

Serum and ascitic fluid hepcidin in HCV positive liver cirrhosis with and without HCC

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Abstract: Background & Aim: Chronic HCV infection suppresses hepatic hepcidin expression which may enhance iron toxicity and lead to disease progression and HCC development. The aim of the study is to investigate the role of hepcidin in HCV+ve liver cirrhosis patients in relation to disease progression and HCC development. Patients and methods: The study population consists of: 20 HCV+ve patients without HCC (HCV patients), 20 HCV+ve patients with HCC (HCV-HCC patients) and 10 controls. In addition to comprehensive clinical examination, they were subjected to laboratory check-up for albumin, bilirubin, PT %, ferritin, AFP and hepcidin. Ascitic fluid hepcidin was done for all patients. Results: There was a significant difference among HCV and HCV-HCC patients and controls with regard serum ferritin and hepcidin ($P = 0.001$ & 0.0001 respectively). Serum hepcidin of HCV and HCV-HCC patients were significantly lower than controls ($P = 0.0001$). Serum and ascitic fluid hepcidin of HCV-HCC patients was significantly lower than HCV patients ($P = 0.01$ & 0.02 respectively). Serum ferritin was significantly higher in HCV and HCV-HCC patients than controls ($P = 0.001$). Serum ferritin of HCV-HCC patients was significantly higher than HCV patients ($P = 0.02$). Ascitic fluid hepcidin was negatively correlated with Child-Pugh score in HCV ($r = -0.55$ & $P = 0.01$) and HCV-HCC patients ($r = -0.53$ & $P = 0.02$). Ascitic fluid hepcidin was negatively correlated with bilirubin in HCV ($r = -0.43$ & $P = 0.04$) and HCV-HCC patients ($r = -0.47$ & $P = 0.04$). Ascitic fluid hepcidin was positively correlated with serum albumin in HCV ($r = +0.44$ & $P = 0.04$) but there was no correlation in HCV-HCC patients ($r = -0.1$ & $P = 0.7$). Conclusion: Low levels of hepcidin may be involved in the pathophysiologic mechanism of iron overload in patients with chronic HCV with and without HCC. Moreover, there is a positive relationship between hepcidin levels and synthetic liver function suggesting that uniform suppression of hepcidin may be linked to disease progression and HCC development. Further analysis is still required to evaluate its usefulness as a marker for early detection of HCC by serial measurement of hepcidin in blood and ascitic fluid.

Keywords: Heparin, HCC, HCV, Ferritin

1. Introduction

Infection with hepatitis C virus (HCV) is a common problem worldwide, affecting millions of people across all populations. Most acutely infected patients develop chronic hepatitis and become a potential source of virus transmission, and as many as 1 in 5 will develop cirrhosis and its complications such as hepatocellular carcinoma (HCC) ⁽¹⁾. HCC is the commonest primary cancer of the

liver. Incidence is increasing and HCC has risen to become the 5th commonest malignancy worldwide and the third leading cause of cancer related death ⁽²⁾. Egypt has the highest prevalence of HCV worldwide and has rising rates of HCC ^(3,4).

Heparin is a cysteine-rich circulating bioactive peptide that is predominantly secreted from the liver and excreted in the urine ⁽⁵⁾. Heparin is a novel peptide due to its dual role as an 'iron-hormone' and 'antimicrobial peptide'.

Hepcidin is likely to play a role in innate immunity⁽⁶⁾.

Hepcidin controls plasma iron concentration and tissue distribution of iron. It inhibits intestinal iron absorption by enterocytes in the duodenum through its binding to ferroportin and inducing its internalization and degradation⁽⁷⁾. It also causes a release of iron by macrophages and hepatocytes⁽⁸⁾. These mechanisms result in a decrease of serum iron concentration and increased intracellular iron content⁽⁹⁾.

Hepcidin was shown to be present in urine and serum. However, its expression by various organs such as salivary glands, tonsils, trachea, lung and prostate has also been reported⁽¹⁰⁾. Hepcidin is in bile and may act along with α -defensins against gut microflora⁽¹¹⁾. Arnold *et al.*⁽¹²⁾ demonstrate the presence of hepcidin in pleural and ascitic fluid with unknown biological activity. Shike *et al.*⁽¹³⁾ showed that in white bass liver, infection with the fish pathogen *Streptococcus iniae* increased hepcidin mRNA expression 4500-fold.

Synthesis of hepcidin is homeostatically increased by iron loading and decreased by anemia and hypoxia⁽¹⁴⁾. Hepcidin regulates serum iron levels during inflammation, infection and possibly also in cancer. Under these conditions iron is shifted from circulation into cellular stores in hepatocytes and macrophages, making it less available for invading microorganisms and tumor cells⁽¹⁵⁾.

At the opposite side of the spectrum, hepcidin deficiency appears to be the ultimate cause of most forms of hemochromatosis, either due to mutations in the hepcidin gene itself or due to mutations in the regulators of hepcidin synthesis^(14, 16).

HCV infection is associated with alterations in body iron homeostasis through a poorly understood mechanism. Chronic HCV infection suppresses hepatic hepcidin expression. The impairment of hepatic hepcidin production occurring with chronic HCV infection may enhance iron toxicity and lead to disease progression⁽¹⁷⁾.

Although, hepcidin mRNA expression was detected in surgical specimens from patients with colorectal cancer⁽¹⁸⁾, and renal cell carcinoma⁽¹⁹⁾, few studies investigated the relation of hepcidin and HCC development.

The aim of our study is to investigate the role of serum and ascitic fluid hepcidin in HCV positive liver cirrhosis patients in relation to disease progression and complications like HCC.

2. Patients and Methods

The study included 40 HCV positive liver cirrhosis patients (previously diagnosed by PCR for HCV-RNA) selected from Internal Medicine and Tropical Medicine Departments in Menofiya University Hospital. The patients were divided into: 20 HCV positive patients without HCC (HCV patients) and 20 HCV positive patients with HCC (HCV-HCC patients). They are 37 males and 3 females ranged between 38-72 years old. In addition, ten healthy individuals are selected as controls.

Patients were diagnosed clinically, by abdominal ultrasonography (liver and spleen status and degree of ascites) and by assessing viral markers (HBsAg, HCV Ab and quantitative PCR for HCV- RNA). HCC is further confirmed by triphasic CT of the liver and AFP. All HCV-HCC patients were newly diagnosed cases and did not receive prior chemotherapy. Exclusion criteria included HBsAg positive patients, patients with concomitant causes of chronic liver diseases and history of alcohol consumption. Patients with hemochromatosis or with previous history of any malignancy are excluded from the study as those are expected to exhibit abnormal hepcidin regulation. All patients and controls gave their written informed consent before participating in the study. The study was approved by our Ethical Committee of Faculty of Medicine.

For all patients and controls, full history taking and complete clinical examination were done. A blood sample was withdrawn by sterile venipuncture and divided into three parts: EDTA was added to the first sample which was used for complete blood count (using the cell counter, Pentra 80, France). A second sample with sodium citrate was used to estimate prothrombin activity. The third sample was put in a plain vacutainer tube which was left to clot at 37°C then centrifuged and serum was separated. Serum was used to estimate AST, ALT, albumin, bilirubin, blood urea and serum creatinine (Beckman Synchro CX5 Clinical System, Minnesota, USA). Serum AFP measurements was done on AxSYM (AxSYM, Abbott Diagnostics, USA). Serum ferritin was measured using Ferritin AccuBind ELISA kit (Monobind Inc. Düsseldorf, Germany). A double-antibody sandwich enzyme-linked immunosorbent assay (ELISA) was used to assay the level of Human Hepcidin (Hepc) in probably diluted serum samples (Glory Science Co., Ltd, USA) Hepcidin in the sample was added to monoclonal antibody Enzyme well which is pre-coated with Human Hepcidin monoclonal antibody, followed by incubation; then, Hepcidin antibodies labeled with biotin, and combined with Streptavidin-HRP to form immune complex was added; then carrying out incubation and washing again to remove the uncombined enzyme. Then Chromogen Solution A, B was added; the color of the liquid changed into the blue and at the effect of acid, the color finally became yellow. The chroma of color and the concentration of the Human Substance Hepcidin of sample were positively correlated. Ascitic fluid hepcidin was done for all patients.

2.1. Statistical Analysis

Results are presented as mean \pm standard deviation (SD) unless otherwise stated. For comparison of two means, the unpaired *t test* and non-parametric *Mann-Whitney test* were used. The *ANOVA test* with post hoc was used to compare among HCV positive patients with and without HCC and controls. *Fisher Exact analysis* was also applied to compare proportions between groups. *Pearson coefficient* was used to study correlations between different parameters. A *P*-

value of ≤ 0.05 was considered statistically significant. All statistical analyses were performed using the Statistical Package for Social Science (SPSS) software version 10.

3. Results

Comparison among HCV patients, HCV-HCC patients and controls (Table 1) showed that there was a significant difference with regard serum ferritin and serum hepcidin levels ($P = 0.0001$ and 0.001 respectively). Serum hepcidin level of HCV and HCV-HCC patients were significantly lower than controls ($P = 0.0001$ for both). Serum hepcidin level of HCV-HCC patients was significantly lower than HCV patients ($P = 0.01$). Serum ferritin level was significantly higher in HCV and HCV-HCC patients than controls ($P = 0.001$ for both). Serum ferritin level of HCV-HCC patients was significantly higher than HCV patients ($P = 0.02$).

Table 2 shows a comparison between HCV and HCV-HCC patients. Ascitic fluid hepcidin was significantly lower in HCV-HCC than in HCV patients ($P = 0.02$). Blood

urea of HCV-HCC patients was significantly higher than HCV patients ($P = 0.003$). HCV-HCC patients had significantly lower serum albumin than HCV patients ($P = 0.0001$). HCV-HCC patients had significantly higher AFP level than HCV patients ($P = 0.0001$).

The severity of the liver disease was graded according to Child-Pugh's criteria. Three patients (15%) belonged to Child's class B in HCV and HCV-HCC patients. Seventeen patients (85%) belonged to Child's class C in HCV and HCV-HCC patients.

Table 3 shows correlation between ascitic fluid hepcidin and different parameters in HCV and HCV-HCC patients. Ascitic fluid hepcidin was negatively correlated with Child-Pugh score in HCV patients ($r = -0.55$, $P = 0.01$) and HCV-HCC patients ($r = -0.53$, $P = 0.02$). Ascitic fluid hepcidin was negatively correlated with total bilirubin level in HCV patients ($r = -0.43$, $P = 0.04$) and HCV-HCC patients ($r = -0.47$, $P = 0.04$). Ascitic fluid hepcidin was positively correlated with serum albumin level in HCV patients ($r = +0.44$, $P = 0.04$) but there was no correlation in HCV-HCC patients ($r = -0.1$, $P = 0.7$).

Table 1. Comparison among HCV patients, HCV-HCC patients and controls (data expressed as mean \pm SD)

	HCV patients (n=20)	HCV-HCC patients (n=20)	Controls (n=10)	P-value	LSD
Age (years)	57.1 \pm 7.9	56.1 \pm 8.2	55 \pm 7.3	0.8	
Gender (male %)	90% (n=18)	95% (n=19)	80% (n=8)	NS	
Serum hepcidin (pg/ml)	16175 \pm 892.6	15250 \pm 723.7	18100 \pm 516.4	0.0001*	*HCV vs Controls *HCV-HCC vs Controls *HCV vs HCV-HCC
Serum ferritin (mg/dl)	244.8 \pm 92.3	299.2 \pm 83.9	145 \pm 41.9	0.001*	*HCV vs Controls *HCV-HCC vs Controls *HCV vs HCV-HCC

n=number, *=significant, PT%=Prothrombin activity, WBCs=White blood count, AFP=Alpha fetoprotein, LSD= Least significant difference

Table 2. Comparison between HCV and HCV-HCC patients (data expressed as mean \pm SD)

	HCV patients (n=20)	HCV-HCC patients (n=20)	P-value
Hemoglobin (gm/dl)	8.7 \pm 1	9.4 \pm 1.6	0.1
WBCs ($\times 10^3$ /mm ³)	8.6 \pm 4.6	10.1 \pm 6	0.7
Platelet count ($\times 10^3$ /mm ³)	94.1 \pm 42.2	84.8 \pm 40.9	0.8
Blood urea (mg/ dl)	41.8 \pm 23.6	122.4 \pm 80.5	0.003*
Serum creatinine (mg/dl)	1.7 \pm 0.5	1.7 \pm 1	0.9
AST (U/L)	55.6 \pm 21.4	81.6 \pm 63.9	0.3
ALT (U/L)	63.6 \pm 29.8	58.8 \pm 45.7	0.2
Serum albumin(gm/dl)	2.68 \pm 0.27	2.15 \pm 0.41	0.0001*
Total bilirubin(mg/dl)	3.6 \pm 1.5	3.9 \pm 2.6	0.6
PT (%)	46.8 \pm 9.9	43.9 \pm 11.8	0.4
Child-Pugh score (points)	11.6 \pm 1.7	11.4 \pm 1.7	0.8
AFP (ng/ml)	8.8 \pm 5.4	629.5 \pm 104.6	0.0001*
Ascitic hepcidin (pg/ml)	24677 \pm 3451	22713 \pm 2983	0.02*

n=number, PT%=Prothrombin activity, WBCs=White blood count, AFP=Alpha fetoprotein, AST= Aspartate transaminase, ALT= Alanine transaminase

Table 3. Correlation between serum and ascitic fluid hepcidin and different parameters in HCV and HCV-HCC patients.

	Serum hepcidin				Ascitic fluid hepcidin			
	HCV patients (n=20)		HCV-HCC patients (n=20)		HCV patients (n=20)		HCV-HCC patients (n=20)	
	<i>r</i>	<i>P</i> -value	<i>r</i>	<i>P</i> -value	<i>r</i>	<i>P</i> -value	<i>r</i>	<i>P</i> -value
Age (years)	+0.07	0.8	-0.28	0.3	-0.08	0.7	-0.26	0.3
Hemoglobin (gm/dl)	-0.2	0.3	+0.09	0.7	-0.11	0.7	-0.24	0.3
WBCs ($\times 10^3/\text{mm}^3$)	-0.2	0.3	-0.27	0.3	-0.2	0.3	+0.2	0.3
Platelet count ($\times 10^3/\text{mm}^3$)	-0.2	0.3	-0.26	0.3	-0.2	0.3	+0.07	0.8
Blood urea (mg/ dl)	+0.3	0.2	+0.15	0.5	-0.006	0.9	-0.009	0.9
Serum creatinine (mg/dl)	+0.14	0.6	-0.27	0.3	-0.14	0.5	+0.14	0.6
AST (U/L)	+0.2	0.3	+0.14	0.6	+0.2	0.3	+0.11	0.7
ALT (U/L)	+0.2	0.3	+0.2	0.4	-0.2	0.3	-0.15	0.6
Serum albumin (gm/dl)	-0.01	0.9	-0.04	0.9	+0.44*	0.04*	-0.1	0.7
Total bilirubin(mg/dl)	-0.2	0.3	-0.09	0.9	-0.43	0.04*	-0.47	0.04*
PT (%)	-0.2	0.3	-0.19	0.3	-0.03	0.8	+0.04	0.9
Child-Pugh score (points)	+0.09	0.7	+0.2	0.4	-0.55	0.01*	-0.53	0.02*
AFP (ng/ml)	-0.05	0.8	+0.04	0.9	-0.14	0.5	+0.13	0.6
Serum ferritin (ng/ml)	+0.3	0.2	+0.17	0.5	+0.03	0.9	-0.16	0.6

n=number, PT%=Prothrombin activity, WBCs=White blood count, AFP=Alpha fetoprotein, AST= Aspartate transaminase, ALT= Alanine transaminase,

4. Discussion

Chronic hepatitis due to HCV infection is a leading cause of liver-related mortality all over the world, due to progression to cirrhosis and HCC. A high prevalence of iron overload has been shown in patients with chronic HCV infection characterized by different genetic backgrounds and exposure to environmental factors, suggesting that several mechanisms are involved, including inflammation, alteration of iron sensing, and deregulation of hepcidin release by hepatocytes^(17, 20). Elevated iron stores have been reported to affect the outcome of antiviral therapy, and to promote fibrogenesis and the risk of HCC⁽²⁰⁾.

Hepcidin, as an antimicrobial peptide, was first described as a liver expressed antimicrobial peptide 1⁽²¹⁾. Hepcidin is synthesized as preprohepcidin and undergoes posttranslational processing before release into circulation in an active form. The peptide hepcidin is proposed to be the key mediator of iron metabolism and systemic distribution. Therefore, hepcidin is considered as a negative regulator of iron release into the system by duodenal enterocytes and reticuloendothelial macrophages⁽²²⁾. Hepcidin is synthesized by hepatocytes in response to both iron overload and inflammatory stimuli, an effect believed to be dependent on cytokine production^(5, 21). Hepcidin exerts its regulatory effect by preventing the efflux of intracellular iron from intestinal enterocytes as well as hepatocytes, macrophages and placental cells⁽⁶⁾. Furthermore, hepcidin expression in the liver was negatively correlated with the total iron stores⁽¹⁷⁾. Iron accumulation in the liver, where hepcidin is exclusively

synthesized, is common in patients with chronic liver diseases, especially in patients with chronic hepatitis C virus (CHC) infection^(23, 24). Excess iron deposition in the liver is known to be hepatotoxic and may exacerbate liver injury in patients with chronic hepatitis C⁽²⁵⁾.

In the present study, Serum hepcidin level of HCV positive patients was significantly lower than healthy controls ($P=0.0001$). The results are in agreement with elegant studies in animal and cellular models that suggest a direct effect of HCV on liver hepcidin expression. Nishina *et al.*⁽²⁶⁾ studied transgenic mice expressing HCV polyprotein, which showed mild progressive hepatic iron accumulation. These mice had reduced hepcidin messenger RNA (mRNA) expression, which was attributed to HCV protein-induced reactive oxygen species (ROS), with consequent upregulation of an inhibitor of the binding of the transcription factor CCAAT/enhancer-binding protein α to the hepcidin promoter. Previously, Fujita *et al.*⁽²⁷⁾ reported that expression of hepcidin in patients with chronic hepatitis C was relatively lower than hepcidin expression in HBV positive patients and noninfected patients. Moreover, Girelli *et al.*⁽²⁸⁾ demonstrated that hepcidin concentration was significantly lower in CHC patients than those of matched controls. Hepcidin downregulation is likely to contribute to liver iron accumulation in this condition. According to the hypothesis of a direct suppressive effect of HCV on liver hepcidin expression, one could anticipate an inverse relationship between viral loading and circulating hepcidin levels. Recently, Liu *et al.*⁽²⁹⁾ further examined the anti-HCV activity of hepcidin by gene over-expression and

knockdown. The data clearly show that over-expression of hepcidin reduced HCV expression, while knockdown of hepcidin induced viral RNA expression. These results indicate that hepcidin might have a broad anti-HCV activity, at least for genotype 1 and genotype 2. The inhibitory effect appears to occur at the viral replication level. Taken together, these results suggest that both exogenous addition of hepcidin peptide and over-expression of hepcidin could attenuate HCV replication in cell models⁽²⁹⁾.

It is possible to develop a therapy using hepcidin. Besides its antiviral effect, the potential advantage of hepcidin therapy for HCV patients is restoration of iron homeostasis. It will be interesting to investigate the therapeutic efficacy of both antiviral activity and iron metabolism in small animal models or possible clinical studies. However, well-designed clinical studies addressing safety and longterm efficacy are needed in order to clarify the risks and benefits of hepcidin-targeted treatment.

On the other hand, Aoki et al.⁽³⁰⁾ reported that hepatic hepcidin expression is increased in response to iron overload in patients with chronic hepatitis C.

In anemia of chronic diseases, hepcidin mRNA expression is influenced by acute-phase reactant cytokines as IL-6⁽³¹⁾, while HCV infection is a cell-mediated immune response that is characterized by a T helper-1 immune response with secretion of IL-2, IL-4, IL-10, tumor necrosis factor- α , and interferon- γ ⁽³²⁾. Thus hepcidin expression may be less influenced by T helper-1 cytokines in chronic HCV infection.

We examined the relationship between parameters reflecting hepatic synthetic function and hepcidin level in blood and ascitic fluid. Serum albumin level was positively correlated with ascitic fluid hepcidin levels in HCV positive patients without HCC ($r = +0.46$, $P = 0.04$). Serum bilirubin was negatively correlated with ascitic fluid hepcidin in HCV positive patients with ($r = -0.47$, $P = 0.04$) and without HCC ($r = -0.43$, $P = 0.04$). Further study is necessary to determine the participation of hepcidin in liver function. Moreover, there was a significant inverse correlation between Child-Pugh score as an indicator of severity of the liver disease and ascitic fluid hepcidin. This correlation may be caused by high level of serum ferritin as an indicator of iron overload in our HCV positive patients. Hepatic iron overload is even more common among patients with end-stage liver disease due to hepatitis C⁽³³⁾. In contrast, Fujita et al.⁽²⁷⁾ reported that serum albumin level was not correlated with hepcidin level and serum bilirubin was positively correlated with hepcidin.

Hepcidin regulates serum iron levels during inflammation, infection and possibly also in cancer. Under these conditions iron is shifted from circulation into cellular stores in hepatocytes and macrophages, making it less available for invading microorganisms and tumor cells⁽¹⁵⁾. Therefore, the role of hepcidin in human cancer deserves to be studied, since there have been only few reports in this context^(18, 19, 34). It is well known that HCC develops in more than 40% of patients with hemochromatosis⁽³⁵⁾.

Furthermore, some studies have indicated that iron overload is a major risk factor for development of HCC⁽³⁶⁾. Iron overload leads to the generation of ROS, which cause chronic inflammation in the liver⁽³⁷⁾. It is noteworthy that iron overload has been found to facilitate oxidative stress, steatosis development and with triggering of hepatic stellate cells thus inducing liver fibrosis and hepatocarcinogenesis in a mouse model transgenic for HCV^(38, 39). In our study, serum ferritin was significantly higher in HCV positive patients with HCC than healthy controls. Serum hepcidin of HCV positive patients with HCC was significantly lower than healthy controls. The results are in agreement with previous studies. Hepcidin transcription was found to be downregulated in hepatoma cell lines expressing HCV core and nonstructural proteins⁽⁴⁰⁾. Moreover, some studies demonstrated that expression of hepcidin mRNA was suppressed universally in HCC, irrespective of the degree of tumor differentiation. Expression of hepcidin mRNA is constitutively suppressed in cancerous, but not in non-cancerous liver tissue of patients with HCC^(34, 41). Therefore, suppression of hepcidin hormone by hepatitis C virus is likely to be an important factor of liver iron accumulation in this condition⁽²⁸⁾. Iron is an essential nutrient for cell growth and particularly required by cancer cells to proliferate⁽⁴²⁾. Therefore, the down-regulation of hepcidin may stimulate tumor progression in chronic HCV infection patients. This finding suggests that, hepcidin may play a role in defending the body against HCC development.

Although the mechanism responsible for suppression of hepcidin mRNA expression in HCC remains unclear, suppression of hepcidin transcription contradicts the previously proposed scheme for iron homeostasis in cancer cells, because cancer cells must retain iron in order to proliferate. However, suppression of hepcidin is rational because duodenal enterocytes transfer iron to plasma, resulting in an increase of total body iron content. One explanation was provided by Weizer-Stern et al.⁽⁴³⁾ who reported that activation of the tumor suppressor gene p53 stimulates the expression of hepcidin. The promoter region of the hepcidin gene (HAMP) contains a putative p53 response element. Inactivation or mutation of the p53 gene has been detected in various types of human cancer⁽⁴⁴⁾, including HCC⁽⁴⁵⁾. Suppression of hepcidin expression may be linked to the altered expression and function of p53.

5. Conclusion

Low levels of hepcidin for the degree of iron burden may be involved in the pathophysiologic mechanism of increased iron overload in patients with chronic hepatitis C with and without HCC. Therefore, suppression of hepcidin by HCV is likely an important factor in liver iron accumulation in this condition. Iron overload has a potential negative effect in CHC in prognosis and development of HCC. The precise mechanism responsible for the suppression of hepcidin in HCC should be

investigated further, focusing on its role in the development and maintenance of this cancer. Supplementation of hepcidin may be beneficial for these conditions.

Moreover, there is a positive relationship between hepcidin levels and synthetic liver function suggesting that a uniform suppression of hepcidin may be linked to disease progression and development of HCC. Further analysis is still required to evaluate its usefulness as a marker for early detection of HCC by serial measurement of hepcidin in blood and ascitic fluid.

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