of Foxp3<sup>+</sup> regulatory T cells in the liver

In HBV cases, the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells was significantly increased in the severe activity group in comparison with the mild activity group (mild:severe = 0.08 ± 0.03:0.11 ± 0.04, *P* = 0.04; Fig. 4, Table 2). In HCV cases, there was no significant difference in either activity group (mild:severe = 0.100:0.05:0.06 ± 0.03, *P* = 0.09; Fig. 4, Table 3). There were no significant differences among the fibrosis groups in the HBV and HCV cases (mild:severe = 0.09 ± 0.04:0.10 ± 0.03, *P* = 0.58, mild:severe = 0.09 ± 0.05:0.10 ± 0.05, *P* = 0.65, respectively; Tables 2 and 3).

##### Association between the ratio of Foxp3<sup>+</sup> regulatory T cells in CD3<sup>+</sup> T cells and the ratio of CD8<sup>+</sup> T cells plus CD56<sup>+</sup> NK cells against CD4<sup>+</sup> T cells in the liver

In HBV cases, a positive correlation was observed between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells and the ratio of CD8<sup>+</sup> T cells plus CD56<sup>+</sup> NK cells in CD4<sup>+</sup> T cells (*R* = 0.53, *P* = 0.005; Table 2). In contrast, in the HCV cases, a negative correlation was observed between them (*R* = −0.508, *P* = 0.005; Table 3).



**Fig. 1.** Immunostaining of CD3<sup>+</sup> lymphocytes and Foxp3<sup>+</sup> Tregs in representative liver sections of patients with CHB and CHC.

**Table 2.** Comparison between Foxp3/CD3 and each parameter in Chronic hepatitis B

|                       |                         | <i>P</i> value* |
|-----------------------|-------------------------|-----------------|
| Sex (male:female)     | 0.09 ± 0.04:0.09 ± 0.03 | 0.97            |
| Age                   | <i>R</i> = -0.118       | 0.57            |
| AST                   | <i>R</i> = 0.402        | 0.025           |
| ALT                   | <i>R</i> = 0.481        | 0.026           |
| Plt                   | <i>R</i> = -0.10        | 0.90            |
| Grading (mild:severe) | 0.08 ± 0.03:0.11 ± 0.04 | 0.04            |
| Staging (mild:severe) | 0.09 ± 0.04:0.10 ± 0.03 | 0.58            |
| Viral load            | <i>R</i> = 0.314        | 0.19            |
| (CD8+CD56)/CD4†       | <i>R</i> = 0.53         | 0.005           |

\*Differences between the groups were analysed using the Mann-Whitney *U* test. Correlations between the groups were determined by the Spearman analysis.

†The ratio of CD8<sup>+</sup> T cell plus CD56<sup>+</sup> NK cell against CD4<sup>+</sup> T cell.

ALT, alanine aminotransferase; AST, aspartate aminotransferase; NK, natural killer.

**Table 3.** Comparison between Foxp3/CD3 and each parameter in Chronic hepatitis C

|  |                         | <i>P</i> value* |
|--|-------------------------|-----------------|
| Sex (male:female)                            | 0.11 ± 0.05:0.07 ± 0.03 | 0.04            |
| Age  | <i>R</i> = 0.255        | 0.20            |
| AST  | <i>R</i> = -0.177       | 0.38            |
| ALT  | <i>R</i> = -0.127       | 0.53            |
| Plt  | <i>R</i> = -0.18        | 0.37            |
| Grading (mild:severe)                        | 0.10 ± 0.05:0.06 ± 0.03 | 0.09            |
| Staging (mild:severe)                        | 0.09 ± 0.05:0.10 ± 0.05 | 0.65            |
| Genotype (1:2)                               | 0.07 ± 0.03:0.11 ± 0.05 | 0.0002          |
| Antiviral treatment (naïve:former treatment) | 0.08 ± 0.05:0.10 ± 0.03 | 0.40            |
| Viral load                                   | <i>R</i> = -0.482       | 0.01            |
| (CD8+CD56)/CD4†                              | <i>R</i> = -0.508       | 0.005           |

\*Differences between the groups were analysed using the Mann-Whitney *U* test. Correlations between the groups were determined by the Spearman analysis.

†The ratio of CD8<sup>+</sup> T cell plus CD56<sup>+</sup> NK cell against CD4<sup>+</sup> T cell.

ALT, alanine aminotransferase; AST, aspartate aminotransferase.

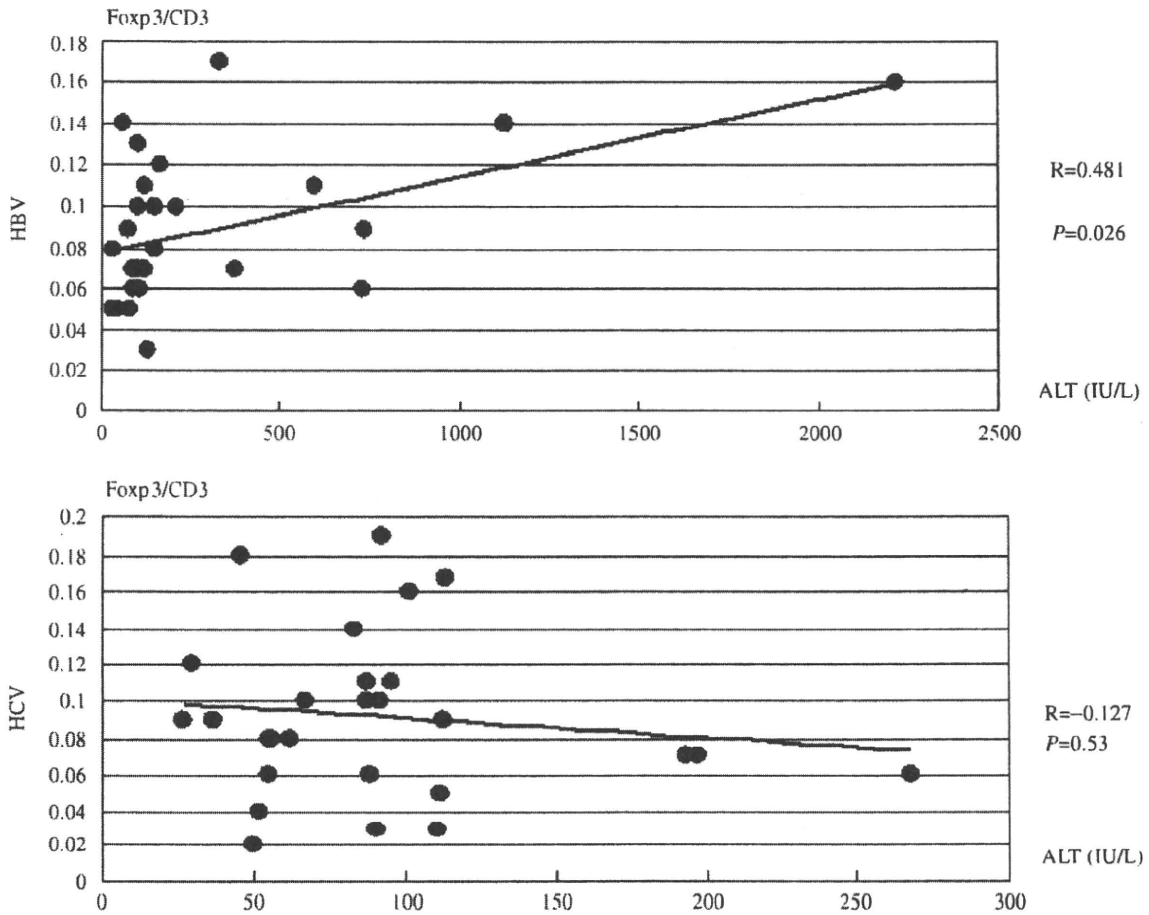
## Discussion

Both HBV and HCV infect the liver and cause acute and chronic liver disease (1, 2). However, the mechanism of persistent infection of these viruses is not clear. Some studies have reported that CD4<sup>+</sup> CD25<sup>high</sup> regulatory T cells (Tregs) play a critical role in persistent virus infection (18–20). Recently, forkhead box P3 (Foxp3) was detected as a specific marker of Tregs (12). The present study examined the distribution of Foxp3<sup>+</sup> Tregs in the liver and clarified the role of Tregs in chronic viral hepatitis.

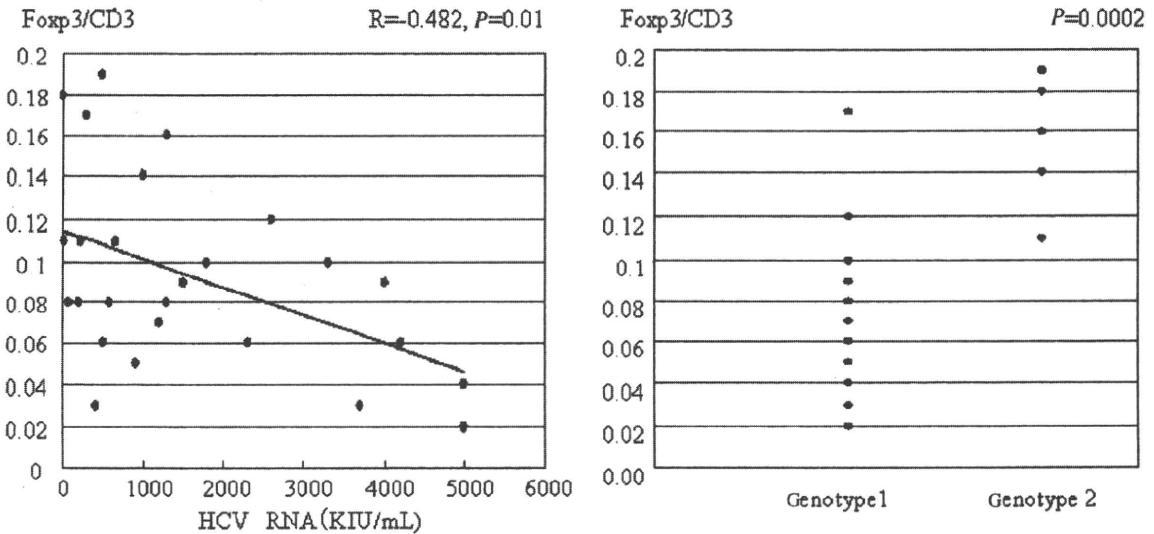
A previous study showed that the normal human liver does not display a significant population of Tregs. This may mean that the tolerance of the liver is not primarily maintained by Tregs. In this study, Foxp3<sup>+</sup> regulatory T lymphocyte populations in the liver constitute 9% of CD3<sup>+</sup> T cell in both HBV and HCV cases. Other studies have shown that Tregs accumulate and expand locally at the site of infection where they exert their suppressive activity (21–23). This suggests that Tregs may be a key factor in the pathogenesis of chronic viral hepatitis.

In HBV cases, the variables that were significantly associated with the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells included the transaminase level (AST and ALT) and pathological grading. These factors are characteristics of the degree of inflammation in the liver. Indeed, the ratio of CD8/CD3 correlated significantly with the transaminase level (AST and ALT) in HBV cases (*r* = 0.428, *P* = 0.029, *r* = 0.495, *P* = 0.010; data not shown). These results indicated that, when severe hepatitis occurred in CHB, the number of Foxp3<sup>+</sup> Treg increases. The ratio of CD8<sup>+</sup> T cell + CD56<sup>+</sup> NK cell against CD4<sup>+</sup> cells significantly correlated with the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells. These observations suggest that Tregs may migrate into the liver during the necroinflammatory reactions induced by the virus-specific effector T cells. If Treg acts too late, the unlimited effector T cells may lead to excessive destruction and the death of hepatocyte. Therefore, it is possible that Tregs suppress the excessive activity of effector T cells in severe hepatitis by HBV.

In contrast, in HCV cases, no correlation was found between alanine aminotransferase or pathological grading and the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells. Moreover, the ratio of CD8<sup>+</sup> T cell + CD56<sup>+</sup> NK cell against CD4<sup>+</sup> cells showed a significant inverse correlation with the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells. In daily medical examination, the patients rarely experienced acute exacerbation of chronic hepatitis in HCV cases.



**Fig. 2.** Comparison between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cell and serum alanine aminotransferase level in chronic hepatitis B (CHB) and in chronic hepatitis C. A significantly positive correlation was observed between the frequency of Foxp3<sup>+</sup> Treg and the serum alanine aminotransferase level in chronic hepatitis B by the Spearman analysis ( $R = 0.402$ ,  $P = 0.025$ ).



**Fig. 3.** (a) Comparison between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells and HCV RNA (KIU/ml). A significantly negative correlation was found between the frequency of Foxp3<sup>+</sup> Tregs and HCV core protein by the Spearman analysis ( $R = -0.482$ ,  $P = 0.01$ ), (b) Comparison between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells and HCV genotype. A significant difference was observed in the genotype by the Mann–Whitney  $U$  test ( $P = 0.0002$ ).

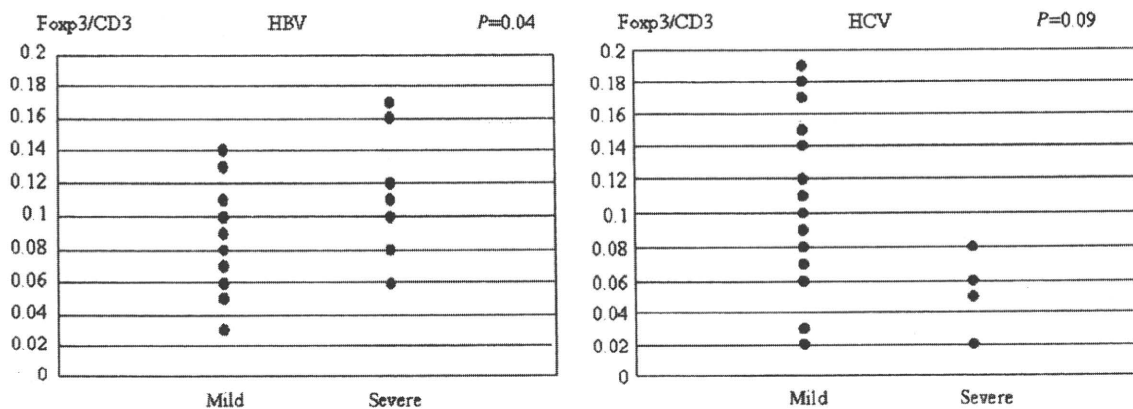


Fig. 4. Comparison between the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells and the histological fibrosis grade in chronic hepatitis B and in chronic hepatitis C. A significant difference was observed in chronic hepatitis B using the Mann–Whitney *U* test ( $P=0.04$ ).

In this study, the transaminase level was > 1000 IU/ml in some of the HBV cases, but in all HCV cases, it was < 300 IU/ml. These results suggest that Treg could constantly suppress the flare of inflammation induced by HCV.

The prevalence of regulatory T cells in the peripheral blood of patients with autoimmune encephalitis or systemic lupus erythematosus (24) is higher in males than in females. Similarly, in the present study, the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells in HCV cases in the liver was significantly higher in females than in males. Sex hormones may act directly on Tregs to alter their functions because immune cells express oestrogen and androgen receptors (25, 26). In the HCV cases, most of the female patients were over 50 years old (76%) and in a postmenopausal state. These findings suggested that oestrogen deficiency is therefore associated with the Tregs in the liver.

Viral genotype, viral load and gender have been shown to be key factors associated with HCV clearance by interferon therapy (27, 28, 29). The mechanism responsible for this difference in interferon treatment outcome is unknown. In this study, in HCV cases, the variables that were significantly associated with the ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cell were genotype2 and male gender. These results suggested that regulatory T cells in the liver may be related to the clearance of HCV by interferon therapy and may be a useful indicator for interferon therapy. The ratio of Foxp3<sup>+</sup> Tregs in CD3<sup>+</sup> T cells showed a significant inverse correlation with the HCV viral load. Tregs were induced by antigen-presenting cells such as dendritic cells (30). Therefore, a decreasing number of Tregs may indicate an impairment of antigen-presenting cells; conversely, the presence of Tregs may represent a normal function of antigen-presenting cells. Taken together, it is possible that, in CHB, regulatory T cells suppress the excessive activity of effector T cells during acute exacerbation, and that, in CHC, the presence of regulatory T cells may be a marker of the normal response of the host immunosystem against HCV.

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