

ORIGINAL PAPER

OMENTIN SERUM CONCENTRATION AND HEPATIC EXPRESSION
IN CHRONIC HEPATITIS C PATIENTS – TOGETHER OR APART?

MICHAŁ KUKLA¹, MAREK WALUGA¹, BRYGIDA ADAMEK², MARZENA ZALEWSKA-ZIOB²,
JANUSZ KASPERCZYK³, ANDRZEJ GABRIEL⁴, RAFAŁ J. BUŁDAK⁵, BARBARA SOBALA-SZCZYGIEL⁶,
LUCJAN KĘPA⁶, KATARZYNA ZIORA⁷, KRYSZYNA ŻWIRSKA-KORCZALA⁵, EDWARD SURMA¹,
TOMASZ SAWCZYN⁵, MAREK HARTLEB¹

¹Department of Gastroenterology and Hepatology, Medical University of Silesia, Katowice, Poland

²Department of General Biology, Medical University of Silesia, Katowice, Poland

³Department of Environmental Medicine and Epidemiology, Medical University of Silesia, Katowice, Poland

⁴Department of Pathomorphology in Zabrze, Medical University of Silesia, Katowice, Poland

⁵Department of Physiology in Zabrze, Medical University of Silesia, Katowice, Poland

⁶Department of Infectious Diseases in Bytom, Medical University of Silesia, Katowice, Poland

⁷Department of Paediatrics in Zabrze, Medical University of Silesia, Katowice, Poland

Chronic hepatitis C (CHC) is accompanied by numerous metabolic disorders, partially associated with altered adipokine system regulation. Omentin (intelectin-1) is a novel adipokine known to play a pivotal role in metabolic regulation in CHC. In a group of 63 CHC patients (29 men/34 women) infected with genotype 1b, aged 6.6 ± 14.6 years, serum omentin levels and its gene expression in liver tissue were examined and their association with metabolic and histopathological features was assessed.

Serum omentin levels were significantly higher in CHC patients compared to controls ($p < 0.001$), regardless of sex, body mass index (BMI), insulin sensitivity and lipid concentrations. There was no correlation between serum omentin and *omentin* hepatic expression. Neither parameter was associated with any histological features. Serum omentin in non-obese CHC patients seems not to be related to metabolic disorders or liver pathology. *Omentin* hepatic expression shows no relationship with either serum omentin levels or histopathological features. This suggests different mechanisms regulating circulating omentin concentration and *omentin* hepatic expression in CHC.

Key words: omentin, adipokine, chronic hepatitis C, fibrosis, insulin resistance, steatosis, liver.

Introduction

Adipo(cyto)kines are fat-derived hormones and cytokines fundamental to the pathogenesis of metabolic syndrome (MS) and liver steatosis that play a central role in insulin homeostasis and necro-inflammatory activity [1, 2]. Adipokines have been suggested to in-

fluence liver fibrosis progression [3, 4]. Thus, they are considered to play an important role in interactions between a variety of systems, including adrenal, immune, and central and peripheral nervous systems [5].

Chronic hepatitis C (CHC), because of concomitant insulin resistance (IR) [6, 7], high prevalence of steatosis [8, 9], increased prevalence of impaired

glucose tolerance and/or type 2 diabetes mellitus (T2DM) [10, 11, 12] and changes in lipid metabolism [13], has been considered not only as a viral disease, but also as a metabolic liver disease. Moreover, IR is a strong predictor of poor response to antiviral therapy, alongside advanced hepatic steatosis and fibrosis [6, 14]. Numerous reports have demonstrated altered levels of some adipokines such as leptin, adiponectin, resistin, visfatin, vaspin, chemerin and fibroblast growth factor (FGF) 21 in CHC patients [4, 15, 16, 17, 18, 19, 20].

Yang *et al.*, looking for molecule(s) regulating omental and mesenteric fat depots, identified a novel 34-kDa adipokine gene which they named omentin [21]. Further analysis established the human *omentin* gene as consisting of 8 exons and 7 introns, presenting 100% identity with human intelectin [22]. It is mainly expressed in the vascular fraction of visceral fat depots but is barely detectable in subcutaneous adipose tissue and mature adipocytes [21, 23]. Omentin enhances insulin-stimulated glucose transport via increase of protein kinase B (PKB/Akt) phosphorylation in human subcutaneous and visceral adipocytes, suggesting the possibility of improving their insulin sensitivity [21]. Activation of adenosine monophosphate-activated protein kinase (AMPK) by omentin may result in inhibition of gluconeogenesis and lowering glucose levels [24]. Omentin has been suggested to exert anti-inflammatory activity. It attenuates C-reactive protein-induced angiogenesis potentially via the nuclear factor κ B signaling pathway, a potent proinflammatory signaling pathway [25]. Additionally, omentin inhibits tumor necrosis factor α (TNF- α)-induced vascular inflammation in human endothelial cells. Omentin-activated AMPK may directly inhibit lymphocyte adhesion to vascular endothelial cells [26]. The major circulating isoform in human plasma is omentin-1, correlating inversely with obesity and IR as determined by homeostasis model assessment for IR (HOMA-IR) yet correlated positively with adiponectin and high-density lipoprotein (HDL) cholesterol levels [27]. Plasma omentin levels were significantly lower in the morbidly obese, and they correlated inversely with glucidic metabolism parameters, while expression in visceral adipose tissue was significantly lower in morbidly obese women than in controls [28]. Circulating concentrations of adipokines, including omentin, were suggested to be sex-dependent, because sex hormones influence adipose tissue metabolism regulation [29]. Schäffler *et al.* noted important *omentin* mRNA expression in omental adipose tissue probes from patients with chronic inflammatory bowel diseases, suggesting that this adipocytokine plays a role in the defense against intestinal bacterial translocation in the context of Crohn's disease [22]. Nassif *et al.* reported elevated serum omentin-1 levels positively associated with

the liver enzymes aspartate and alanine aminotransferases (AST and ALT), postulating its implication in the pathogenesis and metabolic complications of CHC [30]. According to our best knowledge, there has been only one study that assessed serum omentin concentration in CHC patients, but the liver tissue *omentin* expression in CHC patients has not yet been explored.

Thus, the main objectives of this study were to evaluate serum omentin levels and *omentin* mRNA expression in the liver tissue and to assess their relationship with metabolic parameters and liver histology in men and women diagnosed with CHC.

Material and methods

Study group

The study was performed on 63 patients with CHC (29 men/34 women), aged between 19 and 70 years – average 46.6 ± 14.6 years. All of the patients were infected with HCV genotype 1b. Serum HCV-RNA was assayed with the reverse transcription polymerase chain reaction (RT-PCR) method (Amplicor Roche/Promega v.2 Diagnostic Test, Branchburg, NJ, USA) while virus genotype was assessed by a reverse-hybridization line probe assay (LiPA Versant Test, Milwaukee, WI, USA) and viral load by signal amplification nucleic acid probe assay for the quantitation of human hepatitis C viral RNA [Bayer Versant_HCV RNA 3.0 Assay (bDNA); Bayer Diagnostics, Berkeley, CA, USA]. All patients were naive for the antiviral treatment. Patients infected with other HCV genotypes, hepatitis B or HIV co-infection, drug or alcohol abuse and autoimmune, neoplastic, thyroid and psychiatric diseases, as well as diabetes mellitus, and renal or heart failure were excluded from the study.

Study group participants had their systolic and diastolic blood pressure and waist circumference measured. With body mass and height estimations, body mass index (BMI) was calculated. To enable further analysis, subgroups defined as BMI < 25 and BMI ≥ 25 kg/m² were also created. All patients underwent an oral glucose tolerance test for diagnosis of diabetes mellitus or impaired glucose tolerance. On the day of the liver biopsy, a single blood sample was drawn in the morning from all patients subjected to fasting. The samples were centrifuged, and sera were aliquoted and frozen at -70°C until further processing.

The control group consisted of 30 (15 men/15 women) healthy persons, with BMI ranging between 20 and 25, aged 47.9 ± 14.8 years (44.7 ± 14.9 and 51.1 ± 14.5 years men and women, respectively). A single blood sample in the fasting state was taken in the morning. Liver biopsy was not carried out in the control group.

The study protocol was approved by the Ethical Committee of the Medical University of Silesia in Katowice, Poland, and conformed to the ethical guidelines of the Declaration of Helsinki. Informed consent was obtained for all study participants.

Biochemical and serological assays

Biochemical parameters were measured using routine methods. The upper limit of ALT activity was set at 38 IU/l, AST at 40 IU/l, gamma-glutamyltransferase (GGT) activity at 50 IU/l and total serum bilirubin concentration at 17 μ mol/l. Insulin concentration was measured with DiaMetra Insulin EIA Kit, Cat. No DKO076 (DiaMetra, Italy). For IR estimation the homeostasis model assessment for IR (HOMA-IR) was calculated by the formula: fasting insulin level (mUI/l) \times fasting glucose level (mg/dl)/405. With respect to the HOMA-IR value, patients were divided into two subgroups – below and equal to or above 2.5. Serum omentin concentrations were assessed in duplicate by the immunoenzymatic method using the Omentin-1 kit (BioVendor, Czech Republic; cat. no RD 191100200R).

Liver histology

Liver biopsy was performed with the Hepafix kit (B. Braun, Melsungen AG, Germany) as part of the diagnostic routine before the antiviral therapy. Tissue samples were immediately divided into the higher part for histopathological examination and the smaller one stabilized in RNA $later$ (Sigma-Aldrich, St. Louis, USA) and frozen at -80°C for further molecular procedures. Biopsy specimens including at least eleven portal tracts were examined independently by two experienced pathologists. Histopathological features were assessed according to Scheuer's (necro-inflammatory activity, fibrosis), Brunt's (steatosis) and Kleiner's (ballooning degeneration) scales [31, 32, 33].

Liver tissue omentin expression

Total RNA was isolated from liver biopsy specimens of CHC patients using the RNeasy Mini Kit (Qiagen, Hilden, Germany), followed by removing trace amounts of genomic DNA with RNase Free DNase Set (Qiagen, Hilden, Germany). RNA was quantified by measuring the absorbance at 260 and 280 nm (NanoDrop 1000 Spectrophotometer, Thermo Fisher Scientific, Wilmington, USA) and the integrity was assessed by electrophoresis in 1.2% ethidium bromide stained agarose gel.

RNA isolates were used for cDNA synthesis by RT reaction. 2 μ g of total RNA was reverse transcribed into cDNA in a total volume 20 μ l using High Capacity RNA-to-cDNA Kit (Applied Biosystems, Foster City, USA) according to the manufacturer's instructions. Received cDNA was used to determine *omentin*

gene expression level by real-time quantitative PCR (RT-Q-PCR) assay (TaqMan system). TaqMan primers and probes for *omentin* and the housekeeping gene *GAPDH* were bought as ready to use assays: TaqMan Gene Expression Assays (Hs 00914740_g1 for *omentin*) and Human GAPD Endogenous Control (FAM/MGB Probe, Non-Primer Limited) for *GAPDH* (Applied Biosystems, Foster City, USA). RT-Q-PCR for both genes was performed in a volume 20 μ l on the ABI PRISM 7300 Real Time PCR Detection System (Applied Biosystems, Foster City, USA). For each run, a RT-Q-PCR mix was prepared on ice containing 10 μ l of Applied Biosystems Universal PCR Master Mix, 1 μ l of primers and probe mix and 8 μ l of H₂O (Qiagen, Hilden, Germany). To each well of a 96-well plate, 19 μ l of QRT-PCR mix and 1 μ l of cDNA samples were added. All PCRs were performed in duplicate. In all amplification reactions, a negative control was also included. Thermal cycling for both *omentin* and *GAPDH* was initiated with an incubation step at 50°C for 2 min, followed by a first denaturation step at 95°C for 10 min, and continued with 40 cycles of 95°C for 15 s, 60°C for 1 min. The standard curves for the housekeeping gene *GAPDH* and the target gene were generated by serial dilutions of the control cDNA (equivalent to 1 μ g of total RNA) in four 2-fold dilution steps. The expression levels of *omentin* and *GAPDH* genes in every sample were determined from the respective standard curve and the *omentin* gene expression was divided by the *GAPDH* gene expression to obtain a normalized target value (relative expression level).

Statistical analysis

The data are presented as mean \pm SD. Differences between groups were examined through non-parametric tests (Mann-Whitney or Kruskal-Wallis), linear correlation (Spearman) and logistic regression analysis using the Statistica software version 10.0 (StatSoft Inc. USA). For all the analyses, statistical significance was determined for values of $p < 0.05$.

Results

Clinical and demographic data and the comparison of CHC patients with the control group are summarized in Table I. HOMA-IR, but not serum glucose and insulin, was markedly increased in CHC patients compared to controls (Table I). There were no abnormalities in results of serum biochemical parameters in the control group. Women and men with CHC participating in our study did not differ from each other according to age or diastolic blood pressure, but importantly men had higher BMI, waist circumference, systolic blood pressure and GGT. It was also noted that there were no obese CHC patients and none of them had been diagnosed with hypertension before.

Table I. General characteristics of CHC patient group

	CHC PATIENTS N = 63	MEN N = 29 (46%)	WOMEN N = 34 (54%)	p*	CONTROLS	p**
Age (years)	46.6 ± 14.6	48.2 ± 16.6	45.3 ± 12.8	NS	47.9 ± 14.8	NS
BMI (kg/m ²)	24.8 ± 3.0	25.7 ± 2.6	24.0 ± 3.1	0.02	23.9 ± 3.3	NS
Waist circumference (cm)	83.4 ± 11.9	87.8 ± 11.2	79.4 ± 12.5	0.03	85.5 ± 8.7	NS
SBP (mmHg)	129.0 ± 15.2	135.0 ± 18.8	125.0 ± 12.5	0.02	115.0 ± 10.5	NS
DSP (mmHg)	79.1 ± 8.9	79.1 ± 8.9	80.0 ± 9.5	NS	77.1 ± 6.6	NS
Fasting insulin (μU/l)	11.6 ± 9.6	10.6 ± 6.3	12.4 ± 11.8	NS	10.0 ± 4.5	NS
Fasting glucose (mg/dl)	93.7 ± 16.0	95.1 ± 16.4	92.6 ± 15.8	NS	80.8 ± 9.0	NS
HOMA-IR	2.9 ± 2.7	2.5 ± 1.3	3.4 ± 3.5	NS	2.0 ± 0.5	0.03
Total bilirubin (μmol/l)	14.3 ± 8.7	15.1 ± 4.1	13.6 ± 11.2	NS	9.7 ± 3.5	0.02
GGT (U/l)	80.7 ± 86.9	104.8 ± 119.2	60.2 ± 34.5	0.04	26.0 ± 5.5	0.04
ALT (U/l)	75.4 ± 44.5	83.2 ± 50.8	98.8 ± 37.7	NS	25.6 ± 4.3	0.001
AST (U/l)	51.7 ± 25.3	54.2 ± 28.5	59.5 ± 22.5	NS	24.1 ± 3.8	0.004
Total cholesterol (mg/dl)	187.8 ± 117.1	199.5 ± 170.6	179.4 ± 53.8	NS	174.6 ± 33.2	NS
LDL cholesterol (mg/dl)	97.6 ± 45.6	85.0 ± 38.3	107.6 ± 49.0	NS	94.5 ± 20.2	NS
HDL cholesterol (mg/dl)	53.7 ± 24.6	50.3 ± 27.8	56.0 ± 22.3	NS	44.0 ± 12.8	NS
Triglycerides (mg/dl)	131.9 ± 57.8	136.2 ± 56.3	129.0 ± 59.6	NS	139.0 ± 45.0	NS
CRP (mg/dl)	1.7 ± 1.2	1.6 ± 1.1	1.8 ± 1.4	NS	1.5 ± 1.4	NS
HCV viral load (IU/l)	2053301 ± 364614	2781433 ± 487426	1432246 ± 1976597	NS		
Serum omentin (ng/ml)	14.68 ± 4.12	14.55 ± 4.66	14.84 ± 3.48	NS	10.51 ± 3.90	< 0.001
Omentin hepatic expression/GAPDH	0.57 ± 1.27	0.36 ± 0.56	0.75 ± 1.64	NS		

BMI – body mass index; SBP – systolic blood pressure; DSP – diastolic blood pressure; LDL – low-density lipoprotein; HDL – high-density lipoprotein; CRP – C-reactive protein; HCV – hepatitis C virus
*p – men vs. women; **p – CHC patients vs. controls

Serum omentin concentration was significantly higher in CHC patients compared to controls (14.68 ± 4.12 vs. 10.51 ± 3.90 ng/ml, respectively, OR 1.30; 95% CI: 1.14-1.49; p < 0.001) (Table I). There was no difference in serum omentin levels and its hepatic gene expression between men and women with CHC (Table I). Similarly, there was no difference in serum omentin levels between men and women in the control group (10.92 ± 3.99 vs. 10.11 ± 3.91 ng/ml, respectively).

Table II. Linear correlation between serum omentin and omentin expression in the liver tissue

	SERUM OMENTIN [NG/ML]		
	MEN	WOMEN	CHC PATIENTS
Liver <i>omentin</i> / <i>GAPDH</i>	r = 0.23 p = NS	r = -0.16 p = NS	r = -0.09 p = NS

There was no relationship between serum omentin concentration and its gene expression in the liver in CHC patients, either in the group as a whole or men or women analyzed separately (Table II).

The whole study group included 34 (54%) patients with BMI ≥ 25 kg/m². BMI ≥ 25 kg/m² was observed in 17 (50%) females and 17 (58.6%) males. In CHC patients there were no significant differences in serum omentin levels and *omentin* expression in liver tissue according to BMI and HOMA-IR either in the study group as a whole or in men or women analyzed separately (Table III).

There was no significant difference in serum omentin levels and *omentin* hepatic expression in CHC patients with various necro-inflammatory activity, steatosis and hepatocyte ballooning degeneration grades and hepatic fibrosis stage (Table IV). Logistic regression analysis of serum omentin and its gene expression in the liver tissue adjusted for fibrosis, steatosis, necro-inflammatory activity and ballooning degeneration grade showed no significant differences.

Table III. Serum omentin concentration and *omentin* hepatic expression in the study participants according to sex, BMI and HOMA-IR

	MEN		WOMEN		CHC PATIENTS	
	BMI < 25 KG/M ²	BMI ≥ 25 KG/M ²	BMI < 25 KG/M ²	BMI ≥ 25 KG/M ²	BMI < 25 KG/M ²	BMI ≥ 25 KG/M ²
Serum omentin (ng/ml)	14.11 ± 4.38	15.35 ± 2.70	14.66 ± 4.14	14.44 ± 5.25	14.43 ± 4.17	14.89 ± 4.13
Liver <i>omentin</i> /GAPDH	0.40 ± 0.67	0.33 ± 0.49	0.33 ± 0.58	1.17 ± 2.20	0.36 ± 0.61	0.75 ± 1.63
	HOMA-IR < 2.5	HOMA-IR ≥ 2.5	HOMA-IR < 2.5	HOMA-IR ≥ 2.5	HOMA-IR < 2.5	HOMA-IR ≥ 2.5
Serum omentin [ng/ml]	14.05 ± 2.87	15.58 ± 3.83	14.14 ± 5.05	14.28 ± 4.45	14.10 ± 4.11	14.96 ± 4.07
Liver <i>omentin</i> /GAPDH	0.16 ± 0.11	0.52 ± 0.88	1.26 ± 1.97	0.21 ± 0.16	0.76 ± 1.53	0.38 ± 0.65

There were no significant differences between analyzed subgroups with respect to omentin expression and serum omentin concentration.

Table IV. Serum omentin concentration and *omentin* hepatic tissue expression in CHC patients according to liver histology

HISTOPATHOLOGICAL RESULTS		NUMBER OF PATIENTS (%)	SERUM OMENTIN [NG/ML]	P	LIVER OMENTIN/GAPDH	P
Necro-inflammatory activity grade	1	13 (18.8)	13.92 ± 3.94	NS	0.20 ± 0.12	NS
	2	38 (55.1)	15.04 ± 3.98		0.67 ± 1.48	
	3-4	18 (26.1)	14.35 ± 4.77		0.59 ± 1.15	
Fibrosis stage	1	29 (42.0)	14.91 ± 4.43	NS	0.21 ± 0.14	NS
	2	29 (42.0)	14.40 ± 4.26		0.90 ± 1.82	
	3-4	11 (15.9)	14.83 ± 2.94		0.65 ± 0.89	
Steatosis presence	present	34 (49.3)	14.68 ± 4.59	NS	0.79 ± 1.65	NS
	absent	35 (50.7)	14.68 ± 3.73		0.36 ± 0.74	
Ballooning degeneration grade	0-1	21 (30.4)	14.82 ± 4.30	NS	0.69 ± 1.66	NS
	2	37 (53.6)	14.88 ± 4.25		0.59 ± 1.22	

Discussion

Adipokines have been found to exert a positive or negative impact on many biological processes, together with insulin sensitivity and inflammatory activity. The role of adipokines in CHC remains under intensive investigations [4, 16, 17]; thus analysis of omentin levels in serum together with its gene expression in liver tissue in CHC patients is the next step of this research. Omentin-1 (intelectin-1) is a newly identified adipokine that is highly and selectively expressed in visceral adipose tissue [22]. It falls under the category of being a “good adipokine”: its serum levels are significantly decreased in patients with components of the metabolic syndrome (MS) and other disease conditions [34].

In our study serum omentin levels were significantly higher in CHC patients compared to the control group. The results were similar to those obtained by Nassif *et al.*'s study showing significantly higher

serum omentin levels in CHC subjects. Nassif *et al.* additionally showed lowered serum omentin levels in CHC patients with T2DM. However, in those patients serum omentin concentration was still significantly higher than in diabetic patients without CHC [30]. The results of both studies suggest a pivotal role of omentin in CHC.

The previous study by Pan *et al.* revealed significantly lower serum omentin levels in patients with DM. Moreover, serum omentin was negatively correlated with HOMA-IR, fasting insulin, fasting glucose and BMI [35]. Circulating omentin levels were negatively associated with a multiplicity of metabolic risk factors, suggesting that omentin acts as a biomarker of metabolic disorders [36]. Keeping in mind the strong influence of DM and obesity on serum omentin levels [27, 28], to avoid the possible influence of metabolic abnormalities associated with obesity and metabolic syndrome on omentin profile, we decided to include only non-diabetic, normal weight

or overweight but not obese CHC patients. Contrary to Nassif *et al.* [30], in our study there was no correlation between serum omentin and HOMA-IR, fasting insulin and glucose. Our results indicate that HCV infection may be a predictor of increased omentin levels.

An additional aspect of the study is the direct impact of the virus on metabolic and inflammatory pathways – hence the choice of a homogeneous group of CHC patients consisting only of those infected with genotype 1b, in whom insulin resistance is the most distinct [37]. HCV genotype 1b evokes insulin resistance by the direct influence on the intracellular insulin signaling pathway or by indirect mechanisms related to metabolic disturbances [37, 38]. Steatosis in genotype 1b infected patients results mainly from metabolic abnormalities. On the other hand, in genotype 3 infection steatosis is mainly viral derived, with less emphatic metabolic disturbances [38]. Therefore it is difficult to compare the influence of both 1b and 3 genotypes on adipokine profile. In the study by Nassif *et al.* [30] there is no information about HCV genotype in analyzed patients. Therefore, this discrepancy may result from a different group of study participants. Contrary to the study by Nassif *et al.*, our CHC patients also included overweight patients. The study by Nassif *et al.* analyzed Egyptian patients. This raises two further issues. The first is that in Egypt HCV genotype 4 is dominant [39]. We can suppose that patients analyzed by Nassif *et al.* were mainly infected with genotype 4. The second issue is the racial divergence in serum omentin concentration. Mean plasma omentin-1 concentrations were higher in African Americans than in Caucasians [40]. This aspect may lead to a discrepancy in the obtained results, because our study included only Caucasians, whereas the study by Nassif *et al.* included Egyptian patients.

In relation to the lack of negative association between IR and omentin, which is contrary to the studies on patients with DM and obesity, it should be noted that IR in obese individuals is primarily metabolic and not viral, whereas in our study we preferred to concentrate on virally derived metabolic abnormalities. However, in obese CHC patients co-existence with nonalcoholic fatty liver disease (NAFLD) is common and exacerbates progression of viral disease by increasing IR and steatosis [4].

Important differences in circulating omentin levels related to estradiol and testosterone concentrations found by Luque-Ramírez *et al.* suggest sexual dimorphism of the serum adipokine profile. It is important to note that subjects with excess weight had lower omentin levels regardless of sex [29]. In our study group there were no important differences between men and women either in serum omentin levels or its gene liver expression. Hence, we did not find any

difference in these parameters between men and women according to HOMA-IR or BMI, but in our study group there were no obese subjects, unlike in the study by Luque-Ramírez *et al.*, where obese study subjects and those with excess weight were analyzed together [29]. In light of this, the relationships between sex hormone axes and the adipokine network need further analysis within a larger study group.

Adipokines, excreted primarily by adipose visceral tissue, are postulated to modify liver inflammation and fibrosis, especially by modulation of hepatic stellate cell biology [41]. Yilmaz *et al.* in their study, which included overweight or obese biopsy-proven NAFLD patients, 40% of whom were diabetic and over 60% of whom had metabolic syndrome, found a positive correlation of systemic serum omentin levels with C-reactive protein (CRP) and hepatocyte ballooning degeneration, suggesting its possible role in hepatocyte death [2]. This finding may be explained, at least partially, by the implication of omentin's role in immune and inflammatory responses as an agent generating anti-inflammatory signaling pathways [25, 26]. In our study, serum omentin levels were not associated with histopathological features of liver tissue specimens. Neither hepatocyte ballooning degeneration grade, necro-inflammatory activity and steatosis grade nor fibrosis stage were associated with serum omentin concentration. In our CHC patients, however, liver pathology was induced mainly by HCV infection while in the study by Yilmaz *et al.* the most important factor was abdominal obesity with a high amount of visceral adipose tissue. Thus serum omentin concentration was positively related to hepatocyte ballooning degeneration independent of insulin resistance and other features of the metabolic syndrome, which could suggest that it plays a role in the development and progression of chronic hepatitis. Hepatocyte ballooning degeneration as a hallmark of NAFLD evoking by enhanced oxidative stress is less evident in CHC. Moreover, as mentioned above, in CHC the impact of HCV on metabolic abnormalities and oxidative stress is strong. In NAFLD liver abnormalities result mainly from metabolic disturbances associated with metabolic syndrome. Due to different pathogenesis and various study groups, it seems difficult to compare the two studies, but the role of omentin in cell destruction and its influence on liver pathology in CHC should be further investigated.

Analysis of multiple tissues indicated that omentin is highly expressed in human omental fat, with much less in other organs but not in subcutaneous adipose tissue [21]. Administration of glucose and insulin to human omental adipose tissue explants resulted in a dose-dependent reduction of omentin-1 expression [42]. Thus, this may be a reason why NAFLD patients had higher levels of serum omentin.

Eisinger *et al.* analyzed omentin levels in systemic venous, hepatic venous and portal venous sera in patients with liver cirrhosis and controls. Cirrhotic patients had higher portal vein omentin levels, which also were increased in systemic venous serum compared with controls, but there was no association with portal hypertension complications [43]. The main cause of cirrhosis in those patients was alcoholic hepatitis, with only three out of 34 having cirrhosis as a result of CHC. In our CHC patients, different stages of fibrosis were revealed, but none of them correlated with serum omentin levels. This poses the question as to whether higher serum omentin levels resulted from slower liver metabolism of this molecule in chronic liver impairment.

Omentin expression in CHC patients' liver tissue has not been reported so far. Nevertheless, in our study group no connections have been shown, either with routinely analyzed histopathological parameters or with the sex of the patients. *Omentin* was found highly expressed in human visceral but not in subcutaneous adipose tissue, but Yang *et al.* did not reveal its expression in liver tissue using Northern analysis [21]. The authors applied procedures (TaqMan Gene Expression Assays (Hs00214137_m1)) which resulted in a hybridization product located at the 3rd and 4th exons' boundary, while in our experiment starters (Hs 00914740_g1 for omentin) hybridized with a fragment involving the 1st and 2nd exons' boundary. Moreover, in this assay the 5'UTR (5' untranslated region) is involved, which could stabilize the transcript and probably help in product detection. Further questions remain as to whether *omentin* expression in the liver is an important source of serum omentin in non-obese individuals and whether the HCV infection plays a stimulating/modulating role.

In summary, our study reported for the first time hepatic *omentin* expression in non-obese, non-diabetic CHC patients. The study did not find any important relationship between hepatic *omentin* and serum omentin levels and sex, metabolic features, routinely analyzed laboratory parameters or liver histology. These results suggest that neither serum omentin nor hepatic *omentin* influence metabolic disorders and/or liver pathology regardless of sex in CHC. Surprisingly, there was also no correlation between serum omentin concentration and hepatic *omentin* expression. The lack of such a correlation points to different mechanisms regulating circulating omentin-1 concentration and hepatic *omentin* in CHC. Additional studies are required to determine whether levels of omentin are increased in CHC patients as a compensatory mechanism against increased IR or as a manifestation of poor omentin-1 metabolism in the HCV-infected liver, or both. *Omentin* liver tissue expression in CHC and its potential role in liver pathology require clarification also.

The authors declare no conflict of interest.

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Address for correspondence

Michał Kukla

Medical University of Silesia

Department of Gastroenterology and Hepatology

Medyków 14

40-752 Katowice

tel. +48 32 789 44 01

e-mail: kuklamich@poczta.onet.pl