

Viral Load Among the Sera of Iraqi Hepatitis C Virus Patients

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Summary:

Background:

Hepatitis C Virus is the main causative agent of hepatitis among blood transfused patients, in which most chronic cases result in liver carcinoma.

Materials & Methods:

Enzyme-Linked Immunosorbent Assay for HCV, FNF- and IL-12p40 estimation with Recombinant Immunoblotting Assay (RIBA) as a confirmatory test for HCV, have been applied for HCV detection in 80 HCV patients' samples in comparison with 30 samples for apparently healthy control, while viral load has been estimated using Branched-DNA (b-DNA) technique for 32 randomly selected positive cases for HCV. Liver function test has been applied for patients' sera in comparison with control.

Results & Conclusions:

This study reveals highly specificity & sensitivity of ELISA technique for HCV detection which results in 100% positivity by RIBA methods. Moreover, Viral load estimation shows that (71.9%) of HCV sera samples with viral load >615 IU/ml and only (28.1%) with viral load <615 IU/ml with highly significant difference between them ($P = 0.013$). Furthermore, there is a highly significant variations between liver function test in comparison with control group ($P < 0.01$), while SGPT is the only parameter which significantly affected by viral load ($P = 0.01$). Beside that, neither INF- nor IL-12p40 level has been affected by viral loads more or less 615IU/ml. It was concluded that ELISA technique is still the best accurate reliable method for viral detection and SGPT is a good marker for highly viral loaded samples.

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Introduction:

Viral hepatitis is a systemic disease primarily involving the liver and is caused by many viral agents which are etiologically, immunologically and epidemiologically distinct [1].

Hepatitis C was first recognized in early 1970, after the discovery of hepatitis A virus (HAV) and hepatitis B virus (HBV). It was noticed at that time that most cases of transfusion-associated hepatitis were not caused by either these viruses, this virus was called "non A non B" [2].

In 1989, the responsible virus for most transfusion-associated non A-non B hepatitis was identified through molecular biological techniques and Hepatitis C virus (HCV) was cloned [3, 4].

Hepatitis C is an enveloped virus, with a diameter of about 50-60 nm, positive sense, single stranded RNA virus [5, 6]. The HCV genotype is an intrinsic characteristic of the transmitted HCV strains(s) and does not change during the course of the infection. HCV genotypes from six clades or types (number from 1-6) and are themselves subdivided into number of subclades or subtypes identified by lower case letter (1a, 1b, 1c, ... etc) [7].

Probably as many as 70%-90% of infected people fail to clear the virus during the acute-phase of the disease and become chronic carrier [8, 9].

Chronic hepatitis C is highly heterogenous and many patients present with mild form of

liver disease. However, progression of liver fibrosis can be observed at long-term (>5-7 years) follow-up particularly in those cases with elevated and/or fluctuating transaminase level [10]. In 1990 it was become possible to screen for HCV infection with antibody test or direct amplification assays for the nucleic acid [i.e. reverse transcription polymerase chain reaction (RT-PCR)] HCV has been found to be the causative agent of most post-transfusion NANBH (PT-NANBH) [11].

A new test has recently been developed by Ortho-clinical diagnostics detect the HCV core protein (HCV core antigen) which is coded by the most conserved regions of the virus genome [12].

The assay has better sensitivity and permits the detection of HCV infection earlier than the HCV antibody screening tests and an average of only two days later than quantitative HCV-RNA detection in individual specimens. The performance of the assay correlates well with those of molecular HCV-RNA detection methods, but the lower level of detection (20,000 IU/ml) is significantly higher [13].

Materials and Methods:

Eighty Iraqi patients complained from hepatitis C were involved in this study during the period from November 2005 to August 2006, with ages ranged between (8-80) years. They were sequentially selected from cases

Results:

I. Some Clinical and Demographical Picture of Patients:

It is clear from Table 1 that the majority of patients are the males [55 out of 80] (i.e.

referred to the Hepatology and Gastroenterology Teaching Hospital in Baghdad at first presentation. Their diagnosis based upon the patients medical history, physical examination and laboratory tests which include biochemical tests (bilirubin (TSB) [Spin react, Spain], Alanine amino-transferase, (AIT), Asparate amino-transferase (AST), Alkaline Phosphatase (AIP) [Biomeriex, France; for both enzymes]; and virological testes include {Anti-HCV screening (ELISA, Rand ox, UK) and HCV confirmatory RIBA, Chiron, Irland} and detected HCV RNA by branched DNA assay (b-DNA), [Bayer Healthcare, USA].

From those patients, 60 cases were confirmed by using test by (RIBA) and 32 cases were measure for viral load by branched DNA (b-DNA) assay because no material available on all cases.

After these tests; determination of some serological markers include (IL-12p 40 and INF-) [Bio source, Europe; for both] in the sera of these patients have been performed.

Statistical Analysis:

All the results have been statistically analyzed for comparison of significance using:

- A. Binomial test.
- B. Krushal Wallis test.
- C. Matched paired t-test for repeated measurements.
- D. Pearson correlation coefficient (r) [14]

68.8%) rather than the females [25 out of 80 (i.e. 31.2%) with highly significant differences between both frequencies (P<0.001). The ratio between male to female was 2.2:1.

Table 1: Demographical Picture of HCV patients

<i>Age groups (years)</i>	<i>Patients' No.</i>	<i>Frequency</i>	<i>P value</i>
1-20	9	11.3	0.001
21-40	24	30	
41-60	33	41.3	
61-80	14	17.5	
Gender:	Patients' No.	Frequency	
Male	55	68.8	
Female	25	31.2	
Clinical Presentation	Patients' No.	Frequency	
Acute	24	30	
Chronic	47	58.8	
Unknown	9	11.3	
Total	80	100	

Moreover, Table 1 revealed the frequency of patient's distribution of among different age groups. According to this table, the majority of patients are at the age between 41-60 years (41.3%), while children elicit less frequency of infection (11.3%). There are highly significant differences between the incidences of the different age groups ($P < 0.01$). The chronicity of HCV is very important finding which determines the treatment plane of the disease. Table 4.2 reveals the frequency of different clinical presentations of the disease among 80 patients whom they were studied. It was observed that 47 patients had undergone chronic HCV infection (58.8%) and only 24

patients were at acute stage of the disease (30%). Only 9 patients were miss-diagnosed (11.3%) if the are at acute or chronic stage, it may be explained that they were in between or at the transitional stage of the disease. There was a highly significant differences between acute and chronic frequencies ($P < 0.001$).

II. Methods for HCV Detection:

There are several methods for HCV diagnosis. The first step is the screening method by using ELISA technique for detection of anti-HCV antibody. All the sera samples of the 80 hepatitis patients were found to be positive for this test.

Table 2: Distribution of Chronic HCV patients according to Anti-HCV Screen Test, RIBA Test and Viral Load by (b-DNA) assay (IU/ml) results

Anti-RNA HCV	RIBA* No.=60	b-DNA No.=32	No. of patients (%)	P value
No. of (+)ve = 80	60 (100%)	> 615 IU/ml	23 (71.9)	0.013#
No. of (-)ve = 0	0 (0.0%)	< 615 IU/ml	9 (28.1)	
Total = 80 (100)	60 (100)	32	32 (100)	

* out of 80 randomly selected 60 samples and from those 60 only 32 samples tested for viral load.
 # P value for comparison of difference between the frequencies of less and more than 615 IU/ml.

Another advanced method as a confirmative test such as Immuno-blotting (RIBA) has been applied. All these specimens gave positive results. These findings reflected highly sensitivity and specificity of ELISA technique detection of HCV infection as a screening accurate method while branched-DNA (b-DNA) assay method applied for studying HCV viral load. The last method applied on randomly selected samples from chronic HCV. It was observed that 23 out of 32 had > 615 IU/ml (71.9%) and only nine had viral load < 615 IU/ml (28.1%) with significant difference between them ($P = 0.013$) as shown in Table 4.3. All the control groups, samples gave negative results for all the above tests.

III. The Correlation Between Liver Function and Viral Load:

The correlation between liver function tests and viral load had been listed in table 3. This table revealed that there were significant differences in TSB, S.GPT and S.GOT for those with less or more than 615 IU/ml viral load in comparison with control group.

Serum GPT was observed to be the single marker elicits significant difference between samples with < 615 IU/ml viral load and those with > 615 IU/ml viral load. This means that S.GPT is an important inverse indicator for the viral load.

Table 3: Correlation of liver function with viral load among HCV patients and apparently healthy control

Studied Groups	Liver Function Tests			
	TSB (mmol/l)*	S.GPT (IU/ml)	S.GOT (IU/ml)	S.ALP (IU/ml)
< 615 IU/ml No.=9	128.83 ± 44.14	119.33 ± 74.27	48.00 ± 24.37	104.83 ± 44.03
> 615 IU/ml No.=23	130.94 ± 61.85	74.47 ± 37.71	55.12 ± 20.77	85.12 ± 41.18

Control Group No.=30	9.5 ± 1.73	9.25 ± 2.31	9.40 ± 2.95	66.20 ± 13.36
P value vs Control	0.001	0.001	0.001	0.013
P value < 615 vs > 615 IU/ml	0.917	0.011	0.35	0.198

* Mean ± SD [Standard Deviation]

IV. Correlation Between HCV Viral Load and Some Cytokines:

A. The Correlation between INF-γ and viral load:

The association between INF- and viral load had been listed in table 4.8. This table

showed that there was a highly significant association between the samples with viral load > 615 IU/ml and that for sample with < 615 IU/ml in comparison with control group.

Table 4: Association between INF- level and viral load among HCV patients

Studied Group	Number	Mean ± SD	P value
Control	30	1.585 ± 0.24	-
< 615 IU/ml	9	0.324	-
> 615 IU/ml	23	0.3322 ± 0.256	0.001
Total	62	<615 IU/ml vs. >615 IU/ml = 0.87	

SD = Standard Deviation

B. The Correlation between IL-12p40 and viral load:

The correlation between IL-12 level in the sera of HCV patients and the viral load

appeared in table 4.9. This table showed that there is no significant difference between viral load less and more than 615 IU/ml and the level of IL-12 (P=0.157).

Table 5: The association between IL-12p40 level and viral load among HCV patients

Studied Group	Number	Mean ± SD	P value
Control	30	22.934 ± 7.284	-
< 615 IU/ml	9	187.004 ± 179.329	0.007
> 615 IU/ml	23	103.416 ± 97.335	0.083
Total	62	< 615 vs. > 615 IU/ml = 0.157	

SD = Standard Deviation

Discussion:

Most studied denoted to the prevalence of HCV was among men rather than women which revealed that the male:female ratio is (2:1) [15-19]. Some observed an equal ratio of males:females (1:1) in Iraq [20, 21]. The explanation for these variations may be attributed to the difference in sample's size in addition to the different time of blood collection.

Concerning the effect of age on the incidence of HCV the result of the current study is comparable to that of the others which mentioned that there was high infectivity among Iraqi patients at age range between 27-60 years (65.9%) [17] as well as for the others which was between 40-60 years (70%) [15, 22]. On the other hand, aboard one declared that the highly infected age range between 13-

82 years (62%) [14]. This may be due to so many drug abusers among these communities in comparison with Iraqis. Moreover many bad costumes are prevalent among teenagers such as tattooing [23], besides intra-familial sharing razors, tooth brushes which enhances the disease transmission [24].

The current study revealed that (58.8%) of HCV cases are chronic. This frequency is lower than that of Europeans (70%) [25], though it is higher than the last report for WHO (20%) [26]. While the local studies denoted to similar frequency (58.9%) [27, 28]. These variations may be attributed to the geographical and environmental differences between Iraq and other countries.

Studies referred to an increase in the viral load of patients' sera [29], though it was observed that only (25%) of patients showed

High viral load in comparison with this study (71.9%) revealed High viremia. This variation related to the genotype of HCV particularly, genotype 3 HCV which observed to elicit lower viremia. Furthermore the difference in methods used for viral load estimation added another reason for this variation between this study which detected viral load by Taqman PCR and the other one [29, 30]. Moreover the cut-off value may vary according to the method and company [31-32]. Additionally, viral load was found to be varied according to the duration of the disease; highest viremia was achieved within 4-12 weeks of infection [33]. Most of Iraqi patients, who involved in this study had chronic infection and had HCV infection before months.

Although, there was no correlation between viral load and disease severity since the cytopathic effect attributed to the immune mechanisms rather than the viral pathogenicity. These findings are analogue to some extent with that of other studies which denoted that no correlation between viral load and S.GPT, S.GOT levels [34-35]. While it was well-inversely associated with the cytotoxic T-lymphocyte [36].

There was no significant difference between the levels of INF- according to the viral load of the patients' specimens. These findings were disagree to some extent with another study which referred to an inversely correlation between viral load and INF- in chronic cases of HCV [37]. Perhaps the reason was related to the disease duration and small sample size.

Obviously, serum IL-12p40 concentration inversely related to the viral load. This fact was confirmed by others [38].

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