# THE ROLE OF BCL-2 ONCOPROTEIN IN HEPATITIS C-RELATED HEPATOCELLULAR CARCINOMA

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## ABSTRACT

Viral hepatitis has emerged as a major public health problem worldwide. There is increasing evidence that, several viral oncogenes such as core protein C virus (HCV) is playing a direct role in neoplastic transformation. The tumor development requires both increased proliferation and inhibition of apoptosis. There is increasing experimental evidence to suggest that bcl-2 plays a critical role in blocking apoptosis in cancer cells. In the present study we tried to explore the role of bcl-2 in development of viral hepatitis into hepatocellular carcinoma (HCC). Out of 105 participants; 50 patients were HCV positive infected patients, 30 HCC patients either infected with HCV (HCC/HCV) or not (HCC) and 25 healthy volunteers were also included. We determined the serum level of bcl-2, AFP and HCV-RNA viral load in patients and compared them with the other clinicopathological parameters. Increased serum bcl-2 level was showed in HCC/HCV infected patients more than HCV and controls (P=0.001) and more than HCC patients (P=0.018). A significant positive correlation between AFP and total load of viremia was elucidated (r=0.345, P=0.014) in the HCV infected patients who didn't develop HCC. By analyzing the diagnostic performance of the studied parameters among the different groups. Bcl-2 was found to be a significant prognostic marker for the progression of HCC, as it represented two or more diagnostic values (Sensitivity 80 %, 70% and specificity 60%, 70% respectively).and (OR=6.0, 95% CI=1.74-20.59& OR=5.4 & 95% CI=1.75-16.88).These results make bcl-2 a candidate marker for HCC progression following HCV infection in Egyptian patients.

Key words: Hepatocellular carcinoma (HCC) hepatitis C virus (HCV), bcl-2.

## INTRODUCTION

Hepatocellular carcinoma (HCC) is a primary cancer of the liver that is predominant in developing countries, with nearly 600,000 deaths each year worldwide (El-Serag and Rudolph 2007). HCC normally develops as a consequence of underlying liver disease and is most often associated with cirrhosis (Blum 2005). Hepatitis C virus (HCV) is now the major cause of chronic hepatitis, liver cirrhosis and hepatocellular carcinoma (HCC) in the western world. In addition, end-stage liver disease due to chronic hepatitis C is the leading indication to liver transplantation (Williams 2006). It is estimated that 120-180 million people worldwide are infected with HCV (Shepard et al., 2005), which corresponds to roughly 4 times the number of individuals infected with HIV and about half the number of persons infected with the hepatitis B virus (HBV). The seroprevalence rate is 1-2% in Western Europe and North America, 3-4% in some Mediterranean and Asian countries and up to 20% in Egypt and parts of Central Africa. (Stramer et al., 2004) Chronic hepatitis C virus (HCV) infection is characterized by inflammatory lesions in the liver, often accompanied by intrahepatic lipid accumulation (steatosis) and

progressive fibrosis of variable degrees, and longterm progression to cirrhosis and hepatocellular carcinoma (HCC) (Moradpour and Blum, 2005; Zoulim et al., 2003). HCC incidence has increased sharply over recent decades and has been attributed to chronic HCV infection. Chronic HCV infection, therefore, is a major risk factor for HCC development. and HCV infection increases the risk for HCC development by an estimated 17fold compared to healthy individuals. (Donato et al., 2002; Imazeki et al., 2003; El-Serag 2002; Sun et al., 2003) HCV is the only RNA virus with a predominantly cytoplasmic life cycle. (Moradpour et al., 2007; Lindenbach et al., 2007) All potentially pro-oncogenic events are therefore likely to be restricted to the cytoplasm, suggesting indirect mechanisms of hepatocarcinogenesis. HCV While infection leads to chronic inflammation, steatosis, fibrosis and oxidative DNA damage, several HCV proteins have been shown to have direct oncogenic effects and to upregulate mitogenesis (McGivern et al., 2009; Koike 2007).

The anti-apoptotic factor B-cell leukemia /lymphoma 2 Bcl-2, is an oncogene in follicular lymphoma which is frequently linked to an immunoglobulin locus by the chromosome translocation (McGivern et al., 2009; Vaux et al., 1988; Adams et al., 1998; Gross, et al., 1999). It was the first example of an oncogene that inhibits cell death rather than promoting proliferation. B cells transfected with bcl-2 were shown to be rendered resistant towards apoptosis induced by IL-3 withdrawal: for the first time it was shown that the pathway toward tumorigenesis depends not only on the ability to escape growth control but also depends on the ability to prevent apoptosis (16). The antiapoptotic bcl-2 family members are critical regulators of the mitochondrial apoptotic pathway that are not regulated by the cell surface receptor pathway or the "extrinsic pathway" (Adams et al., 1998; Gross, et al., 1999; Strasser et al., 1999).

For the early detection of sub clinical HCC by AFP measurement and/or ultra sonography (us) has been proposed since the 1980s. Some evidence also indicates that AFP secretion stimulates human hepatoma tumor cell growth and proliferation (Wang et al., 1999). Because the majority of cells losing the ability to silence AFP have a disease phenotype, it is likely that the mechanism by which AFP is re-activated is one way that HBV contributes to the development and progression of HCC.

AFP is a fairly specific but insensitive marker for HCC. Sensitivity of HCC detection by blood markers is improved by combining various other markers with AFP.

The cross relation between HCV proteins and host cell regulatory proteins was extensively studied, but still unable to clear the role of HCV in HCC initiation and/or progression. In this concern, the aim of the present work was to study the interference of the HCV infection in HCC patients with the serological markers trying to find out if there is any correlation with those markers in HCV infected patients and developing malignancy.

## PATIENTS AND METHODS

Patients and grouping: This study was carried out on 105 participants. Eighty patients were recruited between January 2008 and March 2010 from Tropical medicine Department of Alexandria Main University Hospital, Egypt. Twenty five healthy volunteers were also included. This study was approved by the research review committee of the Alexandria faculty of Medicine. And all the studied subjects gave their informed consent. The studied subjects were classified into 3 main groups:- Group I included 50 HCV Positive patients 26 males and 24 females, their ages were ranged from 20 to 71 years old. Group II included 30 HCC Patients who were further categorized into 2 subgroups (A&B), subgroup A, included 20 HCC patients infected with HCV 17 males and 3 females, their ages were ranged from 42 to 69 years old, and subgroup B, included 10

HCC patients who were HCV free 7 males and 3 females, their ages were ranged from 36 to 67 years old. In subgroup B, HCC developed due to Schistosomal infection, alcohol abuse or with unknown etiology rather than HCV infection. All subjects included in the study were negative for HBs Ag.

Group III is the Control group which included 25 healthy volunteers with matched age and sex were chosen carefully, the subjects of this group with no evidence of liver disease and who were negative for HCV antibodies. RT-PCR (Reverse transcription polymerase chain reaction) was performed for the first two groups to confirm HCV infection in group I and subgroup A and absence of HCV infection in subgroup B.

# Methods:

HCC patients were diagnosed by liver fine needle aspiration (Biopsy) whenever possible to confirm the presence of malignant tissues, or by ultrasonography and rectal snip.

All patients were subjected to:

A - Thorough history taking and clinical examination.

B - Laboratory investigations including:

1 - Liver function tests (Total Bilirubin, Albumin, Alanine amino transferase (ALT),

Aspartate amino transferase (AST), alkaline phosphatase and prothrombin time).

2 - Anti HCV antibodies (third generation) by microparticle enzyme immunoassay

(MEIA) technology on Axsym was performed for all participants. Positive cases Followed by RT-PCR (Reverse transcription polymerase chain reaction) for estimation of total viremia.

3 - Quantitative determination of human serum bcl2.

4 - Quantitative determination of alpha-fetoprotein (AFP).

# Samples collection:

Venous blood were withdrawn from all subjects, two ml was in vacutainer trisodium citrate tube for determination of prothrombin time (PT), and the rest in a plain vacutainer tube, sample was allowed to clot for 30 minutes and then centrifuged at 3000 rpm. For 5 minutes, serum was stored in aliquots at -20 °C for (Total bilirubin, Albumin, Alkaline phosphatase, ALT, AST, AFP and Bcl-2) determination.

#### Quantitative determination of α1-fetoprotein (AFP) using chemiluminescence technique on Cobas e 411 from Roche (Germany): (21)

Sandwich principle: 1<sup>st</sup> incubation: 10 µl of sample, a biotinylated monoclonal AFP-specific antibody, and monoclonal AFP-specific antibody labeled with a ruthenium complex (Tris(2.2bipyridyl) ruthenium (II). 2<sup>nd</sup> incubation: After addition of streptavidin-coated microparticles, the complex becomes bound to the solid phase via interaction of biotin and streptavidin. The reaction mixture was aspirated into the measuring cell where the electrode. Unbound substances were then removed with procell. Application of a voltage to the electrode then induces chemiluminescent emission which was measured by а photomultiplier. Results were determined via a calibration curve which was instrument-specificity generated by 2-point calibration and a master curve provided via the reagent barcode.

### Quantitative polymerase chain reaction (PCR) for HCV RNA (total viremia) by Real Time PCR (Taq Man Technology) by Step-One (Applied Biosystems): (22)

The TagMan probe contains a fluorescent reporter dye at the 5`end of the probe and at the 3`end of the probe. When the probe was intact, the proximity of the reporter dye to the guencher dye suppresses the reporter fluorescence. Probe cleavage during the PCR reaction spatially separates the reporter dye from the quencher moiety and allows detection of the reporter dye fluorescence. Pathogen diagnosis by the polymerase chain reaction (PCR) was based on the amplification of specific regions of the pathogen genome. In real time PCR the amplified product was detected via fluorescent dyes. Monitoring the fluorescence intensities during the PCR run (i.e. in real time) allows the detection and quantitation of the accumulating product was allowed without having to re-open the reaction tubes after the run.

# Bcl-2 Immunosorbent Assay: (23)

An anti-human bcl-2 Coating antibody is adsorbed onto microwells. Human bcl-2 present in the sample or standard binds to antibodies adsorbed to the microwells. A biotin-conjugated anti-human bcl-2 antibody was added and binds to human bcl-2.Captured by the first antibody. Following incubation unbound biotin-conjugated anti-human bcl-2 antibody was removed during a wash step. Streptavidin- HRP is added and binds to the biotin- conjugated anti-human Bcl-2 antibody. Following incubation unbound Streptavidin-HRP was removed during a wash step, and substrate solution reactive with HRP was added to the wells.(Bender Med Systems, Austria).

### Statistical analysis:

Statistical analysis was carried out using SPSS version 15. Values were expressed as mean $\pm$ SD Kruskal-Waills and Mann-Whitney tests were used. Pearson's correlation was done between different parameters. ROC curve analysis was done and X<sup>2</sup> was obtained by Chi-Square test. P is considered significant when < 0.05

### **RESULTS AND DISCUSSIONS**

One hundred and five individuals were enrolled in the study. With age ranged from 20-71 years. They were divided into 50 patients with HCV infection (54% males, 46% females) 30 patients with hepatocellular carcinoma associated with HCV infection (HCC/HCV) (A) (n=20) or non associated (HCC) (B) (n=10)(85% males, 15% females and 70% males, 30% females respectively).Twenty five healthy individuals as a control group (68% males, 32% females).

Parameters	GI	C	G III Control	Test	Over	
	HCV(n=50) m±SD	A(HCC/HCV) (n=20) m±SD	B(HCC) (n=10) m±SD	(n=25) m±SD	Value (F)	all P value
Age(yr)	51.02±12.85	56.70±7.6	54.80±9.46	53.68±13.81	1.184 <sup>b</sup>	0.320
Bcl- 2(ng/ml)	3.77±5.18	13.74±15.61	2.41±1.15	2.07±.60	15.55ª	0.001
AFP(ng/ml)	8.56±12.79	3671.04±5044.1	3790.40±5262.81	2.40±.73	75.24ª	<0.001
viremia (copies/ml)	530±860	1367±778				<0.001
AST (U/L)	58.62±45.66	102.10±49.48	107.2±50.64	28.56±6.7	44.16 <sup>a</sup>	<0.001
ALT(U/L)	58.12±43.18	60.80±24.01	69.00±32.02	26.48±7.42	29.82 <sup>a</sup>	<0.001
Albumin (g/dl)	3.57±.49	2.97±.54	3.14±.42	4.05±.40	21.76 <sup>b</sup>	<0.001
PT(Sec.)	13.06±1.51	16.00±3.13	14.36±2.19	11.38±.33	24.69 <sup>b</sup>	<0.001
Bilirubin (mg/dl)	1.04±.36	2.41±2.99	4.87±5.60	0.60±.07	55.43ª	<0.001
Alk. Phosph. (U/L)	238.66±85.92	477.15±262.00	454.4±126.83	172.68±59.29	55.79ª	<0.001

Table 1: Serum Bcl2, AFP, Total viremia levels and liver function tests among different studied groups.

(a) Significant difference between 2 groups.

(b) Significant difference between more than 2 groups.

Serum levels of all serological parameters were represented in figures 1, 2.

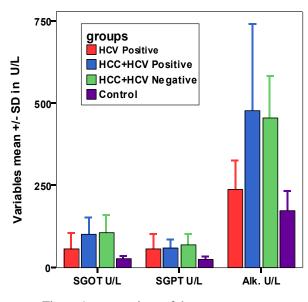
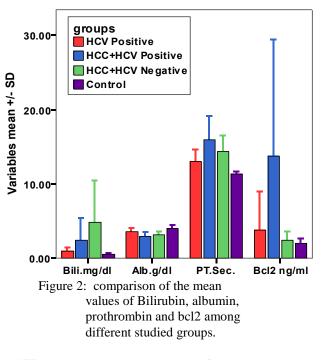


Figure 1: comparison of the mean values of SGOT ,SGPT and alkline phosphatase among different studied groups.

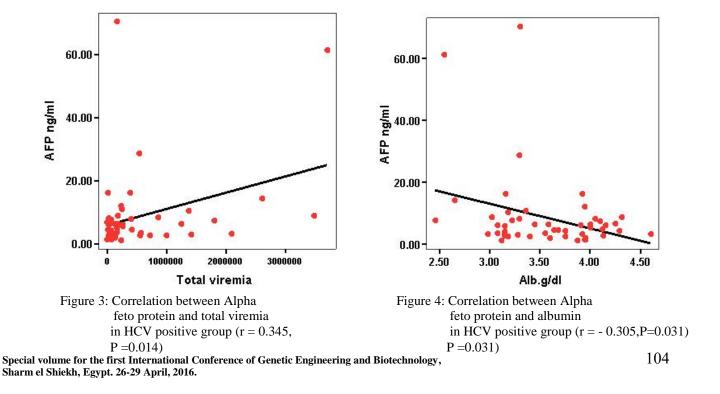
**Bcl-2** level was highly significant elevated in HCC/HCV than HCV infected patients and control (P=0.001) besides it's significantly elevated in comparison to HCC patients (P=0.018).

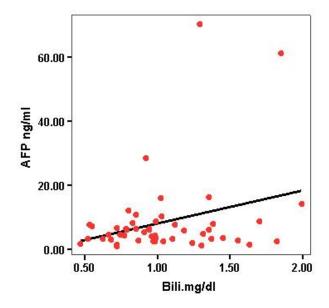
The level was non significantly elevated in HCV and HCC compared to control or between each other. (Table 1)

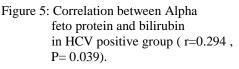


**AFP** and total viremia were significantly correlated in HCV infected patients (P=0.014).

AFP was also significantly correlated with albumin and bilirubin in the same group (P=0.031, P=0.059) respectively. As shown in figures 3, 4, 5.







While in HCC/HCV patients AFP was significantly correlated to alkaline phosphatase enzyme and bilirubin as represented in figures 6, 7.

**Total viremia** was in a negative significant correlation with prothrombin time in HCC/HCV patients (P=0.020) (figure 8).

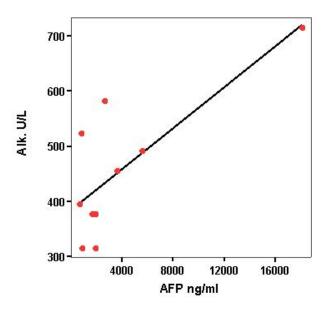
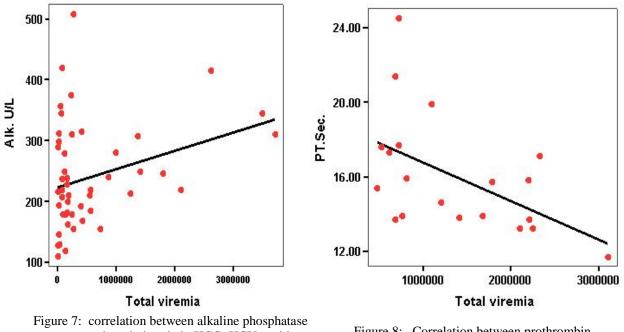
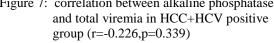
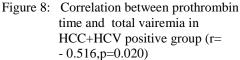


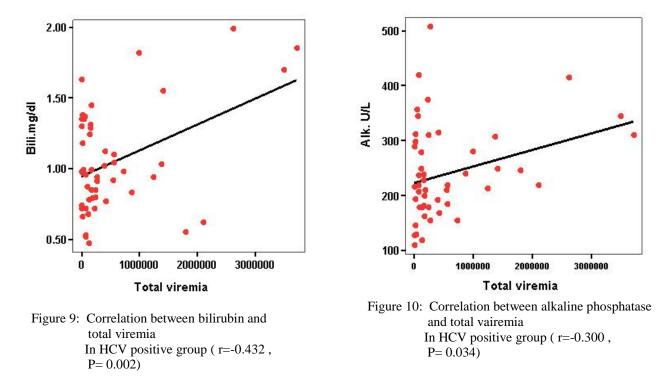
Figure 6: Correlation between Alpha feto protein and alkaline phosphatase in HCC+HCV negative group (r=0.769 P= 0.009).

While it was significantly correlated with bilirubin and alkaline phosphatase in HCV infected patients (P=0.002, P=0.034) (figures 9, 10). While bcl2, was not significantly correlated with other parameters in all patients groups.









Parameters	BCL-2		AFP		Total viremia	
	r	Р	r	Р	r	Р
BCL-2	1	-	0.084	0.562	-0.047	0.747
AFP	0.084	0.562	1	-	0.345	0.014
Total viremia	-0.047	0.747	0.345	0.014	1	-
Age	-0.005	0.974	0.150	0.297	0.198	0.168
AST	0.058	0.691	0.096	0.506	0.063	0.666
ALT	0.018	0.900	-0.072	0.620	0.128	0.374
Albumin	-0.011	0.940	-0.0305	0.031	-0.078	0.590
PT	-0.048	0.742	0.033	0.823	0.195	0.175
Bilirubin	0.217	0.129	0.294	0.039	0.432	0.002
Alk. phosphatase	-0.150	0.298	0.089	0.538	-0.300	0.034

Table 3: Correlation of Bcl-2, AFP levels and total viremia in HCC/HCV patients.

Parameters	BCL-2		AFP		Total viremia	
	r	Р	r	Р	r	Р
BCL-2	1	-	-0.253	0.282	-0.137	0.564
AFP	-0.253	0.282	1	-	0.057	0.810
Total viremia	-0.137	0.564	0.057	0.810	1	-
Age	0.289	0.217	-0.328	0.159	-0.305	0.191
AST	0.232	0.325	-0.135	0.571	0.007	0.976
ALT	0.094	0.693	0.095	0.690	-0.161	0.497
Albumin	0.258	0.272	0.255	0.278	0.258	0.272
PT	-0.150	0.527	-0.258	0.273	-0.516	0.020
Bilirubin	0.111	0.641	-0.250	0.289	0.253	0.281
Alk. phosphatase	0.234	0.321	0.433	0.056	-0.226	0.339

Parameters	BCL	-2	AFP		
	r	Р	r	Р	
BCL-2	1	-	0.422	0.225	
AFP	0.422	0.225	1	-	
Age	0.534	0.112	-0.005	0.990	
AST	-0.081	0.823	0.054	0.882	
ALT	-0.120	0.742	0.225	0.531	
Albumin	-0.031	0.933	-0.333	0.347	
РТ	0.434	0.210	0.240	0.504	
Bilirubin	0.446	0.196	0.911	0.000	
Alk. phosphatase	0.201	0.578	0.769	0.009	

Table 4 : Correlation of Bcl-2, AFP levels and total viremia in HCC patients

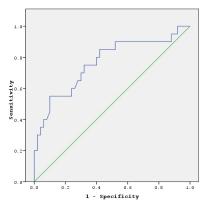
Receiver Operating Characteristic (ROC) Curve analysis (figure 11):

ROC curve analysis was designed for BCL-2 in HCV infected patients against HCC/HCV group of patients Roc curve was performed as cut off for disease progression. The area under the curve and the best cut off values (c.o.v.) and youden index of bcl-2 levels were obtained from the sensitivity and specificity % in progression of malignancy (24).

The ROC curve analysis showed that BCL-2 presented a significant area under the curve of

0.765 and achieved sensitivity (80%, 75%, 70%) and specificity (60%, 68%, 70%) respectively. Table 5: represented the risk estimate at different relative bcl-2 cut off values (2.19, 2.67, 2.76), after calculating the positive and negative predictive values corresponding to each cut off.

A significantly elevated relative risk by 5.4 - 6.3 folds more for developing malignancy after HCV infection



Biochemical parameter	ROC Area under the curve	Sig	Youden's index	Sensitivity	Specificity	Cut off Value		
Bcl2	0.765	0.001	0.40	80%	60%	<u>&gt;</u>		
						2.19		
			0.43	75%	68%	<u>&gt;</u> 2.67		
			0.40	70%	70%	<u>&gt;</u> 2.76		
Youden index = (Specificity + Sensitivity) -1								

Table 5: Frequency distribution and risk estimate at different bcl-2 cut off values in HCV infected patients and HCC/HCV patients.

Group	BCL-2	+ve	-ve	X²	OR	95% CI	Р
	cut off	predictive	predictive				
	value	value	value				
HCV		20 (40%)	30 (60%)				
	≥ 2.19			9.150	6.000	1.748-20.591	0.002
HCC/HCV		16 (80%)	40 (20%)				
HCV		16 (32%)	34 (68%)				
	≥2.67			10.7	6.375	1.971-20.615	0.001
HCC/HCV		15 (75%)	5 (25%)				
HCV		15 (30%)	35 (70%)				
	≥2.76			9.420	5.444	1.756-16.881	0.002
HCC/HCV		14 (70%)	6 (30%)				

 $X^2$ : obtained by chi-square test. P: is considered significant < 0.05 OR: odds ratio ,95%CI: 95%confidance interval

HCV-associated hepatocellular carcinoma is a common neoplasm in Egypt. Since tumor development requires both increased proliferation and inhibition of apoptosis; HCC was exclusively related to HCV infection. bcl-2 gene is known to provide a survival advantage to rapidly proliferating cells. Physiologically bcl-2 protein blocks the apoptotic process by inhibiting the release of cytochrome C from mitochondria whereas it locates at the cytoplasmic surface of mitochondrial membrane, nuclear membrane and endoplasmic reticulum (Brown 1997).

Chung Y L and his team (Chung et al., 2003) stated that HCV mimics the effect of bcl-2 and maintains the growth of HCC, where the non-structural 5A (NS5A) protein of hepatitis C virus region contains three bcl-2 homology domains (BH3,BH1,BH2) which can interact with Bax and lead to the inhibition of apoptosis.

The present work aimed to assess whether HCV infection in HCC patients affects the levels of bcl-2 in comparison to non-infected HCC patients and HCV patients who didn't develop HCC the data revealed that bcl-2 level is influenced by HCV infection, as it is up regulated whenever HCV infection is developed into HCC. The diagnostic performance of bcl-2 estimated an increased risk by 5-6 folds more for developing HCC after HCV infection. Bcl-2 appeared to be over expressed in patients with HCV/HCC patients compared to those with chronic hepatitis and HCC-non associated HCV. It suggests that bcl-2 may have

a role in development or progression of HCC in patients with chronic hepatitis C. It was not correlated with total viremia or AFP or ALT levels in all patient groups either infected with HCV or not. Our data indicated that total viremia is related to AFP and bilirubin serum level in patients with chronic hepatitis C. This finding correlates with the previous reports (Sultan et al., 2006) which stated that high viral load associated frequently with HCC in patients with HCV. A study by Sultan. et al. reported that although there is a significant increase in the bcl-2 level in HCC patients associated with HCV infection compared to HCC patients alone, the results suggest that HCV might not be the direct cause of bcl-2 elevation, since not all HCV-infected patients had increased level of bcl-2. Previous reports have indicated that the expression of bcl-2 mRNA was up regulated in HCC, but low level of bcl-2 protein was detected (Ravazoula et al., 2002).

A contradictory study by Vail and his coworkers (Vail et al., 2001) postulated that bcl-2 does not play a role in the progression of HCC in patients with progressive tumors (GII and G III). While, another study (Chung et al., 2003) found that in HCC patients with GI a significant relation ship between bcl-2 protein in HCV infected and HCVfree patients. It is speculated that HCV may exert an anti-apoptotic effect and thus enhance tumorigenesis. A previous study (Ali et al., 2004) revealed a positive correlation between bcl-2 and AFP in HCC patient infected with HCV concluding that the increased level of bcl-2 in HCC may be involved in hepatocarcinogenesis.

A recent study (Alenzi et al., 2010) was done on Egyptian patients with chronic HCV infection either developed HCC or not. As they evaluated the expression of bcl-2 by ELISA, Their results were in some way in a good agreement with ours, as bcl-2 level increased in HCV patients who developed HCC. They collectively concluded that Apoptosis is dysregulated and proliferation activity perturbed. In the present study, bcl-2 protein expression showed a dramatical highly significant increase in HCV patients who developed HCC, in comparison to other groups (P=0.001). Over expression of bcl-2 gene could lead to apoptosis inhibition and enable cell survival (Anatol et al., 2005).

A study by Hamazaki and co-workers (Hamazaki et al., 1995) in Japan was done to examine the expression of Fas antigen and bcl-2 protein in normal liver, diseased liver (chronic hepatitis or liver cirrhosis) and HCC. They found a difference in bcl-2 expression among the groups, but it was not significant. They suggested the induction of apoptosis may be a possible therapy against HCC. The conflict here is that in the present study, the Over-expression of bcl-2 in HCV patients who developed HCC was much higher than the bcl-2 up regulation in HCC patients without HCV infection (almost the same like the characteristics of HCC group in Hamazaki K. et al. (Hamazaki et al., 1995). This provided us to use HCV infection and bcl-2 related expression as a control arm to predict developing HCC.

The diagnostic performance of the Roc curve, in the present study, by examining the different studied parameters; revealed that bcl-2 was found to be a significant prognostic marker for the progression of HCC associated with HCV infection (sensitivity 70-80 % and specificity 68-70 %). This finding makes bcl-2 beside AFP; are good markers for detecting the progression of hepatocellular carcinoma after HCV infection.

Conclusions: The results in the present study predict a significant role of HCV infection on the circulating bcl-2 protein in both HCV and HCC/HCV patients. Bcl-2 could be a candidate marker for HCC progression following HCV infection. Bcl-2 determination may play a significant role in diagnosis, monitoring therapy, and evaluating malignant progression in HCV infected patients.

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