A clinical and Molecular Genetic Study on Egyptian Children with Hypertrophic Cardiomyopathy

THESIS

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Abstract

Hypertrophic cardiomyopathy (HCM) is a primary genetic disease of the myocardium characterized by hypertrophy of left ventricle and interventricular septum and inherited in an autosomal dominant fashion. It is one of the leading causes of sudden cardiac death especially in the young. This study included 10 patients with isolated HCM and 10 normal children as a control group. The aim of the work is to determine whether exon 13 in MYH7 gene is a site for mutations among Egyptian children in the studied sample. Detailed History, clinical examination, echocardiography were done to all patients. Molecular screening for mutations in exons 13 & 14 of MYH7 gene was done to the patients' & control groups. Screening results revealed; normal amino acids sequence in exon 13 among studied sample, normal polymorphism detected among both patients' & control groups. Conclusion: Mutations in exons 13 & 14 of MYH7 may not be the cause of HCM among studied sample.

Key words:

- Hypertrophic Cardiomyopathy.
- Egyptian children.
- MYH7 gene.
- Exon 13 & 14 mutations.

CONTENTS	
♦LIST OF TABLES	I
♦LIST OF FIGURES	
♦LIST of ABBREVIATIONS	
♦INTRODUCTION	
♦ AIM OF WORK	
♦REVIEW Of LITERATURE	
Chapter I: Cardiomyopathies	4
Chapter II: Hypertrophic cardiomyopathy	
• Definitions & prevalence	8
■ Nomenclature	
Pathology	10
 Manifestations of HCM 	
 Mortality rate & risk factors 	14
 Genetic Basis of HCM 	16
Diagnosis of HCM	22
Management of HCM	26
 Prenatal diagnosis of HCM 	36
♦ SUBJECTS & METHODS	37
♦ RESULTS	41
♦ DISCUSSION	
♦SUMMARY	73
♦ CONCLUSION & RECOMMENDATIONS	75
♦ REFERENCES	
◆ APPENDICES	
♦ ARABIC SUMMARY	

List of Tables

No.	Title	Page
Table (1)	Risk factors for sudden cardiac death in	16
	patients with hypertrophic cardiomyopathy	
Table (2)	Causal genes for hypertrophic	18
	cardiomyopathy	
Table (3)	Surgical myectomy versus transcatheter	32
	septal ablation with ethanol	
Table (4)	The Sequence of PCR primers of MYH7 gene	39
Table (5)	Characteristics of HCM and control groups	41
Table (6)	Echocardiographic measurments of our cases	46
Table (7)	length of MYH7 gene fragments	48

List of Figures,

List of Figures

No.	Title	Page
Figure (1)	Transverse section of both ventricles at the	10
	mid-septal level. The hypertrophy involves	
	both the free and septal walls of the left	
	ventricle with marked thickening of the	
	septum—asymmetrical hypertrophic	
	cardiomyopathy (ASH)	
Figure (2)	Endocardial thickening in the subaortic area of	11
	the left ventricle due to contact of the anterior	
	cusp of the mitral valve with the hypertrophied	
	basal septum in a heart from a patient with	
F: (2)	hypertrophic cardiomyopathy (HCM)	
Figure (3a)	showing pinwheel configuration	12
Figure (3b)	showing herringbone pattern	12
Figure (4)	Echocardiographic long-axis view, showing	23
T: (5)	variable HCM types	40
Figure (5)	Consanguinity of the parents of our patients	42
Figure (6)	Family history of similar condition	43
Figure (7)	Family pedigree of cases no. 3 & 4	44
Figure (8)	Different clinical presentations	45
Figure (9)	Different pathological phenotypes	47
Figure (10)	PCR amplified products of exon 13 of index	48
	patients fragment size 190	
Figure (11)	PCR amplified products of exon 14 of index	49
	patients fragment size 258	
Figure (12)	Partial sequence of MYH7 (exon 13) showing	49
	normal sequence of 403 position.	
Figure (13)	Partial sequence of MYH7 (exon 13) showing	50
	normal sequence of 390 position	
Figure (14)	Partial sequence of MYH7 (exon 13) showing	50
	normal sequence of 383 position	
Figure (15)	Partial sequence of MYH7 (intron 14) showing	51

List of Figures,

No.	Title	Page
	homo SNP C instead of A	
Figure (16)	Partial sequence of MYH7 (exon 14) showing normal sequence of 453 position	51
Figure (17)	Partial sequence of MYH7 (exon 14) showing normal sequence of 443 position.	52
Figure (18)	Partial sequence of MYH7 (exon 14) showing normal sequence of 428 position.	52
Figure (19)	Family Pedigree of case no.10 demonstrating 4 successive generations	53
Figure (20)	Two-D Echocardiogram of the case before surgical myomectomy operation: long axis view showing marked hypertrophy of IVS compared to P	54
Figure (21)	Two-D echocardiogram of the patient's father showing; hypertrophy of IVS	55
Figure (22)	Two-D Echocardiogram of the patient after the surgery, long axis view showing the site of myomectomy in subaortic area, hypertrophy of IVS & PW and speckled myocardium	56
Figure (23)	Family pedigree of cases no. 1 & 2 showing double consanguinity of their parents	57
Figure (24)	2-Echo. Case no.(1); long axis view showing asymmetrical LV hypertrophy	58
Figure (25)	Echocardiogram of case no. 2; long axis view showing ASH with midcavitary obstruction	59
Figure (26)	Two-D echocardiogram, long axis view showing subaortic muscular ridge (tunnel-like narrowing)	60
Figure (27)	Family pedigree of case no. 7	61

List of Abbreviations,

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- ACE: angiotensin converting enzyme
- AD: autosomal dominant
- AHA: American heart association
- AR: autosomal recessive
- ARVC: arrhythmogenic right ventricular cardiomyopathy
- ASH: asymmetrical septal hypertrophy
- AV: atrioventricular
- BP: blood pressure
- BSA: body surface area
- CPB: cardiopulmonary bypass
- CUCH: Cairo University Children's Hospital
- DCM: dilated cardiomyopathy
- DM: diabetes mellitus
- DNA: deoxyribo nucleic acid
- HCM: hypertrophic cardiomyopathy
- HOCM: hypertrophic obstructive cardiomyopathy
- HR: heart rate
- ICD: intracardiac defibrillator
- IHSS: idiopathic hypertrophic subaortic stenosis
- IVS: interventricular septum
- LV: left ventricle
- LVEDD: left ventricular end diastolic dimension
- LVEF: left ventricular ejection fraction
- LVH: left ventricular hypertrophy
- LVOTO: left ventricular outflow tract obstruction

- M: month
- MRI: magnetic resonance imaging
- MyBPC: myosin binding protein c
- MYBPC3: myosin binding protein c gene
- MYH7: myosin heavy chain gene
- PG: pressure gradient
- PPI: permanent pacemaker implantation
- PWT: posterior wall thickness
- RCM: restrictive cardiomyopathy
- RVOTO: right ventricular outflow tract obstruction
- SAM: systolic anterior motion
- SCD: sudden cardiac death
- SD: standard deviation
- TDI: tissue Doppler imaging
- WHO: World Health Organization
- Y: year
- Z score: is the SD for echocardiographic measurements in relation to age & BSA
- ACTC1: alpha cardiac actin gene
- B-MyHC: beta myosin heavy chain protein

Introduction

Introduction

Hypertrophic cardiomyopathy (HCM) (OMIM #192600) is a primary cardiac disease of the myocardium of autosomal dominant inheritance. It is characterized by left ventricular hypertrophy without chamber dilatation, in the absence of either a systemic or other cardiac disease, which may cause a similar magnitude of hypertrophy (Wang et al., 2010).

It is a genetic disease that results mainly from mutations in genes encoding proteins of the Sarcomere, the most common of which is MYH7 gene that accommodates for about 25 % of HCM mutations (Alcalai et al., 2008).

HCM is a complex and confusing disorder that has been a subject of intense scrutiny for the past 50 years and of great interest to cardiologists, genetists and pathologists. And that is attributed to its diverse pathological, clinical and molecular heterogeneity (Maron et al., 2003).

The estimated prevalence of HCM in the general population is 1:500. It equally affects males and females. Moreover, it is increasingly detected in many races and countries (Maron et al., 1995).

HCM has got variable clinical presentations; from completely asymptomatic to severe symptoms (arrhythmias, angina, syncope...etc.) & even death. It is considered as one of the leading causes of sudden cardiac death (SCD) of young apparently healthy individuals (including athletes) (Maron et al., 2009).

It can present at any age and may be detected in infancy, during childhood and adolescence, or as an accidental finding in the elderly, consequently the pathologist may be the first to encounter a case of HCM at autopsy (Hughes, 2004).

The pathological hallmark of the disease is myocyte hypertrophy and disarray (Shirani et al., 2000).

It can be divided into 2 types; non obstructive HCM or hypertrophic obstructive cardiomyopathy (HOCM) (Davies and McKenna, 1994).

Echocardiography is the cornerstone in diagnosis of HCM (Losi et al., 2010), which is clinically diagnosed by the presence of primary cardiac hypertrophy and a preserved or enhanced LVEF (Maron, 2002).

Predictive genetic testing has an important role in early identification of asymptomatic relatives of HCM patients in attempt to decrease the risk of SCD (Christiaans et al., 2008).

Variable management strategies are being implied according to patients' manifestations; pharmacological therapy (as β -blockers & Ca channel blockers) & non-pharmacological therapy as transcatheter septal ablation or surgical myomectomy of the hypertrophied septum. The later is considered as the "gold standard" for therapy of HOCM with an excellent functional outcome (Marian, 2009).

Aim of the Work

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Clinical assessment of cases with hypertrophic cardiomyopathy to detect familial cases.

Molecular screening for exon 13 of MYH7 gene to define whether it is the most common mutational hot spot among studied HCM cases or not.

Providing genetic counseling concerning nature, inheritance, recurrence risk and implications of the disease to cases proved to have a documented mutation.