# Insulin resistance is a cause of steatosis and fibrosis progression in chronic hepatitis C

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**Background:** Insulin resistance is a frequent feature of chronic hepatitis C. Whether insulin resistance could be the cause or consequence of steatosis and fibrosis is unknown. The ability of HCV genotype 3 to promote steatosis by itself provides an unique opportunity to answer this question.

**Aims:** The aim of the present study was to assess the relationships between insulin resistance, steatosis, and fibrosis according to genotype in 141 non-diabetic patients with biopsy proven non-cirrhotic chronic hepatitis C.

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Revised version received 1 October 2004 Accepted for publication 26 October 2004 **Methods:** All patients had fasting serum glycaemia and insulinaemia measurements. Insulin resistance was evaluated using the homeostasis model assessment (HOMA IR) method. Liver steatosis was determined according to hepatitis C virus genotype (1 or 3). Logistic regression and multivariate regression analysis were used to identify variables independently associated with insulin resistance, fatty liver, and fibrosis. **Results:** Although steatosis and fibrosis were more severe in genotype 3 patients, median HOMA IR was significantly higher in patients with genotype 1 related steatosis than in those with genotype 3 related steatosis (2.1 v 1; p = 0.001). Independent risk factors for steatosis were insulin resistance in genotype 1 patients (p = 0.001) and viral load in genotype 3 patients (p = 0.003). Among genotype 1 patients, independent parameters associated with insulin resistance were age (p = 0.04) and steatosis (p = 0.004). Steatosis was associated with more severe fibrosis whatever the genotype (p = 0.002). Among genotype 1 patients, although there was a significant relationship between circulating insulin level and fibrosis stage (p = 0.006), only steatosis and inflammatory score were independently associated with fibrosis. **Conclusion:** This study shows that insulin resistance is the cause rather than the consequence of steatosis

and fibrosis in genotype 1 patients and that increased circulating insulin is a risk factor for fibrosis through insulin resistance induced steatosis.

iver steatosis is common in chronic hepatitis C virus infection (HCV).<sup>1-3</sup> Evidence supporting a detrimental role for steatosis in liver fibrosis progression in chronic hepatitis C has been provided recently. Several studies have shown that either fat accumulation in the liver or worsening are strong and independent predictors of the severity and progression of fibrosis in chronic hepatitis C infection, irrespective of the viral genotype involved.4-10 Two main mechanisms have been proposed to account for the high prevalence of steatosis in chronic hepatitis C. Firstly, in patients infected with genotype 3, the degree of steatosis is correlated with the level of viral load,4 11 suggesting that HCV could alter fatty acid metabolism and/or export in hepatocytes.<sup>12 13</sup> Secondly, type 2 diabetes and more generally insulin resistance is highly frequent in chronic HCV infection, as established by several recent epidemiological studies.14-21 Whether or not insulin resistance is a player in the development of the histological lesions of chronic hepatitis C—as in alcoholic or non-alcoholic steatohepatitis—or simply the consequence of viral infection and liver injury is debated. Recently, Shintani et al have shown that insulin resistance preceded the occurrence of steatosis in transgenic mice expressing HCV core protein, suggesting that insulin resistance is not a consequence of hepatic steatosis in these mice.<sup>22</sup> This issue is of obvious practical importance because appropriate treatment should be based on the pathophysiological process involved in the disease. In the present study we addressed the following questions: is insulin resistance the cause or the consequence of steatosis and fibrosis in chronic hepatitis C? In other words, is insulin resistance, at least in part, responsible for the occurrence and worsening of steatosis and fibrosis? What are the determinants of insulin resistance in chronic hepatitis C? The unique ability of HCV genotype 3 to promote by itself steatosis provides the opportunity to clarify and unravel the complex relationships between insulin resistance, steatosis, and fibrosis.

### PATIENTS AND METHODS Study population

Between June 2001 and May 2002, 192 consecutive patients with chronic hepatitis C underwent liver biopsy in our liver unit. All subjects had positive HCV viraemia, as detected by reverse transcription-polymerase chain reaction (RT-PCR) (Amplicor; Roche Diagnostic Systems, Branchburg, New Jersey, USA) and increased alanine aminotransferase (ALT) activity. Patients were retrospectively screened according to the following criteria: (1) the presence of non-cirrhotic chronic hepatitis on liver biopsy specimens greater than 10 mm; (2) no previous antiviral treatment; (3) no personal history of diabetes or a fasting serum glucose <7 mmol/l; (4) alcohol intake <20 g/day; (5) no regular use of drugs known to induce steatosis (for example, corticosteroids, valproic acid, amiodarone, perhexiline maleate, tamoxifen); (6) absence of serum hepatitis B surface antigen; (7) absence of co-existing chronic liver disease, such as autoimmune hepatitis, primary biliary cirrhosis, sclerosing cholangitis, haemochromatosis, Wilson's disease, or  $\alpha_1$  antitrypsin deficiency; and (8) absence of human immunodeficiency virus antibodies. A total of 141 patients met these criteria.

Abbreviations: HCV, hepatitis C virus; RT-PCR, reverse transcriptionpolymerase chain reaction; BMI, body mass index; ALT, alanine aminotransferase; HOMA, homeostasis model assessment; IRS-1, insulin receptor substrate 1

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Excluded patients had one or more exclusion criteria: cirrhosis in 17, alcohol intake >20 g/day in 34, personal history of diabetes in six, fasting serum glucose >7 mmol/l in 16, previous antiviral treatment in five, and inadequate biopsy in 31.

### **Clinical parameters**

The following data were recorded at the time of liver biopsy: age, sex, duration of HCV infection, estimated as the time elapsed since the date of contamination in patients with known risk factors (for example, date of transfusion or beginning of intravenous drug use) and liver biopsy, body mass index (BMI), and mean daily alcohol intake between the date of contamination and diagnosis of HCV infection. BMI was calculated as body weight (kg) divided by squared height in metres (kg/m<sup>2</sup>). Patients were interviewed using a standard questionnaire for their alcohol intake in the past five years.<sup>23</sup>

### Methods

### Laboratory determinations

After an overnight fast of 10 hours, venous blood samples were drawn to determine serum levels of the following parameters: alanine aminotransferase (ALT), aspartate aminotransferase, plasma glucose concentration, insulin, and C peptide. All studies were performed before 9am. Plasma glucose concentration was measured by a glucose oxidase method on a vitros 750 analyser (Johnson and Johnson Clinical Diagnostics, Rochester, New York, USA). Serum insulin was measured by monoclonal immunoradiometric assay (CIS Bio International, Gif sur Yvette, France). Serum C peptide was estimated by a immunoradiometric assay (CIS Bio International). The index of insulin resistance was calculated on the basis of fasting values for glycaemia and insulinaemia, according to the homeostasis model assessment (HOMA) method.<sup>24</sup> The formula for the HOMA model is as follows:

insulin resistance (HOMA IR) = fasting insulinaemia ( $\mu$ U/ml) × fasting glycaemia (mmol/l)/22.5.

Insulin resistance, determined by this method, correlates closely with more complex techniques, such as the euglycaemic clamp method, in both diabetic and non-diabetic subjects.<sup>25 26</sup> Because serum insulin levels may be increased secondary to advanced liver fibrosis, insulin secretion was compared in patients with and without fibrosis using the serum C peptide/insulin ratio. C peptide and insulin are secreted in equimolar amounts, and serum C peptide is not significantly cleared by the liver.<sup>27</sup>

### Virological methods

All subjects were reactive for anti-HCV antibodies using a third generation enzyme immunoassay EIA (Ortho HCV 3.0 Elisa (Ortho Diagnostic Systems, Raritan, New Jersey, USA) or Monolisa anti-HCV (Sanofi Diagnostics Pasteur, Marnesla-Coquette, France)). All patients were tested for HCV RNA by RT-PCR (Amplicor HCV; Roche Diagnostic Systems). HCV RNA was quantified by Quantiplex HCV RNA 2.0 Assay (Chiron Diagnostics, Emeryville, California, USA). HCV genotypes were determined using the Inno-Lipa II HCV (Innogenetics, Ghent, Belgium).

### Histological evaluation

Paraffin embedded biopsies were analysed by a single pathologist unaware of the clinical and biological data, except for the presence of chronic hepatitis C. This analysis was performed following haematoxylin-phloxin-safran, Perls, and picrosirius red staining, using the METAVIR group items. Steatosis was graded according to the percentage of hepatocytes containing cytoplasmic vacuoles. Only macrovesicular steatosis was taken into account. Steatosis was classified as significant when more than 10% of hepatocytes were affected. Hepatitis was finally graded and staged using the METAVIR scoring system.<sup>28</sup> <sup>29</sup> Hepatic iron concentration was determined according to the Barry and Sherlock method.<sup>30</sup>

Each liver biopsy specimen was analysed twice by the same pathologist in a blinded manner, with good intraobserver concordance.

### Statistical analysis

Qualitative values are expressed as percentages and were compared using Fisher's exact test. Quantitative values are expressed as medians (range) and were compared using the Mann-Whitney or Kruskal-Wallis test, when more than two classes, or Spearman's correlation test. For statistical analysis, patients infected with genotypes other than 3 were

	Genotype 1 (n = 113)	Genotype 3 (n = 28)	p Value*
Age (y)	43 (20–72)	42.5 (30-63)	NS
Sex ratio (M/F)	1.3	1.4	NS
Disease duration (y)†	18 (5-41)	17 (4-25)	NS
Alcohol (g/day)	0 (0-20)	0 (0-20)	NS
$BMI (kg/m^2)$	23.6 (14-39)	23 (19-32)	NS
ALT (UI/I)	47 (11–264)	50 (19–159)	NS
Viral load (10 <sup>6</sup> Eg/ml)	5.9 (0.2-120)	9.1 (0.2–104)	NS
Glycaemia (mm/l)	4.7 (2.8-6.9)	4.7 (3.6-6.7)	NS
Insulinaemia (µU/ml)	5.8 (0.7-51)	5.3 (2-30)	NS
C peptide (nm/l)	0.6 (0.1–3.2)	0.7 (0.3-1.4)	NS
HOMA IR	1.2 (0.1-8.2)	1 (0.3–6.6)	NS
Activity score >1 (%)	31	29	NS
Fibrosis score >1 (%)	30	54	0.03
% of steatosis	0 (0–60)	10 (0-80)	0.0007
Steatosis			
<10 (%)	71	40	< 0.0001
10–30 (%)	25	24	
>30 (%)	4	36	

Values are median (range) or percentage.

BMI, body mass index; ALT, alanine aminotransferase; HOMA IR, homeostasis model assessment for insulin resistance.

†In 80 patients with known risk factors.

\*Fisher's exact test or Mann-Whitney test.

	Genotype 1		Genotype 3		
	Steatosis<10% (n = 80)	Steatosis≥10% (n = 33)	Steatosis<10% (n = 13)	Steatosis≥10% (n = 15)	p Value*
Age (y)	41	49‡	44.5	43	0.0008
Sex ratio (M/F)	1.22	1.54	1	2.75	NS
Disease duration (y)†	18	18.5	16	17	NS
Alcohol (g/day)	0	0	0	0	NS
BMI	22.8	25.3‡	22.8	23	0.0003
ALT (IU/I)	46	60	47	72	NS
Insulinaemia (µU/ml)	4.8	9.4‡	5.4	5.2	< 0.0001
C peptide (nmol/l)	0.6	0.9±	0.5	0.6	0.01
HÓMA IR	1.1	2.1‡	1.1	1	0.001
Viral load (10 <sup>6</sup> Eq/ml)	6.4	3.4	2.6	33.5‡	0.009
Activity score	1	1	1	1	NS
Fibrosis score	1	2‡	1	2‡	0.002
Percentage of steatosis	0	20	1.5	40±	< 0.0001
Hepatic iron (µmol/g)	9.4	12.2	9.2	6.7	NS

Values are medians.

BMI, body mass index; ALT, alanine aminotransferase; HOMA IR, homeostasis model assessment for insulin resistance.

†In 80 patients with known risk factors.

\*Fisher's exact test or Kruskal-Wallis test; ‡Mann-Whitney test, p<0.05.

collected into a single group termed "genotype 1 patients". Risk factors for insulin resistance were assessed in genotype 1 patients by correlating HOMA IR with the following parameters: age at liver biopsy, sex ratio, disease duration, BMI, mean daily alcohol intake, ALT activity, viral load, activity score, fibrosis score, percentage of steatosis, and intrahepatic iron concentration. Risk factors for steatosis were assessed separately in genotype 1 and genotype 3 patients by correlating the percentage of steatosis to the following parameters: age at liver biopsy, sex ratio, disease duration, BMI, mean daily alcohol intake, viral load, HOMA IR, activity score, and intrahepatic iron concentration. Independent parameters were assessed by multiple regression. The role of insulin in fibrosis progression was assessed in genotype 1 patients. The following parameters were included in a logistic regression model, with extensive fibrosis (score >1) as a dependent variable: age at liver biopsy, sex ratio, BMI, fasting serum insulin, activity score, and percentage of steatosis. Differences were considered significant at a p value of <0.05.

### RESULTS

## Characteristics of genotype 1 and genotype 3 infected patients

A total of 141 patients met the inclusion criteria. The distribution of HCV genotypes was as follows: 78 patients were infected by genotype 1, eight by genotype 2, 28 by



Figure 1 Distribution of homeostasis model assessment for insulin resistance (HOMA IR) according to steatosis and hepatitis C virus genotype. \*p<0.05.

genotype 3, 16 by genotype 4, four by genotype 5, one by genotype 6, and six had an undetermined genotype. The 113 patients infected with genotypes other than 3 were collected into a single group termed "genotype 1 patients" and compared with the 28 patients infected with genotype 3. Their characteristics are listed in table 1. Both steatosis and fibrosis were more frequent and more severe in patients infected with genotype 3 than in those infected with genotype 1. Other characteristics, in particular those with a potential impact on both steatosis and fibrosis (namely, age, disease duration, alcohol consumption, BMI, serum ALT activity, HOMA index and histological activity score), did not differ between the two groups.

### Insulin resistance is associated with genotype 1 related steatosis but not with genotype 3 related steatosis

Table 2 shows the main features of patients infected with genotypes 1 and 3 according to the presence or absence of significant steatosis. Genotype 1 patients with a fatty liver were older, had a higher BMI, and were more frequently insulin resistant than those without a fatty liver or those infected with genotype 3, even with marked steatosis. These latter patients had a higher median viral load than that found in the other groups (table 2). HOMA IR distribution in patients is presented in fig 1. Median HOMA IR was significantly higher in patients with genotype 1 related steatosis than in those with genotype 3 related steatosis or patients without steatosis. In genotype 1 patients, univariate analysis showed four parameters significantly linked with the degree of steatosis: age, BMI, HOMA IR, and hepatic iron

Table 3         Risk factors for steatosis in univariate and multivariate analysis in patients infected with genotype 1						
	Univariate o	analysis	Multivariate analysis			
	Coefficient	p Value*	Coefficient	SE	p Value**	
Age	0.39	< 0.0001	0.096	0.085	0.32	
BMI	0.42	< 0.0001	0.18	0.22	0.06	
HOMA IR	0.56	< 0.0001	0.315	0.7	0.001	
Hepatic iron	0.22	0.03	0.027	0.028	0.75	

BMI, body mass index; HOMA, homeostasis model assessment. \*Spearman's test. \*\*Multiple regression.



Figure 2 Percentage of steatosis according to homeostasis model assessment for insulin resistance (HOMA IR) in genotype 1 patients.

concentration (table 3). Multivariate analysis showed that HOMA IR was the only variable independently related to fatty liver in genotype 1 infected patients (table 3, fig 2) while in patients infected with genotype 3, viral load was the only variable associated with fatty liver (r = 0.64, p = 0.003, fig 3).

#### Insulin resistance is specifically associated with steatosis but not with fibrosis in genotype 1 infected patients

Genotype 1 infected patients with extensive fibrosis (histological score >1) had a higher median HOMA IR than others (1.8 v 1.0; p = 0.0001). However, the association was no longer significant after adjustment for the degree of steatosis (fig 4). Serum C peptide/insulin ratio was similar in patients with or without extensive fibrosis (0.1 (0.03–0.6) v 0.1 (0.03–0.4); p = 0.8).

### Risk factors for insulin resistance in patients infected with genotype 1

In univariate analysis, seven parameters were significantly correlated with HOMA IR in patients infected with genotype 1: age at liver biopsy, disease duration, BMI, ALT activity, activity score, fibrosis score, and percentage of steatosis (table 4). In multivariate analysis ( $r^2 = 0.28$ ), only age and degree of steatosis were independently linked to HOMA IR (table 4).

### Risk factors for extensive fibrosis in patients infected with genotype 1

Among genotype 1 patients, 34 (30%) had extensive fibrosis (score>1). In univariate analysis, six parameters were significantly associated with fibrosis: age, male sex, BMI, insulinaemia, activity score, and steatosis (table 5). Alcohol consumption and duration since contamination were not significantly different between patients with or without extensive fibrosis (4.7 (10) v 6.6 (11.8) g/day (p = 0.4) and 18.5 (9.7) v 19.4 (7.6) years (p = 0.35), respectively). When we compared non-drinker patients (n = 89) with those drinking small amounts of alcohol (n = 29), the prevalence of extensive fibrosis was not significantly different between the groups (32 v 24%, respectively; p = 0.04). In multivariate analysis ( $r^2 = 0.21$ ), only steatosis and activity score were independently associated with extensive fibrosis (table 5).

### DISCUSSION

This study confirms that the pathophysiology of fatty liver associated chronic hepatitis C is different in patients infected with genotype 1 and 3. The study also shows that insulin resistance in patients infected with genotype 1 is the cause rather than the consequence of hepatic steatosis and fibrosis, and suggests that increased circulating insulin is a risk factor for fibrosis through insulin resistance induced steatosis.



Figure 3 Percentage of steatosis according to hepatitis C virus viral load in genotype 3 patients.



Figure 4 Distribution of homeostasis model assessment for insulin resistance (HOMA IR) according to steatosis and fibrosis in genotype 1 infected patients. \*p<0.05.

Although steatosis was more severe in patients infected with genotype 3, insulin resistance was associated with steatosis only in patients infected with genotype 1. Insulin resistance depends mainly on age and is a major risk factor for steatosis, independent of BMI. Steatosis was associated with more severe fibrosis, whatever the genotype, supporting the major role of steatosis, whatever its cause, in the progression of fibrosis. Indeed, while univariate analysis identified a significant link between circulating insulin level and fibrosis stage, multivariate analysis revealed that steatosis, but not insulin, was independently associated with fibrosis, suggesting an indirect effect of the circulating

Table 4Parameters significant homeostasis model assessment (HOMA IR) in univariate and m patients infected with genotype	tly associated with for insulin resistance nultivariate analysis in 1
Univariate analysis	Multivariate analysis
Coefficient p Value*	Coefficient SE p Value**

Fibrosis score	0.39	0.002	0.024	0.225 0.8
ALT	0.25	0.008	0.122	0.003 0.24
disease† BMI	0.41	<0.0001	0.103	0.03 0.27
Age Duration of	0.38 0.25	<0.0001 0.04	0.203 0.085	0.012 0.04 0.018 0.33

BMI, body mass index; ALT, alanine aminotransferase; HOMA, homeostasis model assessment.

\*Spearman's test; \*\*multiple regression.

†In 62 patients with known risk factors. Taking disease duration into account did not change the results.

Table 5	<b>Risk factors</b>	for extensive	fibrosis	in univar	riate and	multivariate	analysis i	n
patients i	nfected with	genotype 1						

	Univariate analysis			Multivariate analysis		
	Fibrosis ≤ 1 (n = 79)	Fibrosis >1 (n = 34)	p Value*	p Value**	OR (CI)	
Age >55 yrs (%)	11	35	0.007	0.14	-	
Male sex (%)	50	70	0.06	0.08	-	
BMI >25 (%)	27	53	0.02	0.3	-	
Insulin $>10 \mu\text{U/ml}$ (%)	14	38	0.006	0.7	-	
Activity score >1 (%)	21	53	0.002	0.016	3.4 (1.3-9.4)	
Steatosis ≥10% (%)	19	53	0.0006	0.02	3.6 (1.2-10.8)	

insulin level on fibrosis stage through a steatosis related pathway.

To clarify the intricate relationship between insulin resistance, steatosis, and fibrosis, the study was performed in two groups of patients with chronic HCV infection due to genotype 1 or genotype 3. The results of the present study confirm the cogency of this distinction as two distinct mechanisms appear to operate in HCV associated fatty liver. In genotype 1 infected patients, steatosis was linked to BMI, while in genotype 3 infected patients, steatosis was related to HCV viral load. Moreover, we took care to exclude patients with usual causes of steatosis, such as alcohol, drugs, or diabetes mellitus. In this selected population, the prevalence of significant steatosis (≥10%) was still high, reaching 60% in patients infected with genotype 3. These results are in accordance with previous studies showing a strong association between steatosis and genotype 3 infection.<sup>4-11 12</sup> Because cirrhosis is a well known cause of insulin resistance,<sup>31</sup> patients with biopsy proven cirrhosis were also excluded.

The mechanisms of development of insulin resistance in patients with chronic HCV infection are not well understood. It has been suggested that insulin resistance may result from steatosis, as excess free fatty acids could downregulate insulin receptor substrate 1 (IRS-1) signalling.32 This concept was further supported by recent evidence that reversing hepatic steatosis may improve insulin resistance in rats with diet induced fatty liver.33 Despite the significant relationship in genotype 1 infected patients, the lack of association between steatosis and insulin resistance in genotype 3 infected patients does not support this hypothesis. Shintani et al showed that insulin resistance preceded the occurrence of steatosis in transgenic mice expressing HCV core protein, suggesting that insulin resistance is not a consequence of hepatic steatosis in these mice.22 As expected, BMI was correlated with the degree of insulin resistance in univariate analysis.34 However, in multivariate analysis, the degree of insulin resistance depended mainly on the age of the patient. Finally, genotype 1 infected patients with fatty liver were more insulin resistant than genotype 3 infected patients, probably because they were older and had higher BMI values. It has been suggested that age associated decline in mitochondrial function could contribute to insulin resistance.35 Our results do not exclude other mechanisms. The higher prevalence of type 2 diabetes in patients with chronic hepatitis C suggests implementation of HCV infection itself.14-20 Impaired IRS-1 signalling could be a possible mechanism, as recently shown in non-obese/non-diabetic patients with chronic HCV infection.36 In a large cohort study, Hui et al showed a genotype specific association between chronic HCV infection and insulin resistance.<sup>21</sup> The significant link between genotype 1 related steatosis and insulin resistance in our study population tends to support this hypothesis.

As in previous studies, we found that steatosis ( $\geq 10\%$ ) was common among patients with chronic hepatitis C, occurring in 34% of biopsy specimens.<sup>1-3</sup> Our data confirm the strong correlation between the degree of steatosis and level of HCV viraemia in genotype 3 infected patients.<sup>4</sup> In patients infected with genotype 1, multivariate analysis demonstrated that insulin resistance was a risk factor for steatosis, independent of BMI. These data confirm the existence of two distinct entities: a group of patients infected with genotype 1 that may have steatosis secondary to metabolic causes such as insulin resistance, and a second group infected with genotype 3 that may have steatosis as a direct consequence of HCV infection.

In patients with chronic hepatitis C, we and others have shown a significant relationship between the degree of steatosis and severity of fibrosis.467 In the present study, steatosis was associated with fibrosis, irrespective of viral genotype. Because steatosis was associated with insulin resistance in genotype 1 infected patients, fibrosis could be the result of hyperinsulinaemia. In fact, it has been demonstrated that high levels of insulin and glucose could promote fibrogenesis by stimulating the release of connective tissue growth factor, a fibrogenic growth factor, from hepatic stellate cells.<sup>37</sup> In genotype 1 infected patients, fibrosis stage significantly correlated with circulating insulin levels in univariate analysis. When considering other potential risk factors of fibrosis, such as steatosis, insulin was no longer an independent risk factor. The relationship between steatosis and fibrosis could be explained by several other mechanisms, such as lipid peroxidation.<sup>38-41</sup> According to the "two hits hypothesis", steatosis could increase the sensitivity of hepatocytes to oxidative stress, the second hit being HCV infection itself in patients with chronic hepatitis C.39 Production of reactive oxygen species in an in vitro model expressing HCV core protein is consistent with this hypothesis.42 The second independent risk factor for fibrosis in the present study was the histological score of activity. This result agrees well with the findings of previous longitudinal studies that showed that the necroinflammatory score was predictive of the development of severe fibrosis in patients with chronic hepatitis C.43 44 In our study, neither the duration of contamination nor alcohol consumption was associated with fibrosis. The non-linear progression of fibrosis in chronic hepatitis C probably explains the former result.43 44 In most of the studies showing a significant association between alcohol and fibrosis, the cut off value was 20 g/day.5-7 In our study population, drinking less than 20 g/day, we did not find any significant effect of small amounts of alcohol on fibrosis.

In conclusion, increased circulating insulin is a risk factor for fibrosis in genotype 1 infected patients with chronic hepatitis C through insulin resistance induced steatosis. Accordingly, it may be speculated that intervention strategies to reduce insulin resistance associated with steatosis should peroxisome proliferator activated receptor  $\gamma$  agonists could be interesting therapeutic options for improving steatosis and fibrosis in HCV patients with insulin resistance.45 4

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