

The Journal of the Egyptian

Society of Endocrinology, Metabolism & Diabetes

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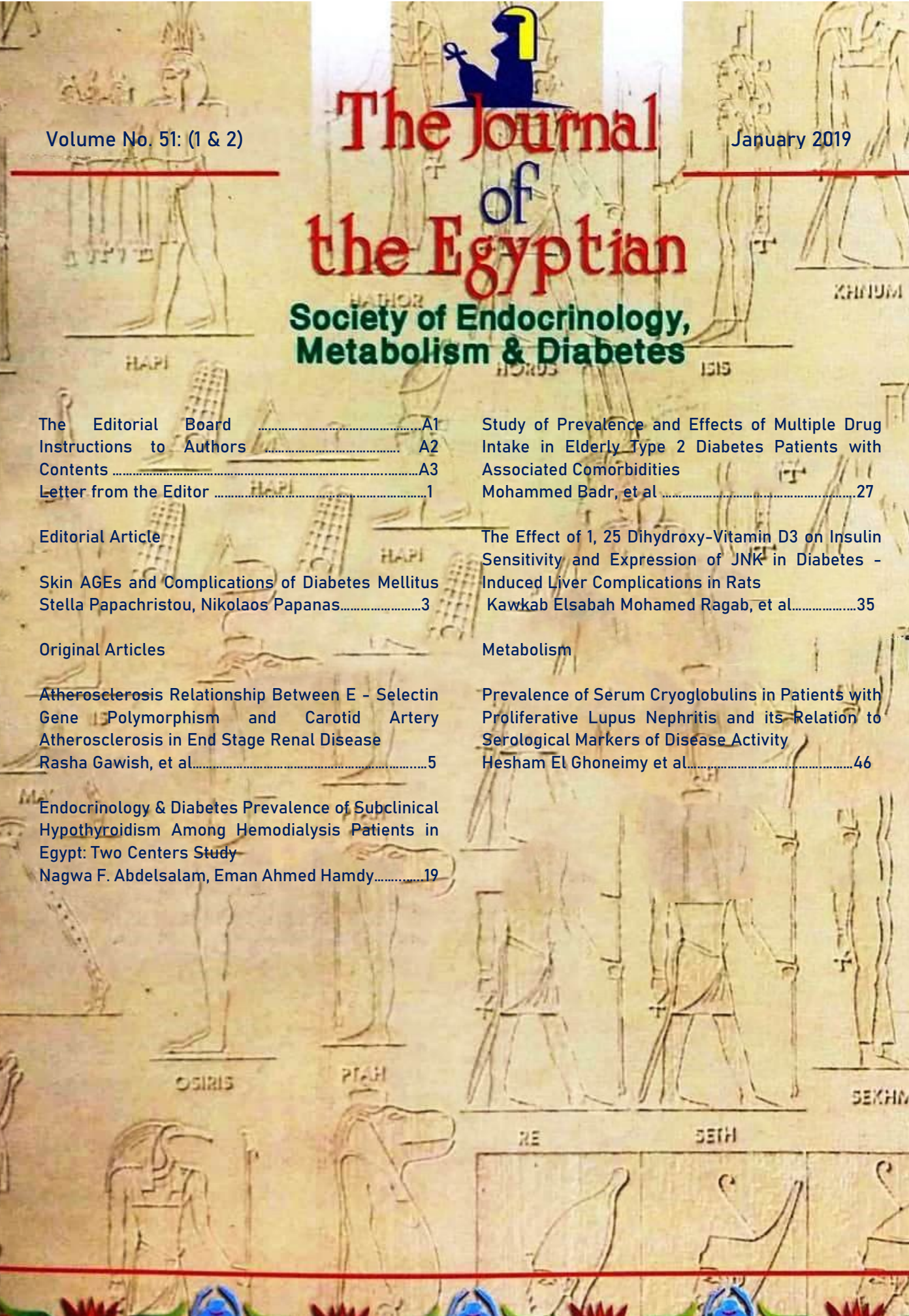
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Journal

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Prof. Samir Helmy Assaad-Khalil,
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Professor of Internal Medicine
Unit of Diabetes and Metabolism
Faculty of Medicine, Alexandria University
Alexandria, Egypt.
Tel.: (203) 4874204
Fax: (203) 4833321
e-mail: assaadkhalil@hotmail.com
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Letter from The Editor

Dear Colleague,

In this volume, we have included a very important editorial article which deals with an important issue of health care for persons with diabetes i.e. Complications of Diabetes Mellitus. Two colleagues from the University Hospital of Alexandroupolis, Athens; Greece Prof. Nikolaos Papanas and Prof. Stella Papachristou are sharing their experience with an article entitled: Skin AGEs and Complications of Diabetes Mellitus. Once again, we hope to meet your expectations, and until we meet in our next issue, deepest regards and best wishes.

The Editor

Prof. Samir Helmy Assaad Khalil

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Cover of the Journal: The gods and goddesses of the ancient Egyptian civilization including: Isis (goddess of prosperity and female fertility), Min (god of male fertility), ect... Designed by: S.H. Assaad Khalil

Skin AGEs and Complications of Diabetes Mellitus

Stella Papachristou, Nikolaos Papanas

Diabetes Centre-Diabetic Foot Clinic, Second Department of Internal Medicine, University Hospital of Alexandroupolis, Alexandroupolis, Greece

Address correspondence to: Prof. Nikolaos Papanas, Diabetes Centre-Diabetic Foot Clinic, Second Department of Internal Medicine, University Hospital of Alexandroupolis, G. Kondyli 22c, Alexandroupolis, Greece, email: papanasnikos@yahoo.gr

According to the International Diabetes Federation (IDF) statistics, it is estimated that every 7 seconds someone is dying due to diabetes mellitus (DM) or its complications.⁽¹⁾ Major complications of DM include cardiovascular disease, diabetic nephropathy, diabetic neuropathy, and retinopathy.⁽²⁾ These complications are also linked with substantial morbidity and poor quality of life.^(1,2) Therefore, timely diagnosis of DM and its complications is crucial. In this context, skin advanced glycation end products (skin AGEs) have received increased attention during the last years.⁽³⁾

During hyperglycaemic state, glucose forms a non-enzymatic process with plasma proteins, lipids, or nucleic acids, which is termed the “Maillard reaction”.⁽⁴⁾ This reaction results in the formation of endogenous AGEs. In addition to them, exogenous AGEs are of extreme importance. Indeed, dietary AGEs are by-products of heat-processed food consumption and are partially absorbed as such.⁽⁵⁾ Smoking is also a major source of exogenous AGEs.⁽⁶⁾ AGEs are metabolised via extracellular proteolysis as well as intracellular uptake and degradation in tissue macrophages, and Kupffer cells in the liver.⁽⁷⁾ Once degraded intracellularly, AGEs are further metabolised to second generation AGEs and excreted by kidneys.⁽⁷⁾

AGEs accumulate with increasing age. Nevertheless, their accumulation is exacerbated by many pathologic conditions, such as DM, cardiovascular disease, Alzheimer’s disease etc. In such conditions, AGEs are not merely a disease

manifestation, but instead a factor contributing to the pathophysiology.⁽⁸⁾

Measurement of AGEs can be accomplished by several different methods. These include ELISA using monoclonal or polyclonal antibodies, high performance liquid chromatography (HPLC), and mass spectrography.⁽⁹⁾ For tissue-bound AGEs, the gold standard remains as tissue biopsy, but it is invasive and very demanding.

In recent years, AGE-related autofluorescence has been used as a non-invasive option to measure AGEs. This method uses the autofluorescence, which is inherent in some major AGEs (e.g. pentosidine).⁽¹⁰⁾ In order to achieve quick, inexpensive and non-invasive measurements, the AGE reader (Diagnoptics, Groningen, The Netherlands) was produced.^(10,11) This is a portable device that illuminates a skin surface of approximately 4 cm² at the volar side of the dominant arm, guarding against surrounding light with an excitation light source. Emission and reflected extraction light are measured with a spectrometer using a glass fibre.^(10,11)

Increased skin autofluorescence (SAF) has been linked with microvascular complications of DM such as nephropathy, neuropathy and to a lesser extent retinopathy.⁽¹⁰⁻¹²⁾ In subjects with type 2 DM, this method was also able to predict likelihood of developing these microvascular complications except for retinopathy.⁽¹²⁾

SAF can contribute to the timely and accurate identification of the future risk for diabetic

peripheral neuropathy (DPN).⁽¹³⁾ In high-risk subjects, close monitoring, multifactorial treatment and education should be intensively offered to avert further risk of complications, notably foot ulcerations.^(13,14)

In conclusion, skin AGEs are increasingly being appreciated for the timely diagnosis of chronic DM complications. For this purpose, they merit further widespread utilisation, to improve patient outcomes.⁽¹³⁻¹⁵⁾

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Relationship Between E - Selectin Gene Polymorphism and Carotid Artery Atherosclerosis in End Stage Renal Disease

*Rasha Gawish¹, Manar Elgelany¹, Mohamed Abd-Elkader¹, Mona Sobhy²,
Omnia Gamal El-din³*

(1) Department of Internal Medicine, Nephrology & Transplantation Unit, Faculty of Medicine Alexandria university.

(2) Department of Medical Biochemistry Department, Faculty of Medicine, Alexandria University.

(3) Department of Radio Diagnosis, Faculty of Medicine, Alexandria University, Alexandria, Egypt

Corresponding author: Rasha Gawish Email: gawishrasha@gmail.com

Abstract:

Background: Adhesion molecules like the members of selectin family participate in the interaction between leukocytes and the endothelium. They are also involved in the pathogenesis of atherosclerotic processes. E-selectin is a cell surface glycoprotein that mediates the adhesion of leucocytes to vessels endothelium, an important early step in the atherosclerotic process. End-stage renal disease (ESRD) is a highly atherogenic disease, but it is unknown whether genetic polymorphism(s) in the E-selectin gene plays a role in the severity of arterial damage in this condition. **Objective:** The aim of the study was to test the relationship between the Leu 554 Phe polymorphism of the E-selectin gene and the severity of carotid atherosclerosis in End stage renal disease patients. **Methods:** In this study, we tested whether the Leu554Phe variant in the E-selectin gene is linked to carotid atherosclerosis as estimated by intima-media thickness (IMT), cross-sectional area and carotid stenosis using carotid ultrasonography in 70 well-characterized ESRD patients. The frequency of this polymorphism was also measured in a population sample of 30 healthy controls of the same geographical area using genetic analysis of Leu554Phe polymorphism of E-

selectin gene by restriction fragment length polymorphism **Results:** A total of 14.3% patients had the CC genotype, 78% had the CT genotype, 7.14% had the TT genotype and this distribution did not differ from that in the control population. Intima-media thickness (IMT) ($P = 0.926$) and cross-sectional area ($P = 0.567$) were not significantly higher in patients with the T-allele than in those without this allele. Furthermore, the degree of carotid stenosis was not significantly higher ($P = 0.593$) in patients with T-allele than in CC patients. On multivariate analyses including the traditional and non-traditional risk factors, the Leu554Phe polymorphism was confirmed as not an independent correlate of IMT ($P = 0.0926$), cross-sectional area ($P = 0.567$) and carotid stenosis ($P = 0.593$). **Conclusion:** In ESRD, the Leu554Phe polymorphism of E-selectin gene is not associated with the severity of carotid atherosclerosis, suggesting that genetically-determined alterations in the E-selectin molecule doesn't render ESRD patients with this gene variant particularly susceptible to the detrimental effects of inflammation on the arterial wall.

Key Words: E- selectin, Leu554Phe variant, carotid stenosis, ESRD,

INTRODUCTION

Atherosclerosis is a multifactorial disease that involves the interaction of environmental and genetic factors; it is the single most common cause of death and disability worldwide ⁽¹⁾. Accelerated atherosclerosis has been noticed in patients with CKD. It was also noticed that atherosclerotic lesions developed in the early stages of CKD ⁽²⁾.

Cardiovascular diseases are 3.5–100 times higher in patients who are on maintenance dialysis than the general population ⁽³⁾. End-stage renal disease patients have also much greater mortality from myocardial infarctions (MIs) ⁽⁴⁾. The mortality of patients receiving dialysis reaches 60 % in the first year after a first myocardial infarction.

Numerous pathophysiologic observations led to the postulation of the response-to-injury hypothesis of atherosclerosis, which initially stated that endothelial desquamation was the key event in atherogenesis ⁽⁵⁾. Now, it is considered that the pathophysiology of atherosclerosis is due to endothelial dysfunction rather than endothelial denudation. Whichever process is at work, each characteristic lesion of atherosclerosis represents a different stage in a chronic process of inflammation that takes place in the arterial wall ⁽⁶⁾.

The first morphologic phenomenon observed in plaque formation is the adherence of monocytes to the endothelial surface ⁽⁷⁾, monocytes adherence are triggered by a number of adhesion molecules such as vascular cell adhesion molecule-1 (VCAM-1), intercellular adhesion molecule-1 (ICAM-1), P-Selectin and E-Selectin. Moreover, there are T-lymphocytes that tend to accumulate in atherosclerotic lesions ⁽⁸⁾.

This adhesion is followed by monocytes migration into the intima, this migration is induced by the presence of certain mediators called chemo attractants in the intima as monocyte chemoattractant protein-1 (MCP-1) which is released from the endothelium ⁽⁹⁾.

In the intima, monocytes are activated, converted to macrophages and may divide. Macrophage cell division is triggered by macrophage colony stimulating factor (M-CSF), which can augment expression of macrophage scavenger receptors leading to uptake of modified lipoprotein particles and

formation of lipid laden macrophages or foam cells ⁽¹⁰⁾.

The macrophage foam cells produce a range of inflammatory cytokines including tumor necrosis factor alpha, metalloproteinases and the pro coagulant tissue factor that potentially stimulates thrombus formation when in contact with blood ⁽¹¹⁾.

Atherosclerosis in chronic kidney disease:

The increased risk of atherosclerosis in CKD patients is the result of interaction between traditional risk factors and specific uremic and dialysis related risk factors ⁽¹²⁾. Moreover, the prevalence of the traditional risk factors is greater in dialysis dependent patients than in the general population ⁽¹³⁾.

Cardiovascular risk factors in chronic kidney disease:

Risk factors can be divided into the following:

I- Traditional risk factors ⁽¹⁴⁾:

Traditional risk factors include the following: age , sex, family history , obesity, cigarette smoking, hypertension, diabetes mellitus dyslipidemia and sedentary life style

II- Non traditional risk factors ⁽¹⁵⁾:

They include homocysteine ,lipoprotein (a), asymmetric dimethyl arginine (ADMA), advanced glycation end products (AGEs) ,oxidative stress , infections and inflammation.

III- Other potential risk factors specific to ESRD

Risk factors related to ESRD include disturbance of glucose metabolism, anaemia ⁽¹⁶⁾, calcium-phosphate disorders ⁽¹⁷⁾, coagulation equilibrium imbalance ⁽¹⁸⁾, vitamin E-deficiency ⁽¹⁹⁾, hypoalbuminaemia ⁽²⁰⁾, haemodynamic overload (plasma volume expansion, arteriovenous fistula) ⁽²¹⁾.

IV- Risk factors related to dialysis ⁽²¹⁾:

Endotoxin, bioincompatibility of membranes, fluctuation in blood pressure and fluctuation in serum electrolytes level.

Inflammation and atherosclerosis

Atherosclerosis is considered as a chronic inflammatory process as suggested by Ross. Inflammation mediates all stages of atherosclerosis from initiation through progression and thrombotic complications.

Inflammation determines the fragility of the fibrous cap of the plaque, and its thrombogenic potentiality. T cells stimulate matrix metalloproteinases production such as proteinase, collagenase and also the production of the pro-coagulant tissue factors. Macrophage-derived matrix metalloproteinase digests the extracellular matrix of the plaque resulting in fibrous cap instability and plaque rupture. This in turn leads to the exposure of the tissue factor-rich lipid core to the circulating blood components, and thrombosis occurs⁽²²⁾.

Adhesion Molecules:

The adherence of circulating leucocytes to endothelial cells and the subsequent trans-endothelial migration to the vascular intima is one of the key events in the initiation of atherosclerosis⁽²³⁾. These processes are mediated chiefly by different cellular adhesion molecules (CAMs), which are released on the surface of vascular endothelial cells in response to pro-inflammatory cytokines, oxidants or oxidized LDL-cholesterol⁽²⁴⁾. Increased expression of CAMs have been described on endothelial cells, vascular smooth muscle cells and macrophages in human atherosclerotic plaques. Soluble types of these adhesive molecules are thought to be shed from activated cells and can be measured in peripheral blood. Plasma concentrations of these soluble forms of CAMs are elevated in septicemia, autoimmune diseases and allograft rejection⁽²⁴⁾.

Cell adhesion molecules fall mainly under four broad families selectins, immunoglobulin superfamily, integrins and cadherins⁽²⁵⁾. The members of immunoglobulin superfamily are related to immunoglobulins and are often expressed on cell surface. They are the biggest group of adhesion molecules including more than 70 members, e.g., intercellular adhesion molecules (ICAM), and vascular cell adhesion molecule-1 (VCAM-1)⁽²⁵⁾.

Members of the selectin family play an important role in the adhesion of leukocytes to activated endothelium. This produces

characteristic rolling motion of the leucocyte on the endothelial surface. The selectins can be classified according to their site of expression into: P-selectin (platelet), L-selectin (leucocyte) and E selectin (endothelium)⁽²⁶⁾.

E-selectin:

E-selectin was described as an antigen that was induced on cultured human umbilical vein endothelium after stimulation by interleukin 1 (IL-1) and that was contributed to the adhesion of neutrophils, previously described as endothelial leucocyte adhesion molecule-1⁽²⁷⁾.

E-selectin has a molecular weight of approximately 94.000 dalton. It is composed of an array of discrete protein domains; an amino terminal, C-type lectin domain, a single epidermal growth factor (EGF) like domain and from 4-9 short consensus repeats (SCR), a single membrane spanning region and a cytoplasmic tail with a 589 amino acids as shown in Figure (1) The terminal lectin domain and an epitope within the EGF like region are important for ligand binding⁽²⁸⁾. The human E-selectin gene (*SELE*) is located on chromosome 1q. It consists of 14 exons and 13 introns, spanning approximately 13kb⁽²⁹⁾.

E-selectin is expressed only by cytokines activated endothelial cells, cytokines such as interleukin-1 and tumor necrosis factor alpha induce ICAM, VCAM, and E-selectin. Interleukin-1 mRNA and tumor necrosis factor protein have been demonstrated in smooth muscle cells and macrophages of atherosclerotic plaques. At the endothelial cells, E-selectin as other selectins mediate leukocyte rolling which is the initial step in leukocytes adhesion to the endothelial cells⁽³⁰⁾.

E-selectin is maximally released 2-4 hours after cell has been activated. Within the next 24-48 hours, E-selectin is removed from the cytoplasmic membrane by shedding into the circulation⁽³¹⁾ Plasma E-selectin levels were increased in patients with hypertension, hypercholesterolemia and diabetes mellitus, all the three disorders are important risk factors for atherosclerosis, it was expressed by endothelial cells overlying atherosclerotic plaques as well as by adventitial vessels⁽³²⁾.

Several lines of evidence indicate that besides environmental factors, the genetic background has an important influence on inflammation-induced arterial damage. The E

selectin genomic location was previously known to be linked to blood pressure. Moreover the polymorphism in the E-selectin gene (Leu554Phe) hinders the E selectin shedding from the endothelium was believed to be associated with coronary atherosclerosis⁽³³⁾.

In patients with ESRD, it is still unknown whether the genetic background plays a role in atherosclerosis and inflammation. It is still a relevant question because in uremia, the factors predisposed to atherosclerosis do not correspond to those in the general population. Adhesion molecules have a crucial role in the development of atherosclerosis, Therefore attention was given on Leu554Phe polymorphism as it enhances the anchoring of E-selectin to the endothelium⁽³⁴⁾. This alteration in amino acids sequence in E-selectin gene that might amplify the inflammation process on the vessel wall.

AIM OF THE STUDY

The aim of the study was to test the relationship between the Leu554Phe polymorphism of the E-Selectin gene and the severity of carotid atherosclerosis as estimated by intima-media thickness (IMT), cross-sectional area and carotid stenosis using carotid ultrasonography in End stage renal disease (ESRD) patients.

SUBJECTS AND METHODS

The present study was conducted on 100 subjects selected from the kidney dialysis unit, El-Mowasah hospital, Alexandria Main University Hospital.

The design of the study consist of two groups:

Group (I): included 70 dialysis patients of age ranged between 20 to 70 years old ,who have been on regular dialysis treatment (RDT) for at least 6 months and who were free of overt infections (fever, infected vascular access , peritonitis or exit-site infection) but suffered from other chronic disease as hypertension , diabetes and cardiac diseases.

Group (II): included 30 healthy age and sex matched as patients who were considered as control group for genetic data analysis, and who were free from acute or serious chronic

diseases such as cancer, hypertension, diabetes and Coronary heart disease.

The following was done to all the enrolled subjects:

I- Thorough history taking and Complete clinical examination including:

1. Body mass index.

The patients were grouped according to their body mass index (BMI) into underweight (<18.5 kg/m²), average weight (18.5–24.9 kg/m²) and overweight (>24.9 kg/m²).

2. Vital signs.

• Heart rate

It is measured by palpation of the radial artery pulsation, number of heart beats per minute.

• Blood pressure

In hemodialysis patients pre dialysis and post dialysis BPs were calculated as the average value of all recordings [12 measurements (i.e. 3 per week)] taken during the month preceding the study. The mean value of pre dialysis and post dialysis BPs was then obtained for each patient and it was considered for global statistical assessment.

II- Laboratory investigations including:

1- Fasting blood glucose is performed during the mid-week non-dialysis day⁽³⁵⁾.

2- Blood urea and serum creatinine.

3- Haemoglobin level⁽³⁶⁾.

4- Fasting Lipid profile (Total cholesterol, serum triglycerides, low density lipoproteins cholesterol and high density lipoproteins cholesterol)⁽³⁷⁾.

5- Serum calcium, phosphorus and PTH⁽³⁸⁾.

6- CRP, measured by nephelometry (high sensitivity)⁽³⁹⁾.

7- Genetic analysis of Leu55Phe polymorphism of E-selectin gene by restriction fragment length polymorphism⁽⁴⁰⁾.

I- Carotid artery ultrasonography for vascular assessment:

Doppler studies were performed in longitudinal and transverse planes using anterior, lateral and posterior approaches by both B-mode imaging and power Doppler imaging. Assessment of atherosclerosis is done by measurements of:

1- Carotid intima media thickness ⁽⁴¹⁾:

Measurement of carotid IMT bilaterally was done by B-mode 7 MHZ probe [ultrasound examination was done using LOGIQ 7 PRO, GE (General Electric Medical System).

2- Degree of stenosis ⁽⁴²⁾.

3- Cross section area, was calculated by the standard formula ⁽⁴³⁾.

Statistical Analysis of the data ⁽⁴⁴⁾:

Data were statistically described in terms of mean \pm SD, median (interquartile [IQR] range) or frequencies (number of cases) and percentages when appropriate. Comparison of numerical variables between the study groups was done using Student's t test for independent samples in comparing two groups when normally distributed and Mann-Whitney U test for independent samples when not normally distributed .

Comparison of numerical variables between more than two groups was done using the one-way analysis of variance (ANOVA) test with post hoc multiple two-group comparisons in normal data, and the Kruskal Wallis test with post hoc multiple two-group comparisons in non-normal data. Association between qualitative data was done using the Chi-square test .

Correlation between various variables was done using the Pearson moment correlation equation for linear relation in normally distributed variables and Spearman's rank correlation equation for non-normal variables. P-Values less than 0.05 were considered statistically significant. All statistical calculations were done using computer programs SPSS (Statistical Package for the Social Science; SPSS Inc., Chicago, IL, USA) version 15 for Microsoft Windows.

RESULTS

I- Demographic data:

This study included 100 subjects, which were divided into two groups.

Group (I): included 70 dialysis patients, who have been on regular dialysis treatment (RDT) for at least 6 months and who were free of overt infections but suffer from other chronic disease as hypertension , diabetes and cardiac diseases, which were considered as risk factors of atherosclerosis. This group was then subdivided according to the genotype distribution of Leu554Phe E selectin gene polymorphism on basis of absence or presence of risk allele (T) into **(TT/CT)** versus **(CC)**.

Group (II): included 30 healthy age and sex matched as patients who were considered as control group for genetic data analysis, and who were free from acute or serious chronic diseases **Table (1)**.

II- Genotype distribution:

The genotype distribution of the Leu554Phe polymorphism of the E-selectin gene did not deviate from Hardy–Weinberg equilibrium either in patients (CC genotype, 14.3%; CT genotype, 78.6%; TT genotype, 7.1%) or in control healthy subjects (CC genotype, 13.3%; CT genotype, 83.3%; TT genotype, 3.3%) **Tables (2)**.

IV- Carotid artery ultrasonography for measuring the indicators of atherosclerosis:

1- Carotid intima media thickness (IMT):

In Patients, IMT ranged 0.60 – 1.6 mm with a mean of 1.08 ± 0.67 mm while in control, it ranged between 0.61 – 0.89 mm with a mean of 0.76 ± 0.10 mm, there were statistically significant difference between two groups as regard IMT, where $P < 0.001$ (P significant as $P < 0.05$). **Table (3)**

2- Cross section area:

In Patients, cross section area ranged 1.227-40.7 mm² with a mean of 21 ± 19.7 mm² while in control, it ranged between 13.85-46.56 mm² with a mean of 30.205 ± 16.355 mm², There were statistically significant difference between two groups as regard cross section area, where $P < 0.001$ (P significant as $P < 0.05$). **Table (3)**

3- Degree of stenosis:

In Patients, degree of stenosis ranged 0.0 – 20.0% with a mean of 2.29 ± 4.87 % while in control ,it ranged 0.0 – 10.0% between with a mean of 0.50 ± 2.01 % ,There were

statistically significant difference between two groups as regard degree of stenosis, where $P=0.040$ (P significant as $P<0.05$).

Table (3)

V- E-selectin Leu554Phe polymorphism and carotid atherosclerosis

When patients were categorized on the basis of the presence/absence of the risk allele (i.e. TT and CT vs CC patients), no significant difference emerged between the two groups as for demographic and somatometric data or for clinical and biochemical parameters. **Table (4)**

Relation of presence of the risk allele (TT/CT) and indicators of carotid atherosclerosis:

1) Intima media thickness:

Patient's intima Thickness in the Patients with CT or TT genotypes ranged between 0.60 – 6.1 mm with a mean of 0.888 ± 0.712 mm while in Patients with CC genotypes it ranged between 0.60 – 1.8 mm with a mean of 0.850 ± 0.375 mm. There were no statistically significant difference between patients with presence of risk allele (TT/CT) and patients with the absence of risk allele (CC) as regard IMT where $P=0.926$ (P significant as $P<0.05$). **Table (5)**.

2-Cross section area:

Patient's Cross section in the Patients with CT or TT genotypes ranged between 1.227-40.7 mm² with a mean of 21 ± 19.7 mm² while in

Patients with CC genotypes it ranged between 5.123-40.1 mm² with a mean of 20.996 ± 19.836 mm². There were no statistically significant difference between patients with presence of risk allele (TT/CT) and patients with the absence of risk allele (CC) as regard cross section area, where $P=0.567$ (P significant as $P<0.05$). **(Table 5)**

3- Degree of stenosis:

Patient's degree of stenosis in the Patients with CT or TT genotypes ranged between 0 – 20 % with a mean of 2.25 ± 4.642 % while in Patients with CC genotypes it ranged between 0 – 20 % with a mean of 2.5 ± 6.346 %. There were no statistically significant difference between patients with presence of risk allele (TT/CT) and patients with the absence of risk allele (CC) as regard degree of stenosis, where $P=0.593$ (P significant as $P<0.05$). **(Table 5)**

E- Selectin gene and carotid atherosclerosis multiple regression analyses:

Since the association between E-selectin genotypes and carotid atherosclerosis could be confounded by other risk factors, multiple regression analyses were performed adjusting for all variables which were associated with the three indicators of carotid atherosclerosis. In these analyses, mentioned risk factors (cofounders) were confirmed as a positive correlate of IMT, cross sectional area and the severity of carotid stenosis **(Table 6)**.

Table (I): Comparison between the two studied groups according to demographic data

	Cases (n = 70)		Control (n = 30)		Test of Sig.	p
	No.	%	No.	%		
Sex						
Male	40	57.1	13	73.3	$\chi^2=$ 1.608	0.205
Female	30	42.9	17	56.7		
Age (years)					t= 1.941	0.055
Min. – Max.	22.0 – 74.0		27.0 – 55.0			
Mean \pm SD.	51.06 ± 13.52		47.30 ± 5.84			
Median	52.50		48.50			

χ^2 : Chi square test

t: Student t-test

p: p value for comparing between the two groups

Table (II): Comparison between the two studied groups according to E-selectin gene polymorphism

	Cases (n = 70)		Control (n = 30)		χ^2	MCp
	No.	%	No.	%		
E-selectin						
TT	5	7.1	1	3.3	0.437	0.918
CC	10	14.3	4	13.3		
CT	55	78.6	25	83.3		
HWE	0.053		0.139			

χ^2 : Chi square test MC: Monte Carlo test

p: p value for comparing between the two groups

HWE: p value for comparing observed and expected data according to Hardy-Weinberg between the two groups

Table (III): Comparison between the two studied groups according to the indicators of atherosclerosis

	Cases (n = 70)	Control (n = 30)	U	p
INT thickness (mm)				
Min. – Max.	0.60 – 1.6	0.61 – 0.89	397.50*	<0.001*
Mean ± SD	1.08 ± 0.67	0.76 ± 0.10		
Median	1.0	0.77		
Cross section (mm²)				
Min.– Max.	1.227– 40.7	13.85– 46.56	3.871*	<0.001*
Mean ± SD.	21 ± 19.7	30.205± 16.355		
Median	16.20	17.0		
Degree of stenosis %.				
Min. – Max	0.0 – 20.0	0.0 – 10.0	863.0*	0.040*
Mean ± SD.	2.29 ± 4.87	0.50 ± 2.01		
Median	0.0	0.0		

U: Mann Whitney testt: Student t-test

p: p value for comparing between the two groups *: Statistically significant at $p \leq 0.05$

Table (IV): Main demographic, somatometric, clinical and biochemical characteristics of the study population divided on the basis of E-selectin Leu554Phe polymorphism

	Patients with CT or TT genotypes (n=60)	Patients with CC genotypes (n=10)	P-Value
Age	51.38±13.581	49.10±13.658	0.817
Sex: n (%)			
Male	35(58.3%)	5(50%)	0.735
Female	25(41.7%)	5(50%)	
BMI	25.258±2.316	25.300±2.611	0.518
Smoker n (%)	10(16.7%)	4(40%)	0.104
DM n (%)	21(35%)	5(50%)	0.483
Systolic	128±18.208	116±21.705	0.988
Diastolic	81.33±9.994	72±12.293	0.878
Duration of RDT	6.277±4.032	6.200±3.190	0.654
HR	65.10±6.262	64.50±4.972	0.510
Hemoglobin	10.682±1.392	10.730±0.874	0.118
WBCs	6.543±3.286	6.350±3.177	0.785
Cholesterol	168.27±45.217	147.30±28.706	0.104
Triglyceride	177.40±109.46	140.00±50.323	0.465
LDL	106.27±24.378	112.20±17.492	0.276
HDL	60.42±9.230	66.30±4.191	0.095
Ca	8.658±0.983	9.190±0.690	0.436
PO₄	5.243±2.072	5.74±2.560	0.361
Ca X PO₄	45.660±19.731	52.625±23.405	0.322
PTH	553.692±405.294	685.840±316.997	0.228
Albumin	3.833±0.275	3.620±0.561	0.617
Creatinine	10.026±2.644	9.630±2.008	0.475
Urea	129.95±26.602	128.5±28.602	0.934
FBG	90.242±16.273	88.00±13.333	0.265
CRP	8.086±8.814	6.455±6.852	0.569
IMT	0.888±0.712	0.850±0.375	0.926
Cross-sectional area	21±19.7	20±19.836	0.567
Degree of carotid stenosis	2.25±4.642	2.50±6.346	0.593

Data are mean ±SD or percent frequency, as appropriate

Table (V): Distribution of studied sample according to different parameters

	Patients with CT or TT genotypes (n=60)	Patients with CC genotypes (n=10)
Int. Thickness(mm)		
Min.	0.60	0.60
Max.	6.1	1.8
Mean	0.888	0.850
S.D	0.712	0.375
P Value	0.926	
Cross section (mm²)		
Min.	1.227	5.123
Max.	40.7	40.1
Mean	21	20.996
S.D	19.7	19.836
P Value	0.567	
Degree of stenosis%		
Min.	0	0
Max.	20	20
Mean	2.25	2.50
S.D	4.642	6.346
P Value	0.593	

Table (VI): Multiple regression analyses

	Beta regression coefficients	Partial R	P-Value
IMT (Multiple R= 0.744, P <0.001*)			
BP	0.007	0.339	0.005*
CaxPO ₄	0.015	0.672	<0.001*
Cross-sectional area (Multiple R= 0.934, P <0.001*)			
Age	0.010	0.545	<0.001*
BP	0.022	0.759	<0.001*
BMI	0.020	0.251	0.043*
FBS	0.254	0.454	<0.001*
Carotid stenosis (Multiple R= 0.909, P <0.001*)			
Smoking	0.633	0.259	0.033*
BP	0.029	0.906	<0.001
LDL	0.022	0.759	<0.001
Serum urea	0.20	0.251	0.043

DISCUSSION

There are 17 mutations that have been detected in adhesive molecules, five of which resulted in an amino acid substitution. In E-selectin, exchange at Ser¹²⁸ Arg in epidermal growth factor (EGF) domain and Leu⁵⁵⁴ Phe in membrane domain, and a DNA mutation from guanine to thymine (position 98) are examples of mutations due to amino acids substitutions⁽⁴⁵⁾.

In this study, Leu⁵⁵⁴Phe E-selectin polymorphism has been studied, in which the mutation in exon 10 leads to the substitution of leucine (Leu) by phenylalanine (Phe) at amino acid 554, (Leu = allele-C; Phe=allele-T). This mutation is believed to delay the shedding of E-selectin from vascular endothelium⁽⁴⁵⁾.

In the present study, the distribution of *SELE* rs5355 C and T genotypes did not show significant difference in both ESRD patients on dialysis and control groups. The genotype distribution of our studied Egyptian population group was identical to those reported by MS Issac et al.⁽⁴⁶⁾ however, they were different from those reported previously in Iranian⁽⁴⁷⁾ and Italian populations⁽⁴⁸⁾.

Three well established indicators of carotid atherosclerosis were studied, namely intima media thickness (IMT), cross-sectional area⁽⁴⁹⁾, and arterial stenosis⁽⁵⁰⁾. These indicators reflect different aspects of the atherosclerotic process. Our statistics showed lack of association between *SELE* rs5355 C>T genotypes and presence of carotid atherosclerosis in ESRD patients. This finding is in concordance with results reported previously by Isaac MS et al.⁽⁴⁶⁾ This result is contrary to results found by Testa et al,⁽⁴⁷⁾ who reported that IMT, CSA, and the degree of carotid stenosis were significantly higher in patients with CT and TT genotypes (i.e., in patients with T allele) when compared to CC patients.

Robert Bernat et al 2012, studied the relationship of Leu554Phe polymorphism for

atherosclerosis and Long-Term Outcome after Percutaneous Coronary intervention (PCI) with Stenting, The study included 190 patients with standardized clinical follow-up over 5 years, which were initially treated with PCI, The long term clinical outcome was defined by the major adverse cardiac events MACE: death, target vessel revascularization (PCI or coronary bypass grafting, CABG) and myocardial infarction. T allele carriers of Leu554Phe polymorphism were shown to have a significantly lower repeat revascularization rate in comparison with the CC genotype. There was no significant difference between the incidence of in-stent restenosis and Leu554Phe polymorphism⁽⁵¹⁾.

Srivastava et al, 2018 analysed the Leu554Phe polymorphism and expression of E-selectin gene in 250 patients with essential hypertension and 250 normal healthy controls. A significant association of E-selectin genotypes (CT + TT) with essential hypertension ($P < .0001$, Odds ratio = 2.2 [1.58-3.24] at 95% CI) was observed⁽⁵²⁾.

Marteau et al 2004, studied the association between E selectin polymorphism (Leu554Phe), blood pressure and obesity. 478 men and 546 women were selected from volunteers for a free health check-up. These individuals underwent two examinations (t_0 and t_{+5}) and they were not taking medications that can affect blood pressure. At t_0 , no relationship was observed between Leu554Phe polymorphism and blood pressure. However at t_{+5} , systolic blood pressure (SBP) was greater in individuals carrying the T allele, and the Leu554Phe polymorphism was associated with SBP in interaction with BMI ($P < 0.001$ in men and $P < 0.05$ in women). There was a steeper increase in SBP with BMI greater than 25 Kg/m² in carriers of the T allele than in CC homozygotes. Similar results were observed for diastolic blood pressure in men ($P = 0.0103$). These results suggest a BMI-specific effect of Leu554Phe polymorphism of the E-selectin gene on blood pressure⁽⁵³⁾.

CONCLUSION

The main conclusion that emerged from the present results was the following:

1. E-selectin Leu⁵⁵⁴Phe polymorphism can be of value as a genetic marker to assess risk of atherosclerosis in ESRD.

2. No significant correlation could be found between E-selectin Leu⁵⁵⁴Phe polymorphism and atherosclerosis markers in the studied group of ESRD patients on HD.

3. E-selectin Leu⁵⁵⁴Phe polymorphism may require an underlying risk factor that predisposes to endothelial dysfunction to exert its effect and be expressed.

4. The genotype distribution of the Leu⁵⁵⁴Phe polymorphism of the E-selectin gene was identical among ESRD patients on HD and control subjects of the same population.

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Prevalence of Subclinical Hypothyroidism Among Hemodialysis Patients in Egypt: Two Centers Study

Nagwa F. Abdelsalam¹, Eman Ahmed Hamdy²

(¹) Department of Internal Medicine (Nephrology Unit), Helwan University, Egypt., (²) Department of Clinical and Chemical Pathology, Helwan University, Egypt.

Abstract:

Background:

Thyroid dysfunction is commonly observed in chronic kidney disease patients. Both subclinical hypothyroidism (SCH) and end stage kidney disease are independent risk factors for cardiovascular mortality. **Aim of the Work:** Our aim has been to study the prevalence of subclinical hypothyroidism in hemodialysis patients and associated risk factors. **Patients & Methods:** The study was conducted at hemodialysis units of two tertiary care centers in Egypt. The study group included 92 end stage renal disease patients on regular hemodialysis 3 times/ week. Demographic data, current smoking, underlying kidney disease, duration of dialysis were obtained from each patient. Also, history taking, clinical examination and laboratory investigations (hemoglobin, serum level of urea before and urea after dialysis, urea reduction ratio, creatinine, calcium,

phosphorus, albumin, C. reactive protein, thyroid stimulating hormone TSH and free thyroxine T4) were done for each patient.

Results: Prevalence of SCH among our 92 patients was 13.1%. Cases of renal failure due to lupus nephritis were significantly higher in SCH group compared to normal TSH group ($p=0.028$). Patients with SCH had significantly higher duration of dialysis compared to the normal TSH group ($p=0.019$) and higher level of CRP compared to the normal TSH group ($P<0.001$). Multivariate analysis (both adjusted and unadjusted) showed that CRP was the most independent variable affecting SCH followed by duration of dialysis. **Conclusion:** The prevalence rate of SCH among HD patients in Egypt is 13.1% and we suggest the presence of an association between inflammatory markers and SCH. We suggest also that SCH is associated with longer HD duration.

Key words: hemodialysis patients, subclinical hypothyroidism, prevalence.

Introduction

Changes in thyroid hormones are commonly observed in chronic kidney disease (CKD) patients and thyroid dysfunction increases as CKD progresses.⁽¹⁾ End stage renal disease (ESRD) affects thyroid function and now formally listed as an established cause of chronic non-thyroidal illness.⁽²⁾

Alterations in thyroid hormones level and / or metabolism have been described in ESRD patients.⁽³⁾ Also, acute and chronic renal disease has its effects on the hypothalamus-pituitary-thyroid axis. Thyroid stimulating hormone (TSH) levels may be normal or increased in CKD but with reduced response to thyrotropin releasing hormone (TRH).⁽⁴⁾

Subclinical hypothyroidism (SCH) defined as high serum TSH that is above the defined upper limit of the reference range and normal free T4 levels that is within the reference range, is frequently seen in CKD patients.⁽⁵⁾ SCH is known to be a risk factor for atherosclerotic cardiovascular disease, hyperlipidemia and low grade inflammation.⁽⁶⁾

Both ESRD and SCH are independent risk factors for cardiovascular mortality. Previous studies have shown an increased prevalence of SCH in CKD patients but there is a little data with respect to prevalence of SCH in ESRD.⁽⁷⁾ Factors associated with SCH in ESRD patients have not yet been clearly identified.⁽⁸⁾ So, our aim in this study is to identify the prevalence of SCH in ESRD patients in two tertiary care centers in Egypt and associated risk factors.

Patients & Methods

Our study is a cross sectional study done in 2 tertiary care centers in Egypt over 2 years period. The first is the hemodialysis unit of Helwan University Hospital and the second is the hemodialysis unit of one of the Ministry of Health Hospitals in Alexandria. The total number of patients from both centers was 103 ESRD patients on regular hemodialysis (HD) 3 times / week. After exclusion of patients that met the exclusion criteria, those who consented were 92 adult ESRD patients on regular HD 3 times /week and those who formed our study cohort. Exclusion criteria included those who are previously diagnosed as overt hypothyroidism, SCH, overt hyperthyroidism, those who take medications affecting thyroid functions and pregnant females.

Demographic data, current smoking, underlying kidney disease, duration of dialysis in months were obtained from all the patients.

Also, history taking, clinical examination, laboratory investigations (Hb, serum level of urea before dialysis, urea after dialysis, urea reduction ratio, creatinine, calcium, phosphorus, albumin and C reactive protein, TSH and FT4) were obtained from all study patients. TSH and FT4 were measured by chemiluminescence immunoassay using the Modular Analytics E170 immunoassay analyzer (Roche Diagnostic, Basel, Switzerland). Diagnosis of SCH was defined as high TSH level but normal free T4 level. The reference range for free T4 was 0.93- 1.7 ng/dl and for TSH 0.27-4.2 μ U/ml. The intra-assay coefficient of variation for these hormones ranged from 1.4% to 2.0% for free T4 and 3.0% to 1.1% for TSH.

Informed consent was obtained from the entire study group and the study was approved by Faculty of Medicine, Helwan University Research Ethics Committee.

Statistical Analysis: Data were fed to the computer and analyzed using IBM SPSS software package version 20.0. (Armonk, NY: IBM Corp). The Kolmogorov- Smirnov test was used to verify the normality of distribution of variables. Comparisons between groups for categorical variables were assessed using Chi-square test (Fisher or Monte Carlo). Student t-test was used to compare two groups for normally distributed quantitative variables. Mann Whitney test was used to compare between two groups for not normally distributed quantitative variables, while Wilcoxon signed ranks test was assessed for comparison between different periods. Univariate and Multivariate logistic regression was assessed. Significance of the obtained results was judged at the 5% level.

Results: Of the 92 ESRD patients recruited, the mean age was 51.4 ± 12.8 years, 38 (41.3%) were males, 54 (58.7%) were

females, 19 (20.7%) were smokers and the mean dry weight was 72.2 ± 15.7 kilogram. The baseline characteristics and different laboratory parameters are listed in table II and III. Based on the TSH level, the studied group was divided into two groups; subclinical

hypothyroidism group and normal TSH, normal FT4 group. 80 patients (86.9%) of the total studied group showed normal TSH and normal free T4. 12 patients (13.1%) showed high TSH $> 4.2 \mu\text{IU/ml}$ and normal free T4. The prevalence of SCH was 13.1%. (Table I)

Table (I): Distribution of the studied cases according to TSH level

	Total (n = 92)	Normal (≤ 4.2) n = 80 (86.9%)	Subclinical hypothyroidism (> 4.2) n = 12 (13.1%)
TSH			
Mean \pm SD.	2.8 \pm 1.7	2.2 \pm 0.9	6.4 \pm 1.4
Median (Min. – Max.)	2.4(0.7–8.9)	2.3(0.7–4.1)	6.4(4.6–8.9)

(n : number)

Table (II): Baseline demographic and clinical characteristics of the patients

	Total (n=92)	TSH		Test of sig.	p
		Normal (n=80)	Subclinical hypothyroidis m (n=12)		
Sex					
Male	38 (41.3%)	34 (42.5%)	4 (33.3%)	$\chi^2=$	^{FE} p=
Female	54 (58.7%)	46 (57.5%)	8 (66.7%)	0.362	0.755
Age (years)					
Median (Min.–Max.)	52.5 (21 – 77)	53.0 (21 – 77)	47.5 (22 – 68)	t=0.946	0.346
Mean \pm SD.	51.4 \pm 12.8	51.9 \pm 12.7	48.2 \pm 13.5		
Weight (kg)					
Median (Min.–Max.)	70.5(45 – 118)	70.0 (45 – 112.5)	74.5 (47 – 118)	t=0.891	0.390
Mean \pm SD.	72.2 \pm 15.7	71.4 \pm 14.3	77.6 \pm 23.3		
Current smoking					
Negative	73 (79.3%)	63 (78.8%)	10 (83.3%)	$\chi^2=$	^{FE} p=
Positive	19 (20.7%)	17 (21.3%)	2 (16.7%)	0.134	1.000
Underlying kidney disease					
Unknown	18(19.6%)	17 (21.3%)	1(8.3%)	$\chi^2=1.106$	^{FE} p=0.448
HTN	34 (37%)	27 (33.8%)	7 (58.3%)	$\chi^2=2.707$	^{FE} p=0.118
D.N	27 (29.3%)	25 (31.3%)	2 (16.7%)	$\chi^2=1.070$	^{FE} p=0.498
G.N	2 (2.2%)	2 (2.5%)	0 (0%)	$\chi^2=0.307$	^{FE} p=1.000
CH pyelonephritis	2 (2.2%)	2 (2.5%)	0 (0%)	$\chi^2=0.307$	^{FE} p=1.000
Lupus nephritis	6 (6.5%)	3 (3.8%)	3 (25%)	$\chi^2=7.729^*$	^{FE} p=0.028*
Obstructive uropathy	8 (8.7%)	8 (10%)	0 (0%)	$\chi^2=1.314$	^{FE} p=0.590
PCKD	2 (2.2%)	2 (2.5%)	0 (0%)	$\chi^2=0.307$	^{FE} p=1.000
Renal amyloidosis	2 (2.2%)	2 (2.5%)	0 (0%)	$\chi^2=0.307$	^{FE} p=1.000
Sepsis	1 (1.1%)	1 (1.3%)	0 (0%)	$\chi^2=0.152$	^{FE} p=1.000
Vasculitis	1 (1.1%)	1 (1.3%)	0 (0%)	$\chi^2=0.1s52$	^{FE} p=1.000
Duration of dialysis(month)					
Median (Min.–Max.)	13.0(1.0– 156.0)	12.0(1.0– 125.0)	48.5(1–156)	U= 278.5*	0.019*
Mean \pm SD.	26.3 \pm 31.2	21.9 \pm 25.3	55.8 \pm 48.6		

χ^2 : Chi square test, FE: Fisher Exact, t: Student t. test, U: Mann Whitney test

p: p value for comparing between the two groups

*: Statistically significant at $p \leq 0.05$

Normal: TSH \leq 4.2 , **Subclinical hypothyroidism: TSH $>$ 4.2**

Table (III): Blood pressure and different laboratory parameters for the total group and two studied groups

	Total (n=92)	Normal (n=80)	TSH Subclinical hypothyroidism (n=12)	Test of sig.	p
Blood pressure					
Systolic					
Median (Min.–Max.)	120 (90 – 160)	120 (90 – 160)	130 (100 – 160)	t=0.894	0.374
Mean ±SD.	125.1 ± 20.3	124.4 ± 20.5	130 ± 19.1		
Diastolic					
Median (Min.–Max.)	80 (50 – 100)	80 (50 – 100)	80 (60 – 90)	t=0.394	0.694
Mean ±SD.	78.7 ± 11.2	78.9 ± 11.5	77.5 ± 9.7		
Hemoglobin					
Median (Min.–Max.)	9.1 (5.9 – 14.1)	9.2 (5.9 – 14.1)	9.1 (7.3 – 10.8)	t=0.698	0.487
Mean ±SD.	9.2 ± 1.8	9.3 ± 1.9	8.9 ± 1.2		
Urea					
Before					
Median (Min.–Max.)	154.5 (47–294)	160 (50 – 294)	131.5 (47 – 240)	U=384.0	0.266
Mean ±SD.	157.8 ± 49.4	160.6 ± 45.7	139.3 ± 68.7		
After					
Median (Min.–Max.)	39 (11 – 124)	39 (11 – 124)	43.5 (15 – 96)	U=457.50	0.794
Mean ±SD.	48.3 ± 24.8	47.7 ± 23.8	51.9 ± 31.9		
p ₁	<0.001*	<0.001*	0.002*		
URR%					
Median (Min.–Max.)	68.9 (6.6–89.9)	69.6 (6.6 – 89.9)	68.4 (49.7 – 78.8)	t=0.616	0.540
Mean ±SD.	68.9 ± 13	69.2 ± 13.6	66.7 ± 7.8		
Cr (mg/dl)					
Median (Min.–Max.)	9.9 (2.3 – 19.8)	10.1 (2.3 – 19.8)	9.2 (2.5 – 16.6)	t=1.524	0.131
Mean ±SD.	10.2 ± 3.3	10.4 ± 3.2	8.9 ± 3.9		
Ca (mg/dl)					
Median (Min.–Max.)	9.2 (6.2 – 12)	9.1 (6.2 – 12)	9.2 (7.3 – 12)	t=0.657	0.513
Mean ±SD.	9.1 ± 1.2	9.1 ± 1.2	9.3 ± 1.4		
Ph (mg/dl)					
Median (Min.–Max.)	5.7 (2.7 – 10.7)	5.8 (2.7 – 10.7)	4.8 (4.4 – 7)	t=1.707	0.091
Mean ±SD.	5.7 ± 1.4	5.8 ± 1.4	5.1 ± 0.8		
Albumin (g/dl)					
Median (Min.–Max.)	3.9 (2.5 – 5.2)	3.9 (2.5 – 5.2)	4.1 (3 – 5.2)	t=0.390	0.697
Mean ±SD.	3.9 ± 0.6	3.9 ± 0.5	4 ± 0.7		
Free T4					
Median (Min.–Max.)	1.2 (1 – 1.7)	1.2 (1 – 1.7)	1.3 (1 – 1.5)	t=0.130	0.897
Mean ±SD.	1.3 ± 0.1	1.3 ± 0.1	1.3 ± 0.2		
CRP					
Median (Min.–Max.)	5.0(3.0 – 15.0)	5.0(3.0 – 12.0)	9.0(8.0 – 15.0)	U= 41.50*	<0.001*
Mean ±SD.	6.1 ± 2.4	5.5 ± 1.7	10.3 ± 2.8		

t: Student t-test U: Mann Whitney test

p: p value for comparing between the two groups

p₁: p for Wilcoxon signed ranks test for comparing between before and after

*: Statistically significant at p ≤ 0.05

Normal: TSH ≤ 4.2 , **Subclinical hypothyroidism: TSH > 4.2**

Patients with SCH had significantly higher duration of dialysis and higher level of CRP compared to patients with normal TSH. (p value = 0.019, <0.001 respectively). Also, the lupus nephritis as an underlying kidney disease was significantly higher in the SCH group compared to the normal TSH group (p= 0.028).

By studying the independent variables affecting SCH by assessing multivariate analysis (table IV), it was found that higher duration of dialysis and CRP were significantly associated with SCH in both unadjusted and adjusted multivariate analysis.

Table (IV): Univariate and multivariate analysis for the parameters affecting TSH (n = 92) for total group

TSH	Univariate		#Multivariate		#Multivariate (adjusted age and sex)	
	p	OR (95%C.I)	p	OR (95%C.I)	p	OR (95%C.I)
Lupus nephritis	0.016*	8.56(1.50 – 48.89)	0.563	2.04(0.18 – 22.98)	0.467	2.71(0.18 – 40.00)
Duration of dialysis (month)	0.003*	1.03(1.01 – 1.04)	0.029*	1.03(1.00 – 1.06)	0.020*	1.03(1.00 – 1.06)
CRP	<0.001*	2.57(1.56 – 4.25)	0.001*	2.56(1.48 – 4.41)	0.003*	2.90(1.45 – 5.79)

OR: Odd's ratio, C.I: Confidence interval,

#: All variables with p < 0.05 were included in the multivariate

*: Statistically significant at p ≤ 0.05

Discussion:

The importance of understanding the thyroid functions in CKD & ESRD patients is highlighted by recent studies indicating that subclinical hypothyroidism is common in CKD patients.⁽⁹⁾ Our results had shown a prevalence rate for SCH in ESRD equal to 13.1% which is concordant with Lim⁽¹⁰⁾ who identified a prevalence rate 9.5 % for SCH in ESRD. Chonchol et al⁽¹¹⁾ also studied the prevalence of SCH in CKD patients but not requiring dialysis and they showed 18% prevalence rate.

Eun et al⁽⁵⁾ found that unresolved SCH was significantly associated with a more rapid decline in renal function and unresolved SCH was an independent factor contributing to the rate of estimated glomerular filtration rate (e GFR) decline and that these individuals had a higher risk of progression to ESRD. Similarly, Khatiwada et al found abnormal thyroid functions in CKD patients and SCH to be associated with CKD progression.⁽¹²⁾

It is known that there is alteration in thyroid hormone physiology in CKD. Baseline TSH may be elevated, there is blunted response to

exogenous TRH, disturbed diurnal rhythm of TRH and sometimes there is an observed reduction in serum T4 level.⁽¹³⁾

Inflammation is known to be more prevalent in SCH than in euthyroidism.^(14, 15) In our study, we identified that CRP was significantly high in SCH group and this is supported by what is mentioned by Zoccali et al who found that biomarkers of inflammation as CRP, IL6 are consistently associated with low T3 levels in ESRD patients on HD.⁽¹⁶⁾ Also, experimental and clinical studies indicate that inflammatory cytokines play a central role in low T3 syndrome.⁽¹⁷⁾

What was also noticed that duration of dialysis was longer in SCH patients in comparison to normal TSH and T4 group. In 2011, Yee et al⁽¹⁸⁾ found that in peritoneal dialysis (PD) duration was a significant risk factor for SCH and it was associated with longer PD duration and this may be due to retention of excess iodide and uremic toxins.^(19,20)

There have been controversies about thyroxine supplementation for SCH in ESRD. Decreased thyroid function can be considered as an adaptation to decrease protein

catabolism in ESRD. Hence, thyroxine replacement in ESRD patients may lead to a negative nitrogen balance.⁽²¹⁾

Conclusions: From previous data, it was found that the prevalence rate of SCH among HD patients in Egypt is 13.1% and we suggest that there is an association between inflammatory markers and SCH. We suggest also that SCH is associated with longer HD duration.

Recommendations: Large trials and follow up may be needed to solve the controversies regarding thyroxin replacement in case of SCH in ESRD. Also, the relation of lupus nephritis as a cause of ESRD and SCH in HD patients needs further research.

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Study of Prevalence and Effects of Multiple Drug Intake in Elderly Type 2 Diabetes Patients with Associated Comorbidities

Mohammed Badr, Abdelrahman Kamel, Mai Badrah, Reem Fathalla

Department of Internal Medicine, Faculty of Medicine, Alexandria University, Egypt.

Abstract

Background: Elderly patients with diabetes are a vulnerable group especially for associated comorbidities. The physicians should be aware of the pitfalls of caring for these patients and prioritize an individualized treatment plan to ensure an optimal glycemic control, without placing the patient at unnecessary risk. **Objective:** Assess the effects of multiple drug intakes in elderly type 2 diabetes patients with associated comorbidities. **Patients & Methods:** This study included 100 patients with type 2 diabetes and other associated comorbidities recruited from the outpatient clinic of Internal Medicine Department, Alexandria Main University Hospital. The studied patients were divided in to two groups: Group I; 50 patients aged 55-64 years and group II: 50 patients ≥ 65 years. Each group was further re-divided to (A) subdivision: patients who were receiving < 5 drugs and (B) subdivision: patients who were receiving ≥ 5 drugs. The effects of polypharmacy in elderly type 2 diabetes patients

with comorbidities especially on drug adherence and the observed adverse drug events were assessed. **Results:** Our results demonstrated that non adherence to drugs was significantly higher in patients who received ≥ 5 drugs in group I compared to those who received < 5 drugs. In comparison between the polymedicated groups (IB and IIB), non adherence was significantly higher in IIB group. Among the studied adverse drug events, arrhythmia, repeated hypoglycemia, orthostatic hypotension, GIT upsets and bleeding events were significantly higher in group (IIB) than group (IB). **Conclusion:** Prescription in elderly diabetics must balance between the need for drugs and hazards of polypharmacy. Medication non-adherence and adverse drug events are major consequences of polypharmacy.

Corresponding Author: Mai Hesham Badrah, E-mail: drmaibadrah@gmail.com

Key Words: Polypharmacy, Type 2 DM, Elderly population.

INTRODUCTION

Recent advances in pharmacological therapy, although offer hope for patients, its inappropriate use carry some risks. ⁽¹⁾ Prescription for the elderly is particularly challenging owing to the high prevalence of comorbidities in these population. ⁽²⁾ In general, polypharmacy means the use of multiple drugs without clear cutoff point. ⁽³⁾ A more practical definition for polypharmacy entails

the use of multiple drugs that are not obviously needed. ⁽⁴⁾

Diabetes is a highly prevalent systemic disease with lots of complications including microvascular complications (retinopathy, nephropathy, neuropathy and dermopathy) and macrovascular ones (cerebrovascular diseases, peripheral vascular diseases and coronary heart diseases).⁽⁵⁾ Many comorbid conditions are frequently encountered in elderly patients with diabetes like hypertension and

dyslipidemia in addition to the high incidence of geriatric syndromes like fall, depression and cognitive dysfunction. ⁽⁶⁾ That's why management of diabetes with its complications and associated comorbidities make these patients prone to hazards of polypharmacy.

PATIENTS & METHODS

This study included 100 patients with type 2 diabetes associated with other comorbidities recruited from the Outpatient Clinic of the Department of Internal Medicine of the Main University Hospital of Alexandria during the period of 2017-2018. The patients were classified into 2 groups each of 50 patients; group I included patients aged between 55 and 64 years and group II included patients aged ≥ 65 years. Each group was further subdivided in relation to total number of drug intake in to (A) subdivision including patients receiving < 5 drugs and (B) subdivision including patients receiving ≥ 5 drugs (polymedicated).

The study protocol was approved by the Ethics Committee of Alexandria University. The study was conducted according to the criteria set by the Declaration of Helsinki and each subject signed an informed consent before participating in the study.

All cases were subjected to complete history taking with emphasis on drug history. Data were obtained about the presence of new symptoms due to drug intake with differentiation whether they were adverse drug events or drug-drug interaction. An Adverse Drug Event (ADE) is defined as 'any physical or mental harm resulting from medication use be it misuse, under-dosing or overdosing' ⁽¹¹⁾.

A drug interaction is defined either as increase or decrease of a medical diagnostic or therapeutic effect of a specific drug caused by another substance, which may be another drug, plant or a dietary supplement ⁽¹²⁾. Full clinical examination was done to all included patients. In addition, glycated hemoglobin (HbA1C %) was done to assess glycemic control.

Statistical analysis

Data were analyzed using Statistical Package for the Social Sciences (SPSS 21.0, IBM/SPSS Inc., Chicago, IL) software. Baseline characteristics of the study population were presented as frequencies and percentages (%) or mean values and standard deviations (SD) and median and range (after testing of normality by Kolmogorov-Smirnov and Shapiro-Wilk's tests).

For comparison of data, Chi-Square test (or Fisher's exact test) was used to compare two independent groups of qualitative data. For quantitative data, independent-Samples t-test and Mann-Whitney U test were used to compare two groups of parametric and non-parametric quantitative data respectively. For all tests, P values < 0.05 are considered significant.

RESULTS:

Table I shows the baseline characteristics of the studied population. There was no significant difference in terms of male/female ratio. Body mass index (BMI) was significantly higher in group I. The mean diabetes duration was 7.04 ± 4.97 and 12.14 ± 6.53 in group I and II respectively.

Table I: Comparison between the two studied groups according to demographic data

	Total (n=100)		Age (years)				Test of Sig.	P
			Group I (n = 50)		Group II (n = 50)			
	No.	%	No.	%	No.	%		
Sex								
Male	52	52.0	28	56.0	24	48.0	$\chi^2=0.641$	0.423
Female	48	48.0	22	44.0	26	52.0		
BMI (kg/m²)								
Min. – Max.	23.0 –36.0		26.0 –34.0		23.0 –36.0		t=2.315*	0.023*
Mean ± SD.	29.80 ±2.21		30.30 ±1.92		29.30 ±2.38			
Median (IQR)	30.0 (28.0 –31.0)		30.0 (29.0 – 32.0)		29.0 (32.0 – 31.0)			
Diabetes duration								
Min. – Max.	1.0 –31.0		1.0 –25.0		31.0		U=642.0*	<0.001*
Mean ± SD.	9.59 ±6.32		7.04 ±4.97		12.14 ±6.53			
Median (IQR)	9.0 (4.0 –14.0)		7.0 (3.0 – 10.0)		12.50 (7.0 – 15.0)			

χ^2 : Chi square test t: Student t-test
 p: p value for comparing between the studied groups
 *: Statistically significant at $p \leq 0.05$

The detected comorbidities in the included patients were: hypertension (HTN), dyslipidemia, arthropathies, gout, ischemic heart disease (IHD), stroke, cognitive problems, urinary incontinence and hepatitis C virus (HCV). Among them, arthropathies, IHD and stroke were significantly

higher in group II ($p=0.003$, $p<0.001$, $p=0.006$) respectively.

As shown in table II 80% of the studied population were receiving ≥ 5 drugs. The number of patients who were receiving ≥ 5 drugs (polymedicated patients) were significantly higher in group II.

Table II: Comparison between the two studied groups according to total number of drugs

Total number of drugs	Total (n=100)		Age (years)				Test of sig.	P
			Group I (n = 50)		Group II (n = 50)			
	No.	%	No.	%	No.	%		
<5 (A)	20	20.0	16	32.0	4	8.0	$\chi^2=9.0^*$	0.003*
≥ 5 (polymedicated (B))	80	80.0	34	68.0	46	92.0		
Min. – Max.	3.0–14.0		4.0–14.0		3.0–13.0		U=749.0*	<0.001*
Mean ± SD.	6.98±2.58		6.18±2.46		7.78±2.47			
Median (IQR)	7.0(5.0–9.0)		5.0(4.0–8.0)		7.0(6.0–10.0)			

U: Mann Whitney test
 p: p value for comparing between the studied groups
 *: Statistically significant at $p \leq 0.05$

The relation between the total number of drugs and drug adherence is demonstrated in Table III. Non adherence to drugs was significantly higher in patients who received ≥ 5 drugs in group I

compared to those who received < 5 drugs. In comparison between the polymedicated groups (IB and IIB), non-adherence was significantly higher in IIB group.

Table III: Relation between total number of drugs and adherence

	Total number of drugs							
	Group I (n = 50)				Group II (n = 50)			
	<5 (IA) (n = 16)		≥ 5 (IB) (n = 34)		<5(IIA) (n = 4)		≥ 5 (IIB) (n = 46)	
	No.	%	No.	%	No.	%	No.	%
Adherence								
Yes	16	100.0	15	44.1	2	50.0	8	17.4
Non adherence	0	0.0	19	55.9	2	50.0	38	82.6
$\chi^2(p_1)$	14.421*(<0.001*)				2.446 (FEp=0.174)			
$\chi^2(p_2)$	6.817* (0.009*)							

χ^2 : Chi square test, FE: Fisher Exact

p_1 : p value for comparing between < 5 and ≥ 5 in each group

p_2 : p value for comparing between group I and II for ≥ 5 group

*: Statistically significant at $p \leq 0.05$

Regarding the adverse drug events there was significant difference in the incidence of arrhythmia, repeated hypoglycemia, orthostatic hypotension, gastrointestinal (GIT) upsets and bleeding tendency between the patients in the two studied groups being higher in the older patients (group II) ($p = 0.007, 0.012, 0.007, 0.001, 0.008$) respectively. Other adverse events did not reveal significant difference between the two groups.

Table (IV) shows the relation between the total number of drugs and the detected drug adverse events in the studied groups. Arrhythmia, repeated hypoglycemia, orthostatic hypotension, GIT upsets and bleeding events were significantly higher in elderly polymedicated group (IIB) than polymedicated patients less than 65 year old (IB) group.

Table IV: Relation between total number of drugs and adverse drug events

Adverse drug effect	Total number of drugs							
	Group I (n = 50)				Group II (n = 50)			
	<5 (IA) (n = 16)		≥5 (IB) (n = 34)		<5 (IIA) (n = 4)		≥5 (IIB) (n = 46)	
	No.	%	No.	%	No.	%	No.	%
Arrhythmia	0	0.0	2	5.9	0	0.0	11	23.9
$\chi^2(p_1)$	0.980 (FEp=1.000)				1.226 (FEp=0.563)			
$\chi^2(p_2)$	4.670*(0.031*)							
Repeated hypoglycemia	0	0.0	0	0.0	0	0.0	7	15.2
$\chi^2(p_1)$	-				0.708 (FEp=1.000)			
$\chi^2(p_2)$	5.670*(FEp=0.019*)							
Orthostatic Hypotension	0	0.0	2	5.9	0	0.0	11	23.9
$\chi^2(p_1)$	0.980 (1.000)				1.226 (FEp=0.563)			
$\chi^2(p_2)$	4.670*(0.031*)							
GIT upsets	6	37.5	13	38.2	2	50.0	34	73.9
$\chi^2(p_1)$	0.002 (0.960)				1.044 (0.310)			
$\chi^2(p_2)$	10.269*(0.001*)							
Bleeding tendency	0	0.0	1	2.9	0	0.0	9	19.6
$\chi^2(p_1)$	0.480 (FEp=1.000)				0.954 (1.000)			
$\chi^2(p_2)$	4.940*(0.038*)							
Hyperuricemia	0	0.0	4	11.8	0	0.0	4	8.7
$\chi^2(p_1)$	2.046 (FEp=0.292)				0.378 (FEp=1.000)			
$\chi^2(p_2)$	0.205(FEp=0.717)							
Dry cough	0	0.0	0	0.0	0	0.0	4	8.7
$\chi^2(p_1)$	-				0.378 (FEp=1.000)			
$\chi^2(p_2)$	3.112 (FEp=0.133)							
Lower limb edema	0	0.0	0	0.0	0	0.0	5	10.9
$\chi^2(p_1)$	-				0.483(FEp=1.000)			
$\chi^2(p_2)$	3.942(FEp=0.069)							
Dry mouth	0	0.0	0	0.0	0	0.0	1	2.2
$\chi^2(p_1)$	-				0.089 (FEp=1.000)			
$\chi^2(p_2)$	0.784 (FEp=1.000)							
Headache	0	0.0	6	17.6	0	0.0	4	8.7
$\chi^2(p_1)$	3.209 (FEp=0.159)				0.378(FEp=1.000)			
$\chi^2(p_2)$	1.432(FEp=0.310)							

χ^2 : Chi square test, FE: Fisher Exact

p₁: p value for comparing between <5 and ≥ 5 in each group

p₂: p value for comparing between group I and II for ≥5 group

*: Statistically significant at p ≤ 0.05

Regarding glycemic control, the mean glycosylated hemoglobin (HbA1c) were significantly higher in IB (8.26 ± 0.83) than IA subdivision (7.46 ± 0.55) of group I (P=0.001), while there was no significant difference between IIA and IIB subdivisions. On comparing the polymedicated subdivisions (IB, IIB) in both groups, no significant difference was detected regarding the mean HbA1c (p=0.290).

DISCUSSION:

Diabetes Mellitus (DM) is an important health issue for the elderly population. (7) More than half of patients with diabetes mellitus have two or more associated diseases. (8,9) Multimorbidity, commonly characterized as the co-existence of two or more chronic health conditions, is common within the elderly. (10) Polypharmacy is defined as "the administration of many drugs at the same time or the administration of an excessive number of drugs". (11) But unfortunately, there's no standard cut point with respect to the number of drugs that's

agreed upon for the definition. The foremost commonly utilized term was polypharmacy which was characterized as five or more drugs. ⁽¹²⁾ Comprehensive medication review and hazard evaluation ought to be carried out by interdisciplinary team to distinguish the polypharmacy and its unfavorable impacts. ⁽¹³⁾

Medication nonadherence may be a major cause of morbidity, particularly in elderly patients. Roughly 10% of hospitalizations may be a result of medicine nonadherence. ⁽¹⁴⁾ Our results demonstrated that non adherence to drugs was significantly higher in patients who received ≥ 5 drugs in group I compared to those who received < 5 drugs. In comparison between the polymedicated groups (IB and IIB), non-adherence was significantly higher in IIB group.

Pasina L. et al. ⁽¹⁵⁾ reported that low medicine adherence may be a real, complex issue for geriatric patients receiving polypharmacy. They found that the increasing number of drugs endorsed at hospital discharge is related to non-adherence and a high rate of patients did not understand the reason of their medicines. Simplification of drug regimens and decrease of pill burdens as well as way better clarifications of the reason for the medicines ought to be targets for intervention. Delamater ⁽¹⁶⁾ reported that among the factors associated with medication adherence, adherence with a simple prescription is higher than that for a more complex prescription.

On the contrary, Grant RW et al. ⁽¹⁷⁾ which demonstrated that in their study sample, patients reported very high medicine adherence rates regardless of number of medications endorsed. Among patients on numerous drugs, most patients with suboptimal adherence were perfectly adherent to all but one medication. Unreported side effects and a lack of confidence in immediate or future benefits were significant indicators of suboptimal adherence. Doctors ought to not feel hindered from endorsing multiple agents in arrange to achieve

satisfactory control of hyperglycemia, hypertension, and hyperlipidemia.

Moreover, Cárdenas-Valladolid et al ⁽¹⁸⁾ study revealed that poor therapeutic adherence in homebound elderly patients receiving polypharmacy may be a serious issue influencing one of each three individuals concerned, and is directly related to caregiver burden, regardless of age, gender, cognitive status or number of drugs administered.

The variation in results were due to problems with tools for the evaluation of adherence as different studies used different tools such as questionnaires, tests, pill count or judgment by research nurse. ⁽¹⁸⁻²¹⁾ The same variability was in defining polypharmacy, where the researchers used different definitions. ^(20,22)

Adverse drug reactions happen nearly day by day in health care institutions and can unfavorably influence a patient's quality of life, often causing significant morbidity and mortality. ⁽²³⁾ Regarding the adverse drug events, our results showed that there was significant difference in the incidence of the following ADEs; arrhythmia, repeated hypoglycemia, orthostatic hypotension, GIT upsets and bleeding tendency between the patients in the two studied groups being higher in the older patients (group II) ($p= 0.007, 0.012, p=0.007, 0.001, 0.008$) respectively. Moreover, these ADEs were significantly higher in elderly polymedicated group (IIB) than (IB) group.

These results go hand in hand with the findings of various studies carried out by Rajesh et al. and Pirmohamed et al who reported that the percentage of ADRs found was higher with age and geriatric population. ^(24,25) Akhideno P et al. ⁽²⁶⁾ showed that the elderly age group, presence of polypharmacy and the presence of multiple comorbidities found to be associated with and predisposed to ADRs in this study. Moreover, Shareef J et al. ⁽²⁷⁾ revealed that the geriatric patients with diabetes had more

associated comorbidities which forces them to get multiple medications leading to more ADRs. In addition, Subeesh VK et al. ⁽²⁸⁾ revealed that the prevalence of polypharmacy is very high among geriatric population in the study site. The study supported the consequences of polypharmacy and was closely related with multiple comorbidity and advanced age.

On the contrary, Min Zhang et al. ⁽²⁹⁾ in their study of variables that anticipate repeat admission for ADRs in geriatric population uncovered that comorbidity, but not advancing age, predicts repeat admission for ADRs.

Khan LM et al. ⁽³⁰⁾ in his retrospective and prospective studies detailed that the highest incidence of ADR (retrospective 15% and prospective 14.5%) was watched in both groups in patients getting more than 10 medications. The frequency of ADR in relation to age in both groups was highest in patients of age >60 years; it was 52.7% in retrospective study and 54.5% in prospective study. The system most commonly included in ADR was gastrointestinal tract 47.4% in retrospective study and 57.6% in prospective study. This studies recognized gastrointestinal system as the foremost habitually affected system by ADRs in both studies.

Conflict of Interest:

We confirm that there are no known conflicts of interest associated with this publication

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The Effect of 1, 25 Dihydroxy-Vitamin D3 on Insulin Sensitivity and Expression of JNK in Diabetes - Induced Liver Complications in Rats

Kawkab Elsbah Mohamed Ragab, Hala Salah Ibrahim, Gehan Yassin Soliman, Rasha Mohamed Adel Nassra, Aml Gomaa Ali Kobbas

Department of Medical Physiology, Faculty of Medicine, University of Alexandria

Abstract

Introduction: NAFLD one of most common complication of diabetes mellitus. Several studies tried to understand its mechanism or to prevent it. The current study aim is to study the possible effect of treatment of vitamin D on the expression of inflammatory cytokine C-Jun N-terminal kinase (JNK) in rat models of diabetic liver. **Methods:** thirty male albino rats were assigned into two groups group I (healthy control) that were fed normal diet ($n= 10$), group II ($n=20$) received HFD for 8 week and then injected with STZ 30

mg/kg. After detection of hyperglycemia this group was randomly divided into subgroups, group IIA received vitamin D for 8 weeks in a dose of ($0.03 \mu\text{g/kg/day}$), and group IIB was kept without treatment. **Results & Conclusion:** vitamin D significantly improves glucose intolerance, insulin sensitivity, decreased HOMA-IR, decreased serum ALT and decreased AST, increased SOD and decreased liver JNK expression.

Key Words: diabetic liver, NAFLD, JNK, 1,25dihydroxy vitamin D3, insulin sensitivity, SOD, Interleukine-1 β .

Introduction

Diabetes mellitus affects people worldwide and poses major public health and socio-economic challenges.⁽¹⁾ Monitoring of blood glucose is essential in optimizing long-term outcomes towards reducing the co-morbid conditions that may arise in patients with T2DM⁽²⁾. High Fat Diet (HFD) has also been utilized to model chronic inflammation, which is an important pathogenic mechanism of T2DM. The HFD - mediated chronic inflammation is marked by increased TNF- α , IL-1 β , and IL-6 in the circulation.⁽³⁾

T2DM is strongly associated with obesity and insulin resistance⁽⁴⁾ as well as defects in pancreatic β -cell function and mass. Insulin is important in carbohydrate, fat and protein metabolism⁽⁵⁾.

Insulin resistance acts as a key link between obesity and T2DM, It is likely that the development of obesity-induced insulin resistance involves a complex interplay of genetic and environmental factors.⁽⁶⁾ The Insulin Receptor (IR) signaling pathway is triggered by its binding of the IR a receptor that belongs to tyrosine kinase receptor family. Hormone engagement stimulates the intrinsic tyrosine kinase activity of the IR resulting in the auto phosphorylation on the β subunit cytoplasmic tails.⁽⁷⁾

JNK: JNKs are serine threonine protein kinases that phosphorylate Jun family members, components of activator protein 1 (AP-1) transcription factors JNK also known as stress-activated Mitogen Protein Kinase (AMPK).⁽⁸⁾

Complication of diabetes: Chronic complications of diabetes which involve coronary artery disease, renal, liver and ophthalmologic diseases are the primary cause of disability and mortality in DM patients. It is reported that the standardized mortality rate from end-stage liver disease (i.e. cirrhosis) is higher than that for cardiovascular disease among patients with diabetes.⁽⁹⁾

Liver complications: NAFLD the most common chronic hepatopathy worldwide, reaching a prevalence of above 70% in patients with T2DM.⁽¹⁰⁾ Inflammation is a crucial part of almost all acute and chronic liver disorders including (NAFLD).⁽¹¹⁾

Interleukine-1 β : IL-1 β is a Pro-inflammatory cytokine not expressed in healthy liver. It acts via specific receptors involved in all inflammatory processes in the liver including regulation of insulin resistance and fibrosis.⁽¹²⁾

Oxidative stress is defined as an imbalance in the oxidant-to-antioxidant ratio, causing the generation of free radicals⁽¹³⁾. **Super oxide-dismutase (SOD):** SOD is a key antioxidant enzyme that dismutates superoxide anion (O₂⁻) to continuously form H₂O₂ and O₂.⁽¹⁴⁾

Vitamin D: Vitamin D is a lipophilic molecule essential to maintain calcium and phosphate balance and osteometabolic system regulation. The steroid hormone Vitamin D (Vit D) can be produced by the body or obtained through the diet.⁽¹⁵⁾ Evidence from various directions, including observational and experimental studies, indicates the protective capacity of 1,25-dihydroxyvitamin D3 (active vitamin D) in chronic low-intensity inflammation and IR in T2DM and its complications; although the underlying mechanism is yet to be clarified⁽¹⁶⁾.

Aim of the Work

The purpose of the present study is to develop an appropriate stable animal model which

is analogous to the human T2DM through a combination of high-fat diet with a low-dose STZ injection to test the hypothesis that vitamin D may improve the diabetes-induced liver complications by suppressing the expression of inflammatory cytokine and C-Jun N-terminal kinase (JNK) in the liver of this rat models of diabetes.

Materials & Methods

The study was conducted on thirty healthy male albino rats with body weight (100–150 g). Rats were housed in standard polypropylene cages (three rats/cage).⁽¹⁷⁾ The period of experiment was 16 weeks. The Experiment was conducted in accordance with the guidelines set by the Research Ethics Committee of the Faculty of Medicine Alexandria University. After a week of acclimatization, rats were randomly divided into two groups:

Group (I): [(Control):(n=10)] rats in this group were fed a regular standard chow diet (60% carbohydrates, 5% fat and 18% protein).⁽¹⁸⁾

Group(II): (n=20) rats of this group were fed a high fat diet HFD (22% fat, 48% carbohydrate, 20% protein) with a total caloric value of 44.3 kcal /kg.⁽¹⁷⁾

After 8 weeks from the start of the diet regimen, all rats of group II received a single low dose of STZ 30 mg/kg (dissolved in 0.1 ml sodium citrate buffer at pH (4.4) intra peritoneal injection (IP).⁽¹⁷⁾

Rats of the Control group (Group I) received a single IP injection of ml of sodium citrate buffer only. One week later, random blood glucose level of all rats was measured in blood samples collected from tail vein, using a glucometer.⁽¹⁹⁾ Diabetes was confirmed by the detection of hyperglycemia (random non-fasting glucose level, >300mg/dl),⁽²⁰⁾ T2DM model rats (n=10) were randomly divided equally into 2 groups IIA and IIB each of 10 .

Group IIA (Vit D treated group):(n=10)

Ten diabetic rats received vitamin D (1,25-dihydroxyvitamin D₃) with a dose of (0.03 µg/kg/day) dissolved in corn oil, administered orally using a gavage needle. The dose was 1 µg =40 IU and this treatment was continued for 8 weeks.⁽²¹⁾

Group IIB (untreated group): (n=10)

These ten diabetic rats received the vehicle oral gavage per day for (8 weeks) .

Both control and diabetic untreated rats received an equivalent administration of corn oil for 8 weeks.⁽²¹⁾

Body Weight of rats was measured at the beginning of the experiment and at week 8 and then after 16 weeks at the end of the experiment.

At the end of the experiment, rats were fasted overnight and measuring the weight plexus of the rat by capillary under light ether anesthesia.⁽²²⁾ The serum was separated by centrifugation at 3000 rpm for 15 minutes and was stored at -20°C until assayed for measuring the following parameters:

1. Fasting blood glucose level by colorimetric method.⁽¹⁹⁾

2. Serum insulin level by enzyme linked immunosorbent assay (ELISA).⁽²³⁾
3. Insulin resistance via the (HOMA-IR).⁽²³⁾
4. Serum Liver enzymes; alanine aminotransferase (ALT) and aspartate aminotransferase (AST) activity by colorimetric method.⁽²⁴⁾
5. Serum super oxide dismutase (SOD) activity by colorimetric method.⁽²⁵⁾
6. Serum IL1β level by ELISA.⁽²⁶⁾

Liver tissue preparation

The rats were sacrificed with decapitation, the anterior abdomen was incised to expose the liver. The liver from each rat was excised immediately after perfusion and rinsed with ice-cold saline solution. The ventral median lobe of the liver was fixed in 10% neutral buffered formalin for histopathological study. The remaining portion was homogenized in 0.9% saline, centrifuged at 400×g for 15 minutes and the supernatant was decanted and stored at 70°C until used for 7.Reverse transcription real time quantitative-polymerase chain reaction of (RT-qPCR) was performed on liver tissue to quantify gene expression level of JNK.^(27, 28)

Results

Table (I): Comparison between the three studied groups regarding body weight (g) .

Body weight	Group I (n = 10)	Group IIA (n = 10)	Group IIB (n = 10)	Test of sig.	P
At the beginning of the experiment1	Group I (n = 10)	Group II (n = 20)			
Min. – Max.	130.0 – 150.0	130.0 – 150.0		t= 0.231	0.819
Mean ± SD.	139.7 ± 5.93	139.15 ± 6.23			
Median	140.5	139.0			
At 8 weeks					
Min. – Max.	138.0 – 155.0	202.0 – 339.0	200.0 – 230.0	F= 40.869*	<0.001*
Mean ± SD.	147.5 ± 5.87	268.6 ^a ± 50.66	218.2 ^{ab} ± 10.76		
Median	147.0	270.0	220.5		
Sig. bet. grps	p ₁ <0.001*, p ₂ <0.001*, p ₃ =0.002*				
At the end of the experiment					
Min. – Max.	165.0 – 238.0	177.0 – 299.0	171.0 – 218.0	F= 10.485*	<0.001*
Mean ± SD.	189.8 ± 20.95	246.5 ^a ± 46.32	193.4 ^b ± 17.29		
Median	186.5	244.5	191.0		
Sig. bet. grps	p ₁ =0.001*, p ₂ =0.964, p ₃ =0.002*				

t: Student t-test

F: F for ANOVA test, Pairwise comparison bet. each 2 groups was done using Post Hoc Test (Tukey)

p: p value for comparing between the three studied groups

p₁: p value for comparing between group I and group IIA

p₂: p value for comparing between group I and group IIB

p₃: p value for comparing between group IIA and group IIB

*: Statistically significant at $p \leq 0.05$

Group I: Control

Group IIA: Vitamin D treated

Group IIB: Untreated

a: significant with Group I

b: significant with Group IIA

Table (II): Comparison between the three studied groups regarding fasting blood glucose levels (mg/dl).

Fasting blood glucose(mg/dl)	Group I (n = 10)	Group IIA (n = 10)	Group IIB (n = 10)	F	P
At 9 weeks					
Min. – Max.	68.0 – 95.0	311.0 – 365.0	310.0 – 370.0		
Mean ± SD.	77.80 ± 9.27	336.80 ^a ± 22.49	333.3 ^a ± 20.32	658.84*	<0.001*
Median	74.50	341.50	328.0		
Sig. bet. grps	p ₁ <0.001*, p ₂ <0.001*, p ₃ =0.904				
At the end of the study					
Min. – Max.	63.0 – 97.0	63.0 – 100.0	130.0 – 185.0		
Mean ± SD.	74.80±10.14	86.90±12.80	151.1 ^{ab} ±17.93	85.774*	<0.001*
Median	72.0	93.50	148.5		
Sig. bet. grps	p ₁ =0.149, p ₂ <0.001*, p ₃ <0.001*				
p₄	0.558	<0.001*	<0.001*		

F: F for ANOVA test, Pairwise comparison bet. each 2 groups was done using Post Hoc Test (Tukey)

p: p value for comparing between the three studied groups

p₁: p value for comparing between group 1 and group 2A

p₂: p value for comparing between group 1 and group 2B

p₃: p value for comparing between group 2A and group 2B

p₄: p value for comparing between the same groups at the 9 weeks and at the end of the experiment

*: Statistically significant at p ≤ 0.05

Group 1: Control

Group 2A: Vitamin D treated

Group 2B: Untreated

a: significant groups 2A , 2B versus group1

b: significant with Group 2A

Table (III): Comparison between the three studied groups regarding Serum Insulin and HOMA-IR

Sugar picture	Group I (n = 10)	Group IIA (n = 10)	Group IIB (n = 10)	F	p
Serum Insulin (Pg/ml)					
Min. – Max.	1857.0 – 2354.0	2203.0 – 2927.0	2315.0 – 3560.0		
Mean ± SD.	2053.6 ± 181.8	2551.3 ^a ± 220.1	2900.0 ^{ab} ± 440.8	19.658*	<0.001*
Median	1973.0	2557.0	3003.0		
Sig. bet. grps	p ₁ =0.003*, p ₂ <0.001*, p ₃ =0.041*				
HOMA-IR					
Min. – Max.	7.98–11.25	8.51–17.04	19.51–33.69		
Mean ± SD.	9.37±1.09	13.66 ^a ±2.61	26.76 ^{ab} ±4.38	90.431*	<0.001*
Median	9.03	14.09	27.64		
Sig. bet. grps	p ₁ =0.010*, p ₂ <0.001*, p ₃ <0.001*				

F: F for ANOVA test, Pairwise comparison bet. each 2 groups was done using Post Hoc Test (Tukey)

p: p value for comparing between the three studied groups

p₁: p value for comparing between group I and group IIA

p₂: p value for comparing between group I and group IIB

p₃: p value for comparing between group IIA and group IIB

*: Statistically significant at p ≤ 0.05

Group I: Control

Group IIA: Vitamin D treated

Group IIB: Untreated

a: significant with Group I

b: significant with Group IIA

Table (IV): Comparison between the three studied groups regarding serum ALT (U/ml), serum AST(U/ml) .

Investigation laboratory	Group I (n = 10)	Group IIA (n = 10)	Group IIB (n = 10)	F	p
Serum ALT (U/ml)					
Min. – Max.	15.0 – 37.0	11.0 – 28.0	22.0 – 28.0		
Mean ± SD.	21.0 ± 8.64	17.60 ± 6.13	25.0 ^b ± 2.16	3.520*	0.044*
Median	19.0	16.0	24.50		
Sig. bet. grps	p ₁ =0.453, p ₂ =0.339, p ₃ =0.034*				
Serum AST (U/ml)					
Min. – Max.	32.0 – 70.0	32.0 – 49.0	43.0 – 98.0		
Mean ± SD.	45.20 ± 14.32	41.10 ± 7.28	61.70 ^b ± 20.33	5.315*	0.011*
Median	43.0	43.0	56.0		
Sig. bet. grps	p ₁ =0.814, p ₂ =0.051, p ₃ =0.013**				

F: F for ANOVA test, Pairwise comparison bet. each 2 groups was done using Post Hoc Test (Tukey)

p: p value for comparing between the three studied groups

p₁: p value for comparing between group I and group IIA

p₂: p value for comparing between group I and group IIB

p₃: p value for comparing between group IIA and group IIB

*: Statistically significant at p ≤ 0.05

Group I: Control

Group IIA: Vitamin D treated

Group IIB: Untreated

a: significant with Group I

b: significant with Group IIA

Table (V): Comparison between the three studied groups regarding Serum SOD (U/ml) and IL-1β Eliza (Pg/ml)

	Group I (n = 10)	Group IIA (n = 10)	Group IIB (n = 10)	F	p
Serum SOD (U/ml)					
Min. – Max.	210.0 – 233.0	152.0 – 273.0	142.0 – 203.0		
Mean ± SD.	222.4 ± 8.40	219.5 ± 38.13	178.4 ^{ab} ± 23.51	8.744*	0.001*
Median	223.0	226.5	177.0		
Sig. bet. grps	p ₁ =0.967, p ₂ =0.002*, p ₃ =0.005*				
IL-1β Eliza (Pg/ml)					
Min. – Max.	513.7 – 618.4	410.5 – 844.7	571.8 – 923.4		
Mean ± SD.	563.5 ± 41.27	678.9 ± 128.2	761.7 ^a ± 121.4	9.041*	0.001*
Median	560.0	716.9	804.2		
Sig. bet. grps	p ₁ =0.052, p ₂ =0.001*, p ₃ =0.199				

F: F for ANOVA test, Pairwise comparison bet. each 2 groups was done using Post Hoc Test (Tukey)

p: p value for comparing between the three studied groups

p₁: p value for comparing between group I and group IIA

p₂: p value for comparing between group I and group IIB

p₃: p value for comparing between group IIA and group IIB

*: Statistically significant at p ≤ 0.05

Group I: Control

Group IIA: Vitamin D treated

Group IIB: Untreated

a: significant with Group I

b: significant with Group IIA

Table (VI): Comparison between the three studied groups regarding ΔΔCT (unit ?)

	Group I (n = 10)	Group IIA (n = 10)	Group IIB (n = 10)	F	P
ΔΔCT(unit ?)					
Min. – Max.	0.02 – 0.10	0.04 – 1.05	1.0 – 1.0		
Mean ± SD.	0.07 ± 0.03	0.25 ± 0.31	1.0 ^{ab} ± 0.0	76.407*	<0.001*
Median	0.09	0.12	1.0		
Sig. bet. grps	p ₁ =0.091, p ₂ <0.001*, p ₃ <0.001*				

F: F for ANOVA test, Pairwise comparison bet. each 2 groups was done using Post Hoc Test (Tukey)

p: p value for comparing between the three studied groups

p₁: p value for comparing between group I and group IIA

p₂: p value for comparing between group 1 and group IIB

p₃: p value for comparing between group 2A and group IIB

*: Statistically significant at p ≤ 0.05

Group I: Control

Group IIA: Vitamin D treated

Group IIB: Untreated

a: significant with Group I

b: significant with Group IIA

Effects of vitamin D on liver histopathology

In the control group, the structure of liver is normal and clear. Various pathological changes appeared in the untreated group, including narrow liver sinusoid, distortion of liver architecture, hepatocyte swelling, fatty degeneration, cytoplasm rarefaction, spotty necrosis scattering in the hepatic lobule as well as inflammatory cell infiltration. However, these changes described above were markedly improved in the VD treated group

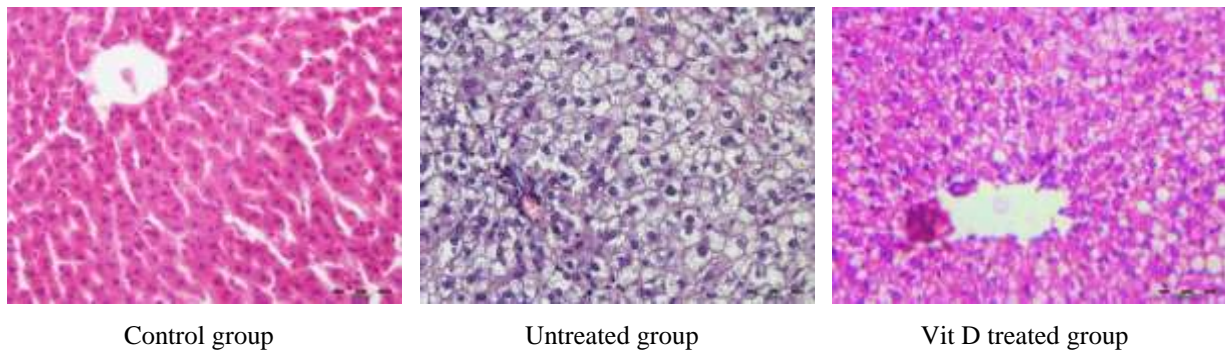


Figure (1): Changes in liver histology (magnification, $\times 400$; stain, hematoxylin and eosin). In the control group, liver cell alignment is normal. In the untreated group, liver cells appear swelled with fatty degeneration and inflammatory cell infiltration. In the vit D treated group, fatty degeneration and inflammatory lesions are alleviated, in comparison with untreated group.

Discussion

It has been demonstrated that inflammation and IR play an initial role in the onset of T2DM and the progression of its complications. Vitamin D is a well-known steroid hormone that has been identified as a regulator of inflammation and IR. Based on available clinical and epidemiological data, the positive effects of vitamin D appear to be primarily related to its action on inflammation and secondary to its action on insulin sensitivity and secretion.

In our study, at the end of the experiment there was significant decrease in body weight in Vit D treated group and untreated group compared by their weight at the 8 weeks. But the BWt was significantly increased in Vit D treated group in relation to untreated group. Most probably the weight loss in untreated group occurred as a result of increased muscle wasting owing to loss of tissue proteins instead of using glucose in aerobic glycolysis. The improvement in BWt in Vit D

treated group is suggested to be due to its effect on insulin sensitivity and glucose metabolism.^(29, 30) Vitamin D influences β -cell insulin secretion through a rise in intracellular calcium concentration via non-selective voltage-dependent calcium channels.^(31, 32) Similar results were found by Rossmeisl et al., [2003](#).⁽³³⁾

Moreover, at the end of the experiment, there was a statistically significant decrease in fasting blood glucose level in group IIB (untreated) and group IIA (Vit D treated group) in relation to control, yet still significantly higher in group IIB than group IIA. Blood glucose return to normal in group IIA may be a result of Vit D improved insulin resistance and improved insulin signaling pathway,⁽³⁴⁾ Benetti et al., 2018⁽³⁵⁾ demonstrated that animals treated with vitamin D showed a significant improvement of glucose tolerance.

In the present study, at the end of the experiment, serum insulin in the untreated group was significantly increased in comparison to control and Vit D treated group. This decrease in serum insulin level in Vit D treated group, supports the evidence that the beneficial effects of vitamin D were mediated through the modulation of VDR a receptor of vitamin D thereby improving signal transduction in the treatment of diabetes-induced liver complications.^(36, 37) **Benetti**, et al⁽³⁵⁾ demonstrated similar results .

In the present work, HOMA-IR in un treated group was significantly increased in relation to control and Vit D treated group. It has been well documented that HFD causes insulin resistance and also reductions in GLUT4 protein levels which are usually associated with increased insulin resistance.^(36, 38) Also significant decreased of HOMA-IR in Vit D treated group suggested that Vit D improves insulin resistance; which was also found by Zhuo et al⁽³⁹⁾

In the present study, serum ALT was significantly increased in the untreated group compared with Vit D treated group. Serum AST showed the same pattern, due to the role of 1, 25 (OH) ₂D₃ in regulating pro inflammatory cytokines and lipid metabolism in a diabetic liver model. ⁽⁴⁰⁾
⁽⁴¹⁾⁽⁴¹⁾⁽⁴¹⁾Ning et al⁽⁴²⁾ demonstrated similar results

In present work serum SOD between significantly increased in the Vit D treated group and significantly decreased in untreated group, suggesting that vitamin D played a vital role in the protection of tissues from damage by free radicals as a non-enzymatic antioxidant compound. ^(43, 44)
Claro da Silva Tet al work is supporting our results. ⁽⁴⁵⁾

In the present study, there was a significantly increased Serum interleukin 1β in untreated group compared to Vit D treated one and in the control group, supporting the role of 1,25-dihydroxy vitamin D as an anti-inflammatory agent .1,25-

(OH)2D3 inhibits the release of the pro-inflammatory cytokine ^(46,47) , Similar result shown by Liu et al⁽²⁰⁾ .

In this study, RT-qPCR analysis showed that the expression level of JNK which are implicated in inflammation and insulin resistance, increased in untreated and decreased following treatment with vitamin D. Based on these results, we speculate that vitamin D exerts therapeutic effects on diabetes-induced liver complications, possibly by downregulating the expression of JNK. This modulation may be in part due to the anti-inflammatory effect of vitamin D; however. the underlying mechanism remains to be elucidated⁽²⁰⁾. Our findings are supported by Liu et al. ⁽²⁰⁾

The histopathological study of liver was performed to assess the direct effects of vitamin D on diabetes-induced liver complications. The results showed that the administration of vitamin D significantly alleviated certain pathological changes, particularly steatosis and inflammatory cell infiltration.

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Prevalence of Serum Cryoglobulins in Patients with Proliferative Lupus Nephritis and its Relation to Serological Markers of Disease Activity

Hesham El Ghoneimy¹, Hanaa Donia², Jasnim Kamel¹, Amr Ebeid¹

(1) Departments of Internal Medicine Faculty of Medicine, Alexandria University; Alexandria Egypt.

(2) Clinical & Chemical Pathology Faculty of Medicine, Alexandria University; Alexandria Egypt.

Abstract

Background: Cryoglobulins are found in patients with systemic lupus erythematosus (SLE) with a variable percentage but their relation to disease activity in patients with proliferative lupus nephritis is not determined yet. **Objectives:** To determine the prevalence of cryoglobulinemia in patients with proliferative lupus nephritis and its relation to disease activity. **Methods:** In a comparative cross-sectional study we investigated 60 candidates, 30 patients with renal biopsy proven proliferative Lupus nephritis (LN), 15 patients with SLE without nephritis and 15 healthy subjects. Serum samples were obtained at 37°C, and cryoglobulinemia was estimated by centrifugation at 4°C after incubation for 7 days in all the studied groups. We also compared between cryo-positive patients and cryo-negative patients according to serological markers of disease activity and activity and chronicity indices in renal biopsy. **Results:** Cryoglobulins were detected in the sera of 2 patients in the LN group (6.7%) , 1 patient in the SLE without nephritis group (6.7%) and non of the

healthy subjects. Rheumatoid factor (RF) titre was significantly higher in cryo-positive patients (p=0.006). Complement levels; C3 and C4 were significantly lower in cryo-positive patients (p=0.006 and 0.011 respectively). There was no statistically significant difference between cryo-positive and cryo-negative patients as regard Anti-double stranded (ds DNA) titre, activity or chronicity indices in renal biopsy. **Conclusion:** Cryoglobulins were positive in 6.7% of cases in our study with no difference between patients with or without nephritis and their presence were not related to pathological markers of disease activity in patients with proliferative LN.

Corresponding Author: Hesham Abdallah El Ghoneimy. Department of Internal Medicine, Nephrology Unit, Alexandria University Faculty of Medicine

Hesham_elghoneimy@yahoo.com

Key Words: Cryoglobulinemia, Lupus nephritis, Lupus activity.

INTRODUCTION

Among patients with lupus, LN affects both sexes equally, is more severe in children and men, and is less so in older adults. The incidence of LN is about 30% in White, 60% in Black and Hispanic, and 40% to 80% in Asian patients with SLE.

Nephritis is more common in African Americans and Hispanics than in whites, and is also more in men than in women. Renal damage is more likely to develop in nonwhite groups.⁽¹⁻⁴⁾

Patients with International Society of Nephrology (ISN) class I biopsies usually have no evidence of clinical renal disease. Similarly, patients with ISN class II may have elevated anti-dsDNA or low complement levels, but usually they do not have active urinary sediment, hypertension is uncommon, the GFR is normal, and proteinuria is rarely above 1 gram/24 hours. Patients with class I and class II LN in renal biopsy have a good renal prognosis unless they transform to another class.⁽⁵⁾

Patients with active ISN class IIIA or IIIA/C usually have microscopic hematuria, hypertension, low complement levels, and proteinuria.⁽⁵⁾ Patients with mild proliferation affecting a few glomeruli usually have good response to treatment, with less than 5% progressing to renal failure during 5 years of follow-up. Others with more glomerular affection or with necrotizing lesions and crescent formation have a prognosis similar to that of class IVA patients.⁽⁶⁾

Patients with ISN class IVA usually have high serologic activity (low serum complement and high anti-dsDNA-binding activity). Clinically, they have active urinary sediment, hypertension, heavy proteinuria, and reduced GFR. Class IV diffuse proliferative disease has the worst renal prognosis in all studies.⁽⁷⁾

Cryoglobulinemia is a clinical disorder characterized by the presence of cryoglobulins in the serum. Cryoglobulins are immunoglobulins that have a specific physical feature of being able to precipitate at cold temperatures and re-dissolve when rewarmed. Cryoglobulinemia may be asymptomatic without end-organ damage and is sometimes accidentally discovered. In type I cryoglobulinemia, the cryoglobulins are monoclonal immunoglobulins (Ig's), usually of the IgG or IgM isotypes and rarely IgA or free immunoglobulin light chains.⁽⁸⁾ Type I cryoglobulinemia develops in the case of protein-secreting monoclonal gammopathies.^(9,10)

In type II cryoglobulinemia, the cryoglobulins are a mix of monoclonal IgM with rheumatoid factor (RF) activity and polyclonal IgG. This type is usually associated with hepatitis C virus (HCV) infection in up to 90% of patients.⁽¹¹⁾ Other causes of type II cryoglobulinemia include other infections (mainly HIV and hepatitis B virus [HBV]), connective tissue diseases (CTDs) like SLE, and lymphoproliferative disorders. Approximately 10% of patients have unknown cause (termed essential mixed cryoglobulinemia). Type III cryoglobulins are by polyclonal IgM with RF activity and polyclonal IgG. This type is seen in CTDs or secondary to infection (mainly HCV). Generally, cryoglobulins in mixed

cryoglobulinemia result from a B-cell proliferation in the setting of chronic immune activation induced by chronic infection, autoimmune disease, or an unknown cause.⁽¹²⁾ However, its role in the pathogenesis and activity of lupus nephritis hasn't been established yet.

PATIENTS & METHODS

We conducted a comparative cross sectional study in which patients were divided into three groups: Group (I): Thirty patients with biopsy proven proliferative lupus nephritis, group (II): Fifteen patients with SLE without lupus nephritis and group (III): Fifteen age and sex matched healthy subjects. We excluded patients with positive hepatitis c virus (HCV) antibodies and antiphospholipid antibodies.

All patients in the study were subjected to thorough history taking with emphasis on symptoms suggestive of cryoglobulinemia as hyperviscosity symptoms (like headache, blurring of vision), skin rash, neuropathy, (as burning sensation and numbness), urinary symptoms (as red or frothy urine) or previously diagnosed thrombotic episodes. They were also well examined for signs of cryoglobulinemia like vasculitic rash and neuropathy.

Ultrasound guided percutaneous renal biopsy with light microscopic examination of obtained specimen and estimation of activity and chronicity index was done.

Complete blood count (CBC), Serum urea & creatinine, complete urine analysis, Urinary protein/creatinine ratio, Anti-nuclear antibody (ANA), Anti dsDNA, Complement factors C3 & C4, Erythrocyte sedimentation rate (ESR) & C-reactive protein (CRP), Rheumatoid factor (RF) and serum cryoglobulins were done.

Blood samples were obtained and kept at 37°C for 30 minutes before separation. Serum was prepared by centrifuging at 37°C for 10 minutes at 2,500 rpm. Fresh centrifugated serum was incubated at 4°C for 7 days after collection and examined for cryoprecipitation.

Statistical Analysis:

We used conventional chi-square and Fisher's exact tests to analyze qualitative differences. For comparison of quantitative parameters, Student t test was used for normally distributed quantitative variables to compare between two studied groups, and the nonparametric Mann-Whitney U test was used for abnormally distributed quantitative variables to compare between two studied groups. F-test (ANOVA) was used for normally distributed

quantitative variables to compare between more than two groups and Kruskal Wallis test for abnormally distributed quantitative variables to compare between more than two studied groups. Post Hoc (Dunn's multiple comparisons test) was used for pairwise comparisons.

A value of P 0.05 indicated statistical significance. This statistical analysis was performed by the SPSS program (SPSS Inc, Chicago, IL) with the information stored in the database program.

RESULTS:

Cryoglobulins were detected in the sera of 3 patients (6.7%), of whom 1 was a female with SLE

without nephritis and 2 were females with proliferative LN.

The duration of SLE ranged between 0.0 – 9.0 years with a median of 2.0 years in group I and 0.20 – 17.0 years with a median of 2.0 years in group II with no statistically significant difference (p=0.608). At the study initiation, the mean age of patients was 31.93 ± 8.68 in group I, 32.60 ± 10.08 in group II and 30.60 ± 8.25 in group III with no statistically significant difference between the three groups (p=0.821).

Table I shows the affected body systems by SLE in the three studied groups.

Table I: Comparison between the three studied groups according to the affected systems

Systems affected	Group I (n = 30)		Group II (n = 15)		Group III (n = 15)		χ ²	P
	No.	%	No.	%	No.	%		
Kidney	30	100.0	0	0.0	0	0.0	–	–
Skin	7	23.3	6	40.0	0	0.0	1.352	^{FE} p=0.304
CNS	2	6.7	2	13.3	0	0.0	0.549	^{FE} p=0.591
Blood	6	20.0	3	20.0	0	0.0	0.00	^{FE} p=1.000
Heart	0	0.0	2	13.3	0	0.0	4.186	^{FE} p=0.106
Joints	10	33.3	8	53.3	0	0.0	1.667	0.197
Serosa	2	6.7	2	13.3	0	0.0	0.549	^{FE} p=0.591
Lung	1	3.3	0	0.0	0	0.0	–	–

χ²: Chi square test FE: Fisher Exact
 p: p value for comparing between the studied groups
Group I: LN
Group II: SLE without nephritis

As shown in table II on comparing between cryo-positive and cryo-negative patients according to serological and pathological markers of the disease activity, we found that C3 had a median of 70.5 mg/dl in cryo negative patients and 26.0 mg/dl in cryo positive patients (p=0.006). C4 had a median of 10.95 mg/dl in cryo negative patients and 4.0 mg/dl

in cryo positive patients (p=0.011). Anti-Ds DNA titre had a median of 134.5 IU/mL in cryo negative patients and 130.0 IU/mL in cryo positive patients (p=0.909). RF titre had a median of 8.0 IU/mL in cryo negative patients and 16.50 IU/mL in cryo positive patients (p=0.006).

Table II: relation between cryoglobulinemia and C3, C4, RF and anti-ds DNA titre

	CRYO		U	P
	Negative (n=42)	Positive (n=3)		
C3				
Min. – Max.	12.0 – 176.0	25.0 – 27.0		
Mean ± SD.	71.05 ± 31.81	26.0 ± 1.0	3.0*	0.006*
Median	70.50	26.0		
C4				
Min. – Max.	3.0 – 47.0	2.10 – 5.0		
Mean ± SD.	14.13 ± 10.60	3.70 ± 1.47	7.0*	0.011*
Median	10.95	4.0		
Anti-Ds DNA				
Min. – Max.	15.0 – 1000.0	32.0 – 173.0		
Mean ± SD.	183.26 ± 210.41	111.67 ± 72.27	60.50	0.909
Median	134.50	130.0		
RF				
Min. – Max.	3.90 – 13.30	11.50 – 24.70		
Mean ± SD.	7.66 ± 2.32	17.57 ± 6.66	2.50*	0.006*
Median	8.0	16.50		

U: Mann Whitney test

p: p value for comparing between **Negative** and **Positive**

*: Statistically significant at $p \leq 0.05$

The activity index (AI) ranged between 1.0 – 12.0/24 with a median of 7.0/24 in cryo negative patients and 6.0 – 10.0/24 with a median of 8.0/24 in cryo positive patients with no significant statistical difference between the two groups ($p=0.901$).

The chronicity index (CI) ranged between 0.0 – 10.0/12 with a median of 3.0/12 in cryo negative patients and 0.0 – 0.0/12 with a median of 0.0/12 in cryo-positive patients with no significant statistical difference between the two groups ($p=0.074$) as shown in table III.

Table III: relation between cryoglobulinemia and activity and chronicity indices in the renal bioosy

	CRYO		U	P
	Negative (n=28)	Positive (n=2)		
AI/24 (n=27)				
Min. – Max.	1.0 – 12.0	6.0 – 10.0		
Mean ± SD.	7.57 ± 2.66	8.0 ± 2.83	26.50	0.901
Median	7.0	8.0		
CI/12				
Min. – Max.	0.0 – 10.0	0.0 – 0.0		
Mean ± SD.	3.54 ± 2.89	0.0 ± 0.0	6.0	0.074
Median	3.0	0.0		

U: Mann Whitney test

p: p value for comparing between **Negative** and **Positive**

*: Statistically significant at $p \leq 0.05$

C3 and C4 were significantly lower in cryo positive patients. RF titre was significantly higher in cryo positive patients. But there was no statistically significant difference between cryo positive and cryo negative patients as regard Anti-DsDNA titre, serum creatinine, urinary protein/ creatinine ratio or activity and chronicity indices in the renal biopsy.

DISCUSSION

In this study, we have investigated 60 subjects, 30 patients with proliferative lupus nephritis, 15 patients with SLE without renal affection and 15 healthy control subjects. Females represented 95% of all cases forming 93% of the LN group and 100% of the group without nephritis. This is consistent with the female predominance shown in previous studies.

The mean age of the three study groups was around 30 years. This is consistent with the literature data that SLE occurs more commonly between 15 and 45 years of age. Group I and group II had a comparable duration of the disease with a mean of 2 years in group I and 3 years in group II.

As regards the affected systems, apart from the kidney that was affected in 100% of group I and not affected at all in group II, the most common presentation of SLE was musculoskeletal affection (40% of cases), followed by mucocutaneous affection (28 % of cases), other affected systems included : blood, lungs, CNS, the heart and serosal surfaces.

In this study, we compared between patients of SLE without nephritis, patients with proliferative LN and healthy controls as regards their routine laboratory investigations, serological investigations for SLE activity and cryoglobulinemia. As ruled in the inclusion and exclusion criteria of each group in the study, all patients in group II had normal renal function tests, normal urinary protein creatinine ratio and normal urine analysis findings.

In group I, they all had renal biopsy proven proliferative LN. The most commonly found pathology is the diffuse proliferative form with or without superadded membranous affection accounting for 63% of cases.

As regards cryoglobulinemia, cryoglobulins were found in 6.7% of SLE patients either with or without LN. There was no relation between the presence of cryoglobulins and the occurrence of LN. Also there was no relation between the presence of cryoglobulins and the activity or chronicity indices in renal biopsy.

Previous studies done to investigate the prevalence of cryoglobulinemia in SLE showed different results ranging from 16% to 83% in small series of cases. A study done in Barcelona in 2001 on a large series of 122 SLE patients found that cryoglobulins were positive in 25% of cases.

In this study neither HCV positive nor antiphospholipid positive patients were excluded. They found that a cryocrit greater than 1% was more frequent in those SLE patients with HCV infection. As our study showed they found a higher frequency of some immunologic markers in cryoglobulinemic SLE patients (RF and hypocomplementemia).

The association between hypocomplementemia and cryoglobulins is well known. Adu and Williams ⁽¹³⁾ described the ability of SLE cryoglobulins to fixate complement in vitro and suggested that these immune complexes can fixate complement in vivo and so, they cause tissue damage in this disease.

Roberts et al., ⁽¹⁴⁾ suggested that, in SLE patients with diffuse proliferative glomerulonephritis, cryoglobulins and glomerular immune deposits can fixate complement via the classic and alternative pathways. They found a higher prevalence of RF in cryoglobulinemic SLE patients, may be due to the RF activity of some cryoglobulinemic component. RF can be used as an immunologic marker that points to the presence of cryoglobulinemia in SLE patients.⁽¹⁵⁾

A cross-sectional study in Isfahan, Iran and was conducted from September 2010 to May 2011. They studied 80 women with SLE, they found a prevalence of 48.8% of cryoglobulins in a large number of SLE patients, all of them showed high titre of circulating cryoglobulins (cryocrit >5%). There was a correlation between Anti-dsDNA, ANA, CRP, and hypocomplementemia and cryoglobulinemia in the studied SLE patients. These finding may identify the SLE patients with cryoglobulinemia. That study helped in the recognition of immunological characteristics of SLE which are involved in systemic inflammation.

In 2016 a case-control study was done. All patients with a cryoglobulinemia between January 2005 and December 2016 in a third level referral centre in Mexico City were included. Cryoglobulinemic SLE patients (cryocrit 1%) were included in the case group, whereas non cryoglobulinemic SLE patients were considered controls.

The investigators studied the demographic, clinical and immunological characteristics at the time of the positive cryoglobulin result, as well as three months earlier, and 6 and 12 months later. Thirty-six SLE patients had a positive test for cryoglobulins throughout the study period. Ten patients had cryocrit of 1% and were included in the case group, whereas 26 patients with a negative test were included as controls. Mean age was 37.7 ± 18.3 in cases and 41.7 ± 19.3 in controls. Women represented 70% of cases and 88.5% of controls.

Among the case group, the cryocrit was 1% in 9 patients, and 3% in one. Regarding clinical and immunological features, a positive lupus anticoagulant and vasculitis were more common in cryoglobulinemic patients ($p=0.004$ and 0.04 , respectively). At the time of the cryoglobulin detection, patients in the case group had lower levels of C3 and C4 ($p=0.026$ and $p=0.003$, respectively), and serum albumin ($p=0.028$). They also had a higher frequency of serositis ($p=0.021$), peripheral oedema

($p=0.034$) and SLICC Damage Index score ($p=0.014$) than controls.

On follow-up, the cryoglobulinemic patients had a higher SLEDAI score after six and twelve months ($p=0.009$ and 0.034 , respectively). After 12 months they had a higher frequency of renal activity ($p=0.004$) and lower C4 levels ($p=0.001$). Among patients with renal activity, 20% of cases and 55% of controls had achieved complete remission after 12 months. So according to that study serum cryoglobulins in SLE patients were relates to positive lupus anticoagulant and hypocomplementemia. At follow-up, patients with cryoglobulinemia had a higher frequency of renal activity, as well as an increased disease activity overall.⁽¹⁶⁾

Comparing our study to the previous studies, the prevalence of cryoglobulinemia found in our study was lower with no difference between LN and non LN patients. Cryoglobulinemia was related to disease activity in SLE represented by hypocomplementemia. There was no relation between the presence of cryoglobulins and activity or chronicity indices in the renal biopsies of LN patients.

These differences between our results and the results of the previous studies may be attributed to exclusion of HCV positive patients who represented some percentage of positive cryoglobulins in SLE patients in the previous studies.

The same also applies to exclusion of patients with positive antiphospholipid antibodies in whom the presence of cryoglobulins are more likely. Other probable explanation includes the use of immunosuppressive therapy in most of our studied patients which may have its impact on the occurrence of seronegativity. Lastly, cryoglobulins may be falsely negative in some patients due to the strict precautions needed for sample collection and analysis.

CONCLUSION

Cryoglobulinemia is not related to pathological markers of disease activity in the renal biopsies of patients with proliferative LN.

Conflict of Interest: Authors declare no conflicts of interest.

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