Cardiac Involvement of Hepatitis B and C Virus Infection

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Abstract

Hepatitis B virus (HBV) and Hepatitis C virus (HCV) are the cause of many different forms of heart disease worldwide and yet few cardiologists are aware of it as an etiology of heart disease, or its treatment. The burden of HBV and HCV derived heart diseases are global, with a higher prevalence in Asia, Africa, and low- and middle-income countries. The study showed that in more than 10% of Japanese patients, their cardiomyopathies are associated with HCV infection. More recently, it is found in the USA that up to 15% of patients with heart failure with myocarditis have been associated HCV infection. In contrast, in China 79% of patients with hepatocellular cancer and 37% of hepatitis C patients have heart disease, as detected by measuring a proven and sensitive biomarker of heart disease, NT-proBNP. In Pakistan, 17% of hepatitis C patients have heart diseases, as measured by this metric (NT-proBNP). Based on these data, 3% of 6.6 billion (198 million) persons worldwide are infected with HCV and 17–37% (34–73 million) persons are suffering from HCV-derived heart diseases. These figures may be comparable to the number of patients with hepatitis C. HCV infection causes only hepatitis in some patients, only heart diseases in some patients, and both hepatitis and heart diseases in other patients. A global network is required to establish methods to detect heart diseases caused by infectious agents. Other goals for the network are the expansion of preventive and therapeutic programs in under privileged countries.

Keywords : Cardiomyopathy; Heart failure; Myocarditis; Virus; Hepatitis B, Hepatitis C

Introduction:

Congestive Heart failure is one of the common clinical conditions encountered in medical practice. Heart failure is rapidly evolving to be a worldwide epidemic. Although most of the adverse outcomes related to cardiovascular diseases have been improving, the only exception is heart failure. As survival for acute cardiac catastrophic events such as myocardial infarction and sudden deaths improves, increasing number of patients suffer from heart failure. There are about 23 million people with heart failure in the world, 5 million in the United States, and 550 thousand new cases are added in 2003^{1} . Meta analysis of 21 studies (n = 37,791) with mild to moderate heart failure showed the prevalence of ischemic heart failure is 24-83% (average 64%)². In contrast, dilated cardiomyopathy is considered 63% of all causes of heart failure. In patients with severe heart failure who were candidates for heart transplantation, about one half had ischemic heart failure and one-half had dilated cardiomyopathy in Western countries. Therefore, dilated cardiomyopathy is the major cause of severe heart failure.

New classification of cardiomyopathies

In 1968, the World Health Organization (WHO) defined cardiomyopathy as a heart muscle disease of unknown etiologies in which the dominant feature is cardiomegaly and cardiac failure; excluded are myocardial dysfunction due to coronary, systemic or pulmonary vascular disease. Distinction was made between secondary cardiomyopathies, in which the heart muscle disease was associated with a known single cause, systemic disease or syndrome, and primary cardiomyopathy, in which the heart muscle disease was of unknown etiology. In 1980, the WHO, jointly with the International Society and Federation of Cardiology, chose to limit the term Cardiomyopathy to heart muscle disease of unknown etiology. The updated WHO definition in 1995 was "diseases of myocardium associated with cardiac dysfunction" and was included for the first time newly recognized Arrhythmogenic Right Ventricular Cardiomyopathy / Dysplasia (ARVC/D) and primary restrictive cardiomyopathy³⁻⁷.

In 2006, the AHA expert consensus panel proposed the definition: Cardiomyopathies are a heterogeneous group

of diseases of the myocardium associated with mechanical and/or electrical dysfunction that usually but not invariably exhibit inappropriate ventricular hypertrophy or dilatation and are due to a variety of causes that frequently genetic⁸. Cardiomyopathies thus are confined to the heart or are part of a generalized systemic disorder, often leading to cardiovascular death or progressive heart failure with related disability. Cardiomyopathies were divided into two major groups based on predominant organ involvement. Primary cardiomyopathies are genetic, nongenetic, acquired and those solely or predominantly confined to heart muscle and are relatively few in number. Secondary cardiomyopathies show pathological myocardial involvement as part of a large number of generalized systemic (multi-organ) disorders. The panel recommended that cardiomyopathies can be most effectively classified as primary : genetic; mixed (genetic and nongenetic), acquired; and secondary⁸ [Table-1]. Recently, the European Society of Cardiology working group on myocardial and pericardial diseases defined, "Cardiomyopathy as a myocardial disorder in which the heart muscle is structurally and functionally abnormal, in the absence of coronary artery disease, hypertension, valvular diseases, and congenital heart disease." Cardiomyopathies were grouped into specific morphological and functional phenotypes; each phenotype was then sub-classified into familial and non-familial forms⁹ [Table-1]. About one-half of hypertrophic Cardiomyopathy occurs familially, and one-half of those patients are associated with gene mutations. Confusion arises when in some cases of nonfamilial hypertrophic Cardiomyopathy, gene mutation are not seen, on the other hand some of the dilated Cardiomyopathy showed gene mutation of hypertrophic Cardiomyopathy. Although myocarditis is believed to be the major cause of dilated cardiomyopathy, myocarditis is often associated with hypertrophic cardiomyopathy which is caused by a viral infection such as hepatitis C virus^{10–13}, Hepatitis B & C virus infection are important causes of hepatitis both in developed & developing country. In Bangladesh Islam MN, Islam KM, Islam N et al, reported HBs positive in 27.2% of adult & 15.4% of children with acute hepatitis, 60% of post transfusion hepatitis patient & 65.5% of doctor. Selimur Rahman et al reported 31.25% cases of acute hepatitis, 76.3% of chronic hepatitis. In a study Hasan Ashraf etal among 1997 participants 452 (22.6%) for anti-HBc and 116 (5.8%) for both HBsAg and anti HBc or by cardiac sarcoidosis as discussed below¹⁴. Therefore, one etiology is not consistent with one phenotype of cardiomyopathy. Because the therapy should be based on the etiology, there will be proposed a new classification of cardiomyopathies (Table-2). For example, when compared with the WHO classification, dilated cardiomyopathy is classified into infectious, genetic, and so forth for etiology, left or right ventricle for anatomy, and systolic dysfunction for physiology. Hypertrophic Cardiomyopathy with obstruction is classified into genetic, infectious etc for etiology, septal hypertrophy for anatomy, and diastolic failure for physiology¹⁰⁻¹⁴.

 Table-I

 Classification or grouping of Cardiomyopathy ^{8, 9}

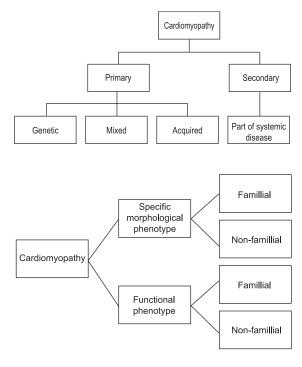


 Table-II

 A new classification of cardiomyopathies^{8, 9}

I.	Etiological classification	A.	Genetic
	-		B. Infectious
			C. Nutritional
			D. Unknown
II. Anatomical (structural)		А.	Dilated
classification			a. LV
			b. RV
		В.	Hypertrophic
			a. Septum
			b. Diffuse
			c. Free wall
			d. Apex
III.	Physiological (mechanical)	А.	Systolic failure/
	classification		dysfunction
		В.	Diastolic failure/
			dysfunction
		C.	Both
IV.	Electrical classification		Ion channel disorders
		В.	Conduction system
			disease
		C.	Others

WHO/ISFC classification Proposed c	lassification ^{8,9}
DCM (with viral infection)	I B, II A a, III A
DCM (with gene mutation)	IA, II A a, III A
HCM (with gene mutation outflow obstruction)	I A, II B a, III B
Apical HCM (with unknown cause)	I D, II B d, III B
Apical HCM (with HCV infection)	I B, II B d, III B
ARVC/D (with gene mutation)	IA, II A b, III B
ARVC/D (with HCV infection)	I B, II A b, III B
Long QT syndrome	I A, IV A

Comparison of classifications of cardiomyopathies WHO/ISFC classification Proposed classification ^{8,9}

Role of viruses in the pathogenesis of cardiomyopathies Myocarditis is one of the cardiological problem encounter in clinical practice, it is caused by many viruses and also it may represent as a part of many systemic diseases. Myocarditis is thought to be most commonly caused by enteroviruses, particularly coxsackievirus B. However, in many cases, when myocarditis has been diagnosed on the basis of clinical characteristics, no definite confirmation of viral origin is obtained, despite extensive laboratory investigations. The evidence is often only circumstantial, and a direct, conclusive proof of cardiac involvement is not available. However, accumulating evidence links viral myocarditis with the eventual development of dilated cardiomyopathy¹⁸⁻²⁰. Affection of the heart is manifested on the variety of pathological lesions e.g. when myocardial necrosis occurs diffusely, congestive heart failure develops and later dilated cardiomyopathy. If myocardial lesions are localized, a ventricular aneurysm may form. When complicated with arrhythmias, myocarditis presents as arrhythmogenic right ventricular Cardiomyopathy²¹. When myocardial necrosis is localized to the subendocardium, restrictive Cardiomyopathy may develop²¹. While it has not been established that hypertrophic cardiomyopathy may be a complication of viral myocarditis, asymmetrical septal hypertrophy has in fact, sometimes been observed in patients with myocarditis²². Hepatitis B antibody was detected in 8.6% of dilated cardiomyopathy and 14.6% of hypertrophic cardiomyopathy patients. Hepatitis C antibody was present in 6.7% of dilated cardiomyopathy and 9.5% of hypertrophic cardiomyopathy patients¹⁷. Recently the importance of hepatitis C virus (HCV) has been noted in patients with hypertrophic cardiomyopathy, dilated cardiomyopathy, and myocarditis^{10–13, 23, 24}.

Burden of cardiomyopathies: incidence and prevalence

There are few epidemiologic data of myocarditis. The Japanese Scientists conducted nationwide epidemiological surveys of cardiomyopathies in Japan^{14–16}. Disorders surveyed included idiopathic dilated Cardiomyopathy (DCM), hypertrophic cardiomyopathy (HCM), restrictive cardiomyopathy (RCM), ARVC/D, mitochondrial disease, Fabry's disease of the heart, and prolonged Q-T interval syndrome. The Scientist estimated that 17,700 persons had dilated Cardiomyopathy (14 per 100,000), 21,900 had hypertrophic cardiomyopathy, 300 had restrictive cardiomyopathy, 520 had ARVD, 640 had mitochondrial disease, 150 had Fabry's disease of the heart, and 1000 had prolonged Q-T interval syndrome. The prevalence of both dilated cardiomyopathy and hypertrophic cardiomyopathy was higher in men than women, the men to women ratios were 2.6 and 2.3; for dilated cardiomyopathy and hypertrophic cardiomyopathy respectively. Detailed data on patients with dilated cardiomyopathy or hypertrophic Cardiomyopathy were collected by a follow-up survey. In one year, more patients with dilated Cardiomyopathy (5.6%) died than with hypertrophic Cardiomyopathy (2.8%). Congestive heart failure and arrhythmias were the leading causes of death for dilated cardiomyopathy and hypertrophic cardiomyopathy respectively. Among patients with dilated Cardiomyopathy (n = 506) and hypertrophic Cardiomyopathy (n = 330) who had myocardial biopsies, significant mononuclear cell infiltrations were seen in 24% with dilated cardiomyopathy and 15% with hypertrophic cardiomyopathy suggesting the presence of inflammation in these patients. Cardiac troponin T was increased in 3% of dilated Cardiomyopathy and 9% of hypertrophic cardiomyopathy patients. Hepatitis B antibody was detected in 8.6% of dilated cardiomyopathy and 14.6% of hypertrophic cardiomyopathy patients. Hepatitis C antibody was present in 6.7% of dilated cardiomyopathy and 9.5% of hypertrophic cardiomyopathy patients. These data suggested that hepatitis B and C virus may cause myocarditis and cardiomyopathies. Five variables, left ventricular dilatation, lower left ventricular ejection fraction, higher NYHA functional class, older age, and male sex were independently related to poor prognosis¹⁷.

Phenotypes of HCV cardiomyopathies

In an initial study, Japanese Scientist found HCV RNA by polymerase chain reaction in the hearts of 19% of patients with dilated cardiomyopathy²⁴. Over a 10-year period, they identified 9.9% of patients with dilated cardiomyopathy had evidence of HCV infection, in contrast to only 2.5% of patients with ischemic heart disease. The main clinical manifestations at initial presentation were heart failure and cardiac arrhythmias. Positive and negative strands of HCV RNA were detected in the hearts of patients by Matsumori²³. Because negative RNA molecules are intermediates in the replication of the HCV genome, the Scientist presume that HCV replicates in myocardial tissues. During the same period, identified 14.1% of patients with hypertrophic cardiomyopathy had evidence of HCV infection. None of these patients had a family history of hypertrophic cardiomyopathy. Apical hypertrophic cardiomyopathy was diagnosed in 8% of patients who had ace of spadeshaped deformities of the left ventricle. Histopathologic studies showed mild to severe degrees of myocyte hypertrophy in the right or left ventricle, mild to moderate fibrosis, and mild cellular infiltration. Positive and negative strands of hepatitis C virus RNA were found in the biopsied hearts of patients²⁵. Teragaki and coworkers²⁶ recently found 18 of 80 Japanese patients with hypertrophic cardiomyopathy (22.5%) had positive HCV antibodies, a prevalence significantly higher than in controls.

Prolonged persistence of hepatitis C virus genomes in paraffin-embedded hearts

A multicenter study by the Scientific Council on Cardiomyopathies of the World Heart Federation (Bernhard Maisch, MD, Chairman) showed HCV genomes in 18% of patients with dilated Cardiomyopathy and myocarditis from Italy, and in 36% from the United States, two from patients with myocarditis and two with ARVC²⁵, which suggests that HCV may cause ARVC. In collaboration with the National Cardiovascular Center and Juntendo University, they have detected HCV RNA in paraffin sections of autopsied hearts from 26% of patients with hypertrophic cardiomyopathy, 12% of patients with dilated cardiomyopathy, and 33% of patients with myocarditis¹³. They also examined autopsied hearts from patients with dilated cardiomyopathy in a collaborative study with the University of Utah and found HCV RNA in 35% of hearts. The sequences of HCV genomes recovered from these hearts were highly homologous to the standard strain of HCV²⁵. However, the rates of HCV genomes detection in the hearts of patients with cardiomyopathies varied widely among different regions of the world. For example, no HCV genome was detected in hearts obtained from St. Paul's Hospital, in Vancouver, Canada. These observations suggest that the frequency of cardiomyopathy caused by HCV infection may be different in different regions or different populations. They analyzed sera stored during

the US Myocarditis Treatment Trial²⁷ of immunosuppression in patients with heart failure and myocarditis. Anti-HCV antibodies were identified in 4.4% of patients, including 5.9% of patients with biopsy-proven myocarditis. According to the US Centers for Disease Control, the prevalence of HCV infection in the general US population is 1.8%²⁸, thus HCV infection is more prevalent in patients with heart failure because of myocarditis. Furthermore, variations between 0% and 15% were found in the prevalence of HCV infection among the different medical centers and regions²⁹.

Genes responsible for development of different phenotypes of HCV cardiomyopathies

The major human histocompatibility complex (MHC) is located on the short arm of chromosome 6 and encodes for several protein products involved in immune function, including complement, TNF-a, and the human leukocyte antigen (HLA) complex, the polymorphisms of which are often proposed as determinants for the susceptibility to various diseases. Recent studies of HCV hepatitis showed that DQB1*0301 was associated with clearance of the virus. DRB1*1101, which is also in linkage disequilibrium with DQB1*0301, was associated with clearance. Several other studies have examined the association of MHC alleles with the progression of liver disease, and DQB1*0401 and DRB1*0405 were more prevalent among patients who developed chronic liver disease. The Scientist found that DPB1*0401 and DPB1*0901 were significantly associated with an increased risk of HCV-hypertrophic cardiomyopathy in the dominant model. The disparity in the gene-dose effect of two susceptible DPB1 alleles may be attributable to the difference between the susceptible and resistant residue-combination consisting of the DPb anchor pocket for antigenic peptide-binding. These results implied that the HLA-DP molecules with a specificity pocket appropriate for an HCV antigen(s) might confer the progressive process of hypertrophic Cardiomyopathy in HCV-infected individuals³⁰. In addition, no significant association was found between the MHC markers and HCV-hypertrophic cardiomyopathy. This marked difference in the MHC-related disease susceptibility for HCV-associated cardiomyopathy strongly suggests that the development of HCV-dilated cardiomyopathy and HCV-hypertrophic cardiomyopathy is under the control of different pathogenic mechanisms³¹.

Treatment of HCV cardiomyopathies

In patients with HCV hepatitis, the success of treatment can be measured by the biochemical and virologic responses. However, therapeutic markers have not been introduced in clinical practice to follow HCV cardiomyopathies. The Scientist have examined the effects of interferon on myocardial injury associated with active HCV hepatitis in collaboration with Shimane University³². They used TL-201-SPECT imaging, because it is more sensitive than electrocardiography or echocardiography to detect myocardial injury induced by HCV. The SPECT scores decreased in 8 of 15 patients (53%) in whom interferon treatment was completed. Circulating HCV disappeared after interferon therapy in all patients who had either a decrease or no change in SPECT scores, and HCV genomes persisted in the blood of two patients whose clinical status worsened³². This preliminary study suggests that interferon is a promising treatment for myocardial disease caused by HCV. They have also reported beneficial treatment with interferon guided by serial measurement of serum HCV RNA and cardiac troponin T in a patient presenting with dilated cardiomyopathy and striated myopathy attributable to HCV infection³³.

Impact of HCV on vascular disease

There are several reports on the association of diabetes mellitus and HCV infection. Also, HCV infection has been reported as a risk factor of atherosclerosis^{39, 40}. Therefore, it is important to clarify the role of HCV infection as a risk factor of atherosclerosis and vascular diseases, especially in the geographic areas with a high prevalence of HCV infection. The anti-viral treatment may cure the diseases, and decrease the number of patients. Thus, the cost to treat these patients and importantly, the burden of disease to the patients can be decreased³⁹⁻⁴².

Conclusion:

In Bangladesh, we are not very aware of Hepatitis virus induced heart disease. Some personal communication reveal that Hepatitis positive case are reported with heart failure. No proven strategy has been identified to reduce the numbers of patients living with, and dying from, cardiomyopathies and myocarditis due to infectious agents. Yet, the morbidity and mortality are high. We can treat these diseases by anti-viral agents, and some patients with cardiomyopathies and heart failure may be cured. Thus, treating these persons much earlier in the disease process—because of the ability to identify and treat HCV infection—will reduce the burden of cardiomyopathies and heart failure in patients and reduce the costs of treatment of these complex and long-term diseases.

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