Study of Apolipoprotein E allele, Coronary Risk Factor and Cardiovascular Function in Asymptomatic Diabetic Elderly Patients

Thesis submitted for partial fulfillment of the PHD Degree in Geriatric and Gerontology medicine

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بني إللهُ البَّمْزَ الرَّحِينَ مِ

[وقُل رَّبِّ زِدْنِي عِلْماً]
سورة طه الآيه رقم ١١٤



Acknowledgement

First of all, all gratitude is due to \mathbf{God} almighty for blessing this work, until it has reached its end, as a part of his generous help, throughout my life.

Really I can hardly find the words to express my gratitude to **Professor Moatassem Salah Amer,** Professor of Geriatric and Gerontology Medicine, faculty of medicine, Ain Shams University, for his supervision, continuous help, encouragement throughout this work and tremendous effort he has done in the meticulous revision of the whole work. It is a great honor to work under his guidance and supervision.

I would like also to express my sincere appreciation and gratitude to **Professor Tarek Khairy Abd Eldayam** Professor of Cardiology Medicine, faculty of medicine, Ain Shams University, for his continuous directions and support throughout the whole work.

Really I can hardly find the words to express my gratitude to **Professor Randa Abdel Wahab Reda Mabrouk**Professor of Clinical pathology, Faculty of Medicine, Ain Shams University for his continuous directions and meticulous revision throughout the whole work. I really appreciate their patience and support.

I feel deeply thankful to **Assistant Professor Mohamed Shawaky Khater**, Assistant Professor of Geriatric and Gerontology Medicine, Faculty of Medicine, Ain Shams University, for his continuous unlimited help, unlimited patience and close supervision throughout the entire work.

Last but not least, I dedicate this work to my family, whom without their sincere emotional support, pushing me forward this work would not have ever been completed.



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List of Abbreviations

ABI Ankle-brachial index AD Alzhiemer disease APO E Apolipoprotien E **AGEs** Advanced glycation end products Acute myocardial infarction **AMI** Amyloid-β deposits APP-Aβ Angiotensin type 1 receptors AT1 AT2 Angiotensin type 2 receptors Body mass index **BMI CAC** Coronary artery calcium Coronary artery disease **CAD** Centers for Disease and Control **CDC** Prevention C-reactive protein **CRP** Cardiovascular diseases **CVD** \mathbf{DM} Diabetes mellitus European Association for the Study of **EASD Diabetes ECHO** Echocardiography **EF** Ejection fraction **FBS** Fasting blood sugar G6PDH Glucose-6-phosphate dehydrogenase **HDL** High density lipoprotein Heart failure HF HTN Hypertension

List of Abbreviations Insulin-like growth factor **IGF** IL-6 Interleukin-6 **IDF** International Diabetes Federation IDL Intermediate density lipids LDL Low density lipoprotein Low-density lipoprotein receptor **LDLR** Lipoprotein-associated phospholipase A2 Lp-PLA2 LRP-1 Low density lipoprotein receptor-related protein-1 LVH Left ventricular hypertrophy **PAFAH** Platelet-activating factor acetylhydrolase **PKC** Protein kinase C Postprandial blood sugar **PPBS** TG **Triglycerides VLDL** Very low density lipoprotein Plasminogen activator inhibitor-1 PAI-1 Phosphoinositol biphosphate PIP2 Renin-angiotensin-aldosterone **RAAS** Receptor for advanced glycation end **RAGE** products Synj1 Phosphoinositol phosphatase synaptjanin1 TNF-a Tumor necrosis factor alpha WHO World Health Organization

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Abstract:

Aim: Study association between Apo E polymorphism and development of heart failure among diabetic patients.

Material and Methods: case control study conducted on 90 elderly participants and they were classified into three groups each had 30 participants first group diabetic with atherosclerotic complications, second was diabetics without any complications while third group was non diabetics as control all of them were subjected to assessment of blood sugar and lipid profile and Apo E allele detection as well as Echocardiography for assessment of heart failure.

Results: study showed that among diabetic patients with atherosclerotic complication there is increased incidence of heart failure more than diabetic without complications or non-diabetic patients and that incidence of heart failure is higher among those carrying Apo E4 allele.

Conclusion: the study concluded that Apo E4 allele is associated with increased risk of development of heart failure among diabetic patient mostly due to effect of Apo E4 on lipid metabolism and atherosclerotic process

Keywords: *Apolipoprotien E allele, heart failure, diabetes*

Introduction

Type 2 diabetes mellitus (T2DM) is one of the most common diseases with a high incidence and prevalence throughout the world. It affects nearly 4% of the world's population and this percentage will supposedly be increasing up to 5.4% by year 2025 (*Chaudhary et al.*, 2012).

Among diabetic patients a high prevalence of coronary heart disease is observed at a relatively young age. Thus, risk factors for atherosclerosis must be defined and avoided in patients with diabetes mellitus. Abnormality of lipids such as high triglyceride levels and low HDL cholesterol levels emerged as residual cardiovascular risks for diabetic patients (*Ehara et al., 2012*).

Patients with type 2 diabetes mellitus have an increased incidence of atherosclerotic cardiovascular disease. This increase is attributable, in part, to associated risk factors, including hypertension and dyslipidemia. The latter is characterized by elevated plasma triglyceride levels, low levels of high-density lipoprotein (HDL) cholesterol, and small, dense low-density lipoprotein (LDL) particles (*Ginsberg et al.*, 2010).

Importance of searching for risk factors causing coronary heart disease (CHD) in diabetics arise from that macrovascular complications represent a major cause of mortality in type 2 diabetes, and MI and stroke accounting for about 80% of all deaths (*Gudbjörnsdottir et al.*, 2011).

T2DM is also known as a major independent risk factor for coronary artery disease (CAD) and is the major cause of morbidity and mortality affecting people with diabetes. To date, several mechanisms such as

Introduction and Aim of the Work

dyslipoproteinemia, obesity, oxidative stress, smoking, exercise, alcohol intake, and genetic factors have been identified as risk factors of both T2DM and CAD. Lack of apolipoprotein E (apoE) gene has been clearly demonstrated as a leading cause of severe hyperlipidemia and spontaneous development of atherosclerosis in mammals (*Chaudhary et al.*, 2012).

Apolipoprotein E (apoE) is a multifunctional protein that plays a key role in the metabolism of cholesterol and triglycerides by binding to receptors on the liver to help mediate clearance of chylomicrons and very low-density lipoproteins from the bloodstream. Although individuals carrying the $\varepsilon 4$ allele have higher and those carrying the $\varepsilon 2$ allele have lower total cholesterol levels than people with the commonest $\varepsilon 3/\varepsilon 3$ genotype, studies of lipid markers have typically involved too few participants to characterize relationships with different lipid subfractions across the 6 common genotypes. Different studies showed that compared with $\varepsilon 3/\varepsilon 3$ individuals, $\varepsilon 4$ carriers have a much greater risk of coronary disease (Benet et al., 2007).

Age and sex have been proposed as potential effect modifiers of the association between *APOE* and CHD risk because there is evidence that genotypic influence on mortality can vary by birth year and that estrogen and *APOE* may jointly affect lipid levels. Cigarette smoking, a well-established risk factor for CHD was also explored as an effect modifier between *APOE* genotype and CHD risk. It has been proposed that *apoe E4* carriers tend to produce a greater amount of LDLs, which makes them vulnerable to smoking-related increases in lipoprotein oxidation (*Ward et al.*, 2009).

Introduction and Aim of the Work

Aim of the work:

To study the association between apolipoprotein E genetic polymorphism, coronary risk factors and cardiovascular function among asymptomatic diabetic elderly patient.

Chapter One

Apolipoprotein E (Apo E)

Introduction:

Human plasma has five different types of apolipoproteins (A, B, C, D, and E) and some of them are further categorized into subtypes (*Eichner et al.*, 2002), (*Anthopoulos et al.*, 2010).

Apolipoprotein (Apo E) is a member of apolipoprotien gene family that was discovered in the 1970s and they are classified to three isoforms encoded by the E2, E3, E4 allele that were further subdivide into subforms giving six common isoforms of Apo E (Meigs et al., 2000), (Eichner et al., 2002), (Volcik et al., 2006), (Anoop et al., 2010).

Apolipoprotein gene polymorphism has different effects on lipid metabolism and it is associated with certain lipid transport disorders, so it can play a role in different diseases that is why many researches were carried out aiming to understand their role in health and disease (*Davignon et al.*, 1988), (*Eichner et al.*, 2002), (*Marrzoq et al.*, 2011).

Shape and structure:

Plasma lipoproteins are spherical bodies composed of a nonpolar lipid core, primarily triglycerides and cholesteryl esters, with an external layer of phospholipids and apolipoproteins (*Eichner et al.*, 2002) (*Anthopoulos et al.*, 2010).

Review of Literature

Apolipoprotein is the only protein component that combine with free cholesterol, phospholipids, cholesterol esters, and some triacylglycerols to form lipoproteins (*Eichner et al.*, 2002), (*Anthopoulos et al.*, 2010).

Apo E is a glycosylated protein it is associated with other plasma glycoproteins, such as high density lipoprotein (HDL), very low density lipoprotein (VLDL), and chylomicrons and it is also linked to the gene for low density lipoprotein (LDL) receptors (*Anoop S et al., 2010*), (*Chaudhary et al., 2012*), (*Elmadbouh et al., 2013*).

The apolipoprotein gene is polymorphic resulting in 3 common alleles and 6 different genotypes which differ in amino acid sequence at positions 112 and 158, Apo E3 contains cysteine at 112 and arginine at 158 while Apo E2 has cysteine at both positions, and E4 has arginine at both sites (Volcik et al., 2006), (Chaudhary et al., 2012), (Ehara et al., 2012), (Elmadbouh et al., 2013).

The amino terminal region of Apo E is responsible for its binding to the LDL receptor and the carboxy terminal mediates the binding of Apo E to surface lipoproteins which is influenced by specific amino acid differences (Anoop et al., 2010), (Marrzoq et al., 2011), (Elmadbouh et al., 2013).

Apo E4 has one more positive charge while Apo E2 has one less, also Apo E2 and Apo E3 have reactive free sulfhydryl groups which can form disulfide bonds with other free sulfhydryl-containing proteins this may have important effects on the function of Apo E (*Davignon et al.*, 1988).