Inflammatory Biomarkers and Liver Histopathology in Non-Uremic and Uremic Chronic Hepatitis C Patients

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ABSTRACT

Background: The aim of this study is to investigate the association between hepatic activity index (HAI) and fibrosis score (FS) with inflammation biomarkers in non-uremic and uremic hepatitis C positive patients. Methods: Fifty chronic hepatitis C (cHepC) positive patients, having a liver biopsy were included in this study. Liver biopsies were scored according to modified ISHAC scoring system. 25 healthy controls of similar age and gender were also enrolled as control group. Serum YKL-40, neutrophil/lymphocyte ratio (NLR), thrombocyte/lymphocyte ratio (PLR), CRP and Immunoglobulin (IgG, A and M) levels were used to determine inflammation. AST to Platelet Ratio Index (APRI) score was also evaluated. According to biopsy findings patients were divided into 2 groups: low (0–2) and severe (3–6) FS. Results: Patients with cHepC had increased inflammation compared to the healthy controls. End-stage renal disease (ESRD) patients had higher levels of inflammation markers (NLR, IgG, CRP and YKL-40) and lower HCV RNA levels, HAI and FS compared to non-uremic patients. When patients were grouped into 2 according to FS as mild and severe, IgG (p < 0.001), YKL-40 (p = 0.02) levels and APRI score (p = 0.002) were significantly higher compared to mild FS (p = 0.002). YKL-40 levels (t value: 3.48; p = 0.001) and APRI score (t value: 3.98, p = 0.002) was an independent associated with FS in ESRD patients. Conclusion: In non-uremic cHepC patients, YKL-40 levels and APRI score may be valuable markers of FS. In ESRD patients, there is not sufficient data for prediction of HAI and FS. In these patients, APRI score may provide better information.

KEYWORDS

APRI score; hepatitis C; end-stage renal disease; inflammation biomarkers; YKL-40; liver histopathology

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INTRODUCTION

It is estimated that 2–3% of the world population has hepatitis C (HepC), varying by geographical location, time period and development status of countries (1, 2). Important risk factors for HepC are blood transfusions, solid organ transplantation from an infected donor, i.v. drug use, non-safe therapeutic injections, occupational percutaneous exposures and familial or dialysis-related exposures (3–5). Despite the fact that HepC incidence and prevalence have declined in the last years, mainly in the hemodialysis population, it is still a major cause of chronic liver disease, cirrhosis and hepatocellular carcinoma (1, 2, 5).

Liver biopsy is the gold-standard procedure for the diagnosis and treatment decision of HepC. However, it is an invasive procedure, leads to various complications, needs to be repeated and requires hospitalization and is costly. In addition, in uremic patients it may be more difficult to perform due to uremia-related coagulation problems (6). Thus, non-invasive markers are under intense investigation (7) with the literature still being limited.

Inflammatory events play an important role during the course of chronic hepatitis C (cHepC) (8–10). Main contributors are cytokine producing CD4+ T and CD8+ T cell proliferation, increased CD5+ B cells, IL-2, IFN- γ and TNF α (8–10). Neutrophil lymphocyte ratio (NLR) and thrombocyte lymphocyte ratio (PLR) are used as markers of inflammation in various diseases including atherosclerotic heart and renal disease. They are more sensitive compared to total leukocyte counts (11–13). However, there are no data to confirm their use in cHepC patients. Similarly, YKL-40 is a glycoprotein synthesized from neutrophils and macrophages that plays a role in inflammation and tissue remodelling. A number of studies have demonstrated the association between YLK-40 and fibrosis in liver disease (14–20).

In this study, we aimed to investigate the association between hepatic activity index (HAI) and fibrosis score (FS) with inflammation markers (NLR, PLR, YKL-40, IgG, IgA and IgM) in patients with cHepC with or without endstage renal disease (ESRD).

MATERIAL AND METHODS

50 cHepC patients, 15 with ESRD, having undergone percutaneous liver biopsy between December 2011 and June 2013 at a tertiary hospital were included. 25 healthy people with similar age and gender were used as controls. Exclusion criteria were co-infection with other viruses, chronic hepatitis due to other reasons, decompansated cirrhosis, primary or metastaic liver cancers, malignancies, serious congestive heart failure and pyschiatric disorders.

Data on age, gender, smoking, alcohol use, systemic diseases, coinfection, medication, height, weight were obtained from patients charts.

Liver biopsy was performed under ultrasound guidance. Informed consent was received from all patients prior to biopsy. Modified ISHAC Scoring was used for histopathological evaluation of the liver biopsies. According to biopsy findings patients were divided into 2 groups: low (0-2) and severe (3-6) FS.

NLR, PLR, IgG, IgA, IgM, CRP and YKL-40 levels were measured as markers of inflammation. Serum YKL-40 was detected by ELISA (Human Chitinase 3-Like 1, EuroimmunAnalayser, USA) according to the manufacturer's protocol. APRI score was calculated from 'increase in AST (patient's AST level/upper limit for AST)/Thrombocyte count × 100'.

In addition to informed consent for liver biopsy, all patients also gave informed consent for this study.

STATISTICAL ANALYSIS

All analysis were performed by using SPSS 15.0 for Windows. All values are reported as mean ± SD. The Pearson's Correlation was used for correlation analysis, student t-test and chi-square to compare two groups and ANOVA to compare more than two groups. Stepwise linear regression analysis was used to find independent predictors for HAI and FS. P value less than 0.05 was considered as statistically significant.

RESULTS

50 patients with cHepC (35 patients were non-uremic cHepC, 15 were ESRD patients) and 25 controls were enrolled. Mean age was 55 ± 11 years and 50% were male. Mean HCV RNA value was $3.6E^{+6}$ IU/mL \pm $6.5E^{+6}$ IU/mL. 39 patients had genotype 1, 1 was genotype 3 and 4 were genotype 4 (6 patients did not have genotype values). Mean HAI and FS scores were 8.26 ± 2.22 and 2.88 ± 1.54 , respectively.

cHepC patients had similar demographical parameters compared to controls. NLR (2.52 ± 1.26 vs. 1.91 ± 0.92 ; p = 0.03), serum IgG levels (1491 ± 459 mg/dL vs. 1143 ± 229 mg/dL; p < 0.001), YKL-40 levels (134 ± 170 ng/mL vs. 18 \pm 9.1 ng/mL, p < 0.001) were higher in cHepC patients. There were no differences with regards to total leukocyte count, PLR, CRP, IgM and Ig A levels, whereas liver function tests were higher.

Non-uremic cHepC patients were older, had higher Body mass index (BMI), AST and ALT levels compared to patients with ESRD. In addition, ESRD patients had higher levels of inflammation markers (NLR, IgG, CRP and YKL-40) and lower HCV RNA levels, HAI and FS. These data are presented in Table 1 and Table 2.

In patients with non-uremic cHepC patients HAI was positively correlated with serum IgG levels (r = 0.400, p = 0.017), serum IgM levels (r = 0.414, p = 0.013), GGT levels (r = 0.511, p = 0.002), YKL-40 levels (r = 0.459, p = 0.006) and FS (r = 0.671, p < 0.001). On the other hand, FS was positively correlated with serum IgG levels (r = 0.519, p = 0.001), AST levels (r = 0.528, p = 0.001), ALT levels (r = 0.427, p = 0.010), GGT levels (r = 0.439, p = 0.008), YKL-40 levels (r = 0.506, p = 0.002), APRI score (r = 0.616, p < 0.001) and HAI (r = 0.671, p < 0.001) and negatively with serum albumin levels (r = -0.493, p = 0.003). When patients were grouped into 2 according to FS as mild and severe, AST (p = 0.005), ALT (p = 0.01), IgG (p < 0.001), YKL-40 (p = 0.02)

	Non-Uremic cHepC patients (N = 35)	cHepC patients with ESRD (N= 15)	Controls (N = 25)	P value
Age (years)	59 ± 9.1	44 ± 10	57 ± 10	<0.001
Gender (M/F)	17/18	8/7	13/12	0.993
BMI (kg/m²)	26 ± 3.8	22 ± 2.4	27 ± 3.6	< 0.001
AST (U/L)	53 ± 28	24 ± 13	24 ± 16	< 0.001
ALT (U/L)	56 ± 34	28 ± 18	30 ± 26	0.001
ALP (U/L)	90 ± 26	163 ± 105	78 ± 25	< 0.001
GGT (U/L)	57 ± 48	44 ± 28	37 ± 70	0.559
Total Bilirubin	0.85 ± 0.35	0.66 ± 0.24	0.65 ± 0.25	0.065
Creatinine(mg/dl)	0.87 ± 0.18	7.2 ± 2.5	0.90 ± 0.15	< 0.001
Uric acid (mg/dl)	5.8 ± 3.7	5.20 ± 1.4	4.9 ± 1.7	0.671
Hemoglobin (g/dl)	13.4 ± 2.0	13.2 ± 2.0	13.4 ± 1.6	0.940
Platelets	208 ± 96	171 ± 74	258 ± 49	0.003
Lökosit (mm³)	6332 ± 1923	6785 ± 2461	6749 ± 1515	0.627
NLR	2.3 ± 1.4	2.8 ± 0.8	1.9 ± 0.9	0.05
PLR	134 ± 109	116 ± 44	129 ± 51	0.777
Serum albumin	4.0 ± 0.5	4.0 ± 0.3	4.0 ± 0.4	0.642
Immunoglobulin G (mg/dL)	1465 ± 432	1557 ± 527	1143 ± 229	0.004
Immunoglobulin M (mg/dL)	124 ± 65	100 ± 58	112 ± 52	0.399
Immunoglobulin A (mg/dL)	234 ± 116	230 ± 141	201 ± 83	0.541
CRP (mg/dL)	0.32 ± 0.27	1.92 ± 2.24	0.63 ± 0.81	< 0.001
Serum YKL-40 (pg/mL)	76 ± 128	271 ± 182	18 ± 9.1	< 0.001

All results are reported as mean ± SD. chepC = chronic hepatitis C, BMI = Body Mass Index, NLR = Neutrophil/Lymphocyte Ratio, PLR = Platelet/Lymphocyte Ratio, CRP = C-reactive protein, ESRD = End stage renal disease

Tab. 2: Comparison of inflammation markers and biopsy parameters
in chepC patients with and without renal failure.

	Non-Uremic cHepC patients (N = 35), Mean ± SD	cHepC patients with ESRD (N = 15), Mean ± SD	P value
NLR	2.3 ± 1.4	2.8 ± 0.8	0.159
CRP (mg/dl)	0.32 ± 0.27	1.92 ± 2.24	0.015
Immunglobulin G (mg/dL)	1465 ± 432	1557 ± 527	0.542
Serum YKL-40 (pg/mL)	76 ± 128	271 ± 182	0.001
APRI score	0.67 ± 0.43	0.36 ± 0.24	0.004
HCV RNA	4.4E ⁺⁶ ± 7.7E ⁺⁶	1.8E ⁺⁶ ± 2.9E ⁺⁶	0.201
Hepatic Activity Index	8.77 ± 2.19	7.07 ± 1.83	0.011
Fibrosis Score	3.26 ± 1.59	2.00 ± 1.00	0.007

All results are reported as mean ± SD. chepC = chronic hepatitis C, NLR = Neutrophil/Lymphocyte Ratio, CRP = C-reactive protein , ESRD = End stage renal disease

levels and APRI score were significantly higher in the latter (p = 0.002).

In stepwise linear regression analysis, serum IgG and GGT levels were found as independent predictors for HAI, whereas YKL-40 levels and APRI score were found as predictors for FS (Table 3).

In patients with ESRD, HAI was positively correlated with age (r = 0.656, p = 0.008), serum IgG levels (r = 0.632, p = 0.011), AST levels (r = 0.538, p = 0.039), APRI score (r = 0.617, p = 0.014) and FS (r = 0.858, p < 0.001). FS was posi-

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	Standardized β coefficient	Unstandardized β coefficient	t value	P value		
Hepatic Activity Index, r ² = 0.414						
IgG levels	0.02	0.392	2.89	0.007		
GGT	0.21	0.524	3.72	0.001		
Fibrosis Score, r ² = 0.550						
YKL 40 level	0.005	0.418	3.48	0.001		
APRI	1.880	0.549	4.57	< 0.001		

Tab. 3: Stepwise linear regression analysis for predictors of Hepatic Activity Index and Fibrosis Score in non-uremic cHepC patients.

APRI1.8800.5494.57<0.001</th>Included variables: Age, IgG, GGT and YKL40 for HAI and age, IgG,
APRI and YKL40 for FS.

tively correlated with age (r = 0.507, p = 0.04), AST levels (r = 0.610, p = 0.016), APRI score (r = 0.733, p = 0.002) and HAI (r = 0.858, p < 0.001) and negatively with thrombocyte count (r = -0.563, p = 0.029). When patients were grouped into 2 according to FS as mild (67%) and severe (33%), AST (p = 0.005), ALT (p = 0.01), IgG (p < 0.001), YKL-40 (p = 0.02) and APRI score (p = 0.002) were higher in the severe group. These patients were older (40 ± 7.7 vs. 53 ± 10; p = 0.01) and had higher APRI score (0.28 ± 0.21 vs. 0.53 ± 0.21, p = 0.06).

In stepwise linear regression analysis (variables: age, IgG, APRI and YKL40), age (t value = 2.97, p = 0.011) and APRI score were independent predictors for HAI (t value = 2.68, p = 0.020). For FS, only APRI score was an independent predictor (t value = 3.98, p = 0.002) (variables: Age, IgG, APRI and YKL40).

DISCUSSION

The results of our study show that YKL-40 levels are independently associated with FS in non-uremic cHepC patients. A similar association could not be confirmed in patients with ESRD.

Many noninvasive biomarkers of fibrosis will be crucial for successful individualized management of disease activity in cHepC patients (7). Currently studies have reported a relation between laminin, hyaluronic acid, type IV collagen, fibronectin, matrix metalloproteinases, metalloproteinase inhibitors, procollagen type III N-terminal propeptide and serum YKL-40 levels with liver fibrosis (7, 15).

YKL-40 is a glycoprotein that plays a role in inflammation and tissue remodelling (14). There remains a controversy regarding the role of YKL-40 in patients with cHepC (7, 16–20). Saitou et al. (16) reported increased levels of YKL-40 in 109 patients with cHepc having severe fibrosis. Hovewer, this could not be confirmed in renal transplant patients or hemodialysis patients (18, 20). In a genetics-based study, Berres et al. reported increased YKL-40 levels in patients with 131G->C polymorphism of the CHI3L1, which were associated with fibrosis. They also reported the protective effect of the G allele (19). In our study, serum YKL-40 levels were correlated with HAI and FS in non-uremic patients and independently associated with FS. However, this finding could not be confirmed in ESRD patients. This may be due to several reasons. Firstly, our study population was relatively small therefore results reaching a statistical power. Secondly, YKL-40 is excreted by the kidneys and the levels are very high compared to non-uremic patients. This may have a limiting effect by interfering with the equations in the statistical analysis. Thirdly, increased inflammation in dialysis patients may be interfering with the YKL-40 measurements (like the abnormal levels of ferritin in the same population) Fourthly but not the last, the fact that the increased levels of YKL-40 are mainly seen in patients with severe fibrosis, less frequency of FS in uremic patients may be limiting the association. Thus, we propose that YKL-40 measurement may provide limited information in ESRD patients.

High IgG levels in cHepC patients have been shown to affect prognosis and response to therapy (21–24). Maruyama et al. (21), reported a correlation between serum IgG levels and the severity of the disease in 102 patients with cHepC and significant decrease in levels by treatment. Gonzàlez-Quintela et al. and Watt et al. also described significantly higher levels of IgG and IgA in patients with cHepC (22, 23). Our results further confirm these studies: significantly higher immunoglobulin levels in cHepC patients compared to controls and correlation of IgG with HAI and FS in non-uremic patients. The association between HAI and IgG was independent. Similar to YKL-40, the association among IgG, HAI and FS could not confirm in ESRD patients.

In ESRD patients, we could not find an association between HAI and FS with the inflammation biomarkers. The most plausible reason may the confounding effect of inflammation due to uremia and in some patients due to comorbid diseases. On the other hand, the decreased viral load by hemodialysis, the release of protective IFN- α and hepatocyte growth factor and relatively lower FS may be affecting the association of inflammatory marker with fibrosis. Thus, the role of these markers may be questionable in ESRD patients.

APRI score is used to predict liver fibrosis in cHepC patients. Schiavon et al. reported a positive predictive value of 66% for fibrosis (METAVIR F2, F3 and F4) when APRI > 0.95 and negative predictive value of 93% when APRI < 0.40 in ESRD patients (17). In another study by the same group, APRI score was found to be superior to YKL-40 with regards to fibrosis in ESRD patients (18). In our study, we found APRI score as an independent predictor for both HAI and FS. Thus, APRI score might be a valuable score to predict FS in ESRD patients.

NLR and PLR are current markers of interest for systemic inflammation (11–13). Data is limited in patients with cHepC. In one study, NLR was proposed as related to viral load and as an independent marker to follow response to therapy, especially in genotype 2 patients (25). We could not confirm any association neither with HAI or FS. Further studies are needed to investigate the role of NLR and PLR in this population.

There are several limitations to our study. The study population is relatively low and this may have prevented some statistical analysis to reach a statistical significance. Also, the majority of the patients were genotype 1 cHepC patients and our results may not be valid for all genotypes. The biopsies were scored according to the modified ISAAC scoring system and therefore, other scoring systems may yield different results. Finally, the relatively low FS observed in the patients may have interfered with the analysis.

As a conclusion, YKL-40 may be a valuable marker to predict fibrosis in non-uremic patients with cHepC. Serum IgG and GGT levels, used frequently in routine practice, may be also good predictors for HAI. However, the role of inflammatory markers in the prediction of HAI and FS may be limited in ESRD patients. In this population, APRI score may provide useful information with regards to HAI and FS.

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CONFLICT OF INTEREST

Declare no conflict of interest.

INFORMED CONSENT

Informed consent was obtained from all individual participants included in the study.

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