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**STUDY OF SERUM IRON AND FERRITIN  
IN PATIENTS WITH ISCHEMIC HEART  
DISEASES**

٥١٠

THESIS

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**Degree of Master of Chemical Pathology**

By

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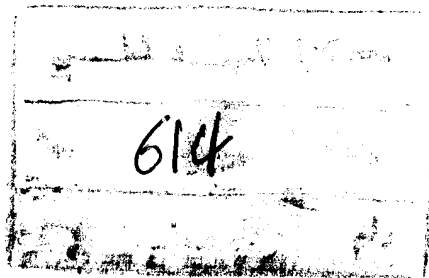
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*Dedicated to*  
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*and*  
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## LIST OF ABBREVIATIONS

AMI	: acute myocardial infarction
Apo A	: Apoprotein A
Apo B	: Apoprotein B
Apo B <sub>100</sub>	: Apoprotein B <sub>100</sub>
Apo C	: Apoprotein C
Apo D	: Apoprotein D
Apo E	: Apoprotein E
HCO <sub>3</sub> <sup>-</sup>	: Bicarbonate
CA	: coronary artery
CHD	: coronary heart disease
CK-MB	: creatine kinase-MB
CU	: copper
DNA	: deoxyribonucleic acid
ECF	: extracellular fluid
ECG	: electrocardiogram
EDTA	: ethylene diamine tetra acetic acid
e <sup>-</sup>	: electron
FBG	: fasting blood glucose
Hb	: haemoglobin
Ht	: haematocrit
HCL	: hydrochloric acid
HRP	: horse radish peroxidase
H	: hydrogen
OH	: hydroxyl radical
H <sub>2</sub> O <sub>2</sub>	: hydrogen peroxide
HDL	: high density lipoprotein
HDL-C	: high density lipoprotein cholesterol
Fe	: iron
IHD	: ischaemic heart disease
LCAT	: lecithin cholesterol acyl transferase
LDL	: low density lipoprotein
LDL-C	: low density lipoprotein cholesterol

<b>LDL-C</b>	: low density lipoprotein cholesterol
<b>MDA</b>	: malondialdehyde
<b>McABs</b>	: monoclonal antibodies
<b>NO</b>	: nitric oxide
<b>NO<sup>+</sup></b>	: nitric oxide radical
<b>O<sub>2</sub></b>	: oxygen
<b>SOD</b>	: superoxide dismutase
<b>TMP</b>	: tetra methyl benzidine
<b>TBA</b>	: thiobarbituric acid
<b>TBARs</b>	: thiobarbituric acid reactive substances
<b>T.Ch.</b>	: total cholesterol
<b>TIBC</b>	: total iron binding capacity
<b>TG</b>	: triglyceride
<b>UV</b>	: ultraviolet
<b>VLDL</b>	: very low density lipoprotein
<b>Zn</b>	: zinc

# INTRODUCTION

# INTRODUCTION

## Ischemic Heart Disease

Ischemic heart disease is defined by the World Health Organization as myocardial impairment due to imbalance between coronary blood flow and myocardial requirements caused by changes in the coronary circulation.<sup>(1)</sup>

The imbalance is usually related to either an absolute reduction in coronary blood flow or inability to increase coronary blood flow relative to the needs of the heart and most often due to atherosclerotic obstruction of large coronary artery.<sup>(2)</sup>

Coronary atherosclerosis is most prominent in the main stem of the coronary arteries, particularly in the segments closest to the ostia of the coronary vessel.<sup>(3)</sup>

### **Risk factors for atherosclerosis:**

Risk factors are certain biological variations or habits of life which can be observed more commonly in coronary patients than in general population.<sup>(4)</sup> They have traditionally been divided into two categories:<sup>(3,5)</sup>

#### ***a- Modifiable risk factors:***

(hyperlipidaemia, hypertension, smoking, diabetes, overweight, physical inactivity and psychological factors).

***b- Non modifiable risk factors:***

(age, gender and hereditary factors).

However more recent data indicate clustering of several metabolic and familial factors that are often related to each other.<sup>(5)</sup>

**1- Hyperlipidemia:**

The association of various lipid parameters with the progression of coronary disease is well established and hyperlipidaemia is a known positive risk factor for coronary arteriosclerosis.<sup>(6)</sup>

**2- Hypertension:**

Coronary disease is now the commonest sequela of hypertension. The incidence of cardiovascular disease and the factors that predispose to it including hyperlipidaemia, hyperuricaemia, glucose intolerance and obesity are more common in hypertensive persons than in the general population.<sup>(7)</sup>

When the blood pressure exceeds 160 mmHg systolic and 95 mmHg diastolic in middle-aged men the risk is five times greater than in normotensive men.<sup>(3)</sup>

**3- Cigarette smoking:**

Smokers are at high risk of developing atherosclerosis and cardiac ischemic disease. Cigarette smoke contains vast amounts of both carbon- and oxygen-centered free radicals, which can directly initiate and propagate the process of lipid peroxidation.<sup>(8)</sup>

Recently, several studies have implicated oxidative modification of LDL by cigarette smoke as a possible explanation.<sup>(9-11)</sup>

#### **4- Glucose intolerance and diabetes mellitus:**

Diabetics show at least twofold increase in the incidence of myocardial infarction, compared with nondiabetics. Younger diabetics have a marked increase in the risk of ischemic heart disease, and diabetic women appear to be even more prone than diabetic men.<sup>(3)</sup>

#### **5- Age:**

“Aging theory of atherosclerosis” supposes that atherosclerosis is an inevitable consequence of ageing and therefore an irreversible process.<sup>(12)</sup>

#### **6- Sex:**

Male gender remains one of the most powerful factors at all levels of risk.<sup>(3)</sup> Epidemiological studies has shown that women loose their protection against CHD after menopause. This cardioprotection in premenopausal women has generally been contributed to endogenous estrogen<sup>(13)</sup> which might influence the development of atherosclerosis by their effect on the distribution or nature of the plasma lipoproteins.<sup>(3)</sup>

#### **7- Genetic factors:**

They are critical in atherosclerosis. The best example of this is the increased incidence of atherosclerosis in individuals with homozygous familial hypercholesterolaemia and familial combined hyperlipidaemia.

Other risk factors such as hypertension and diabetes mellitus can also be inherited and protective factors as increased HDL may also be inherited.<sup>(3)</sup>

#### **8- Obesity:**

When body weight is greater than twenty percent above the normal, there is increased risk of ischaemic heart disease. Obesity is generally associated with hypertriglyceridaemia, hypercholesterolaemia, glucose intolerance and hypertension.<sup>(3)</sup>

#### **9- Physical inactivity:**

Sedentary individuals are more susceptible to atherosclerosis and to sudden death than individuals who maintain an active lifestyle. Increased physical activity may elevate HDL levels.<sup>(3)</sup>

## Pathology

Almost all myocardial infarctions result from coronary atherosclerosis. Generally with superimposed coronary thrombosis.<sup>(14)</sup>

### **Role of acute plaque change:**

Slowly occurring high-grade stenosis of epicardial coronaries may progress to complete occlusion but do not usually precipitate AMI. Probably because of the development of a rich collateral network over time. However, during the natural evolution of atherosclerotic plaques, especially those that are lipid-laden, an abrupt and catastrophic transition may occur characterized by plaque rupture and exposure of substances that promote platelet activation and thrombosis generation. The resultant thrombus interrupts blood flow and leads to an imbalance between oxygen supply and demand and, if this imbalance is severe and persistent, it will lead to myocardial necrosis.<sup>(15-21)</sup>

### **Composition of plaques:**

At autopsy, the atherosclerotic plaque of patients who died of MI is composed primarily of fibrous tissue of varying density and cellularity with superimposed thrombus.<sup>(22-24)</sup>

Calcium, lipid-laden foam cells, and extracellular lipid each constitute 5 to 10 percent of the remaining area.<sup>(22)</sup>

The atherosclerotic plaques that are associated with thrombosis and a total occlusion, located in infarct-related vessels, are generally



more complex and irregular than those in vessels not associated with MI.<sup>(25)</sup>

Histological studies of these lesions often reveal plaque rupture or fissuring.<sup>(16,25,26)</sup>

Angiographic morphology suggestive of plaque rupture has been identified in the majority of stenosis associated with AMI or abrupt onset of unstable angina.<sup>(27)</sup>

Platelet-rich thrombi are often associated with the surface of the most advanced atherosclerotic lesions, called complicated plaque, which are characterized by fibrocalcific degeneration, deposition of lipid, calcium, fibrous tissue, necrotic debris, extravasated blood, and a fibrous cap.<sup>(25)</sup>

Impaired endothelial cell function may contribute to atherogenesis through release of growth factors. Luminal narrowing may potentiate platelet activation through augmentation of shear forces.<sup>(25)</sup>

Young persons with coronary thrombotic events have been described as having a genetic polymorphism in glycoprotein IIb/IIIa, possibly altering platelet-fibrinogen interactions. This observation raises the possibility of screening for patients at increased risk of coronary thrombosis in the event of plaque rupture.<sup>(28)</sup>

**Plaque fissuring and rupture:**

The process of plaque fissuring is an area of intense investigation and is likely to be multifactorial in nature.<sup>(20)</sup>

Lippy has summarized the evidence suggesting that T lymphocytes in human atheroma elaborate the cytokine interferon gamma (IFN- $\gamma$ ) that markedly inhibits the ability of vascular smooth muscle cells to form interstitial collagen in vulnerable regions of the fibrous cap over an atherosclerotic plaque.<sup>(25)</sup>

Furthermore, in atherosclerotic plaques prone to rupture there is an increased rate of formation of metalloproteinase enzymes such as collagenase, gelatinase, and stromelysin that degrade components of the protective interstitial matrix.<sup>(21,25,29)</sup>

These proteinases may be elaborated by activated macrophages and mast cells that have been shown to accumulate in high concentration at the site of atheromatous erosions and plaque rupture in patients who died of AMI.<sup>(19,30,31)</sup>

Reactive oxygen species and free radicals are produced continuously in the body as a result of normal metabolic processes. Free radicals can play a beneficial role in fighting off infection. For example, phagocytes in the blood attack and destroy disease-causing microbes through a mechanism involving free radicals.<sup>(32)</sup> However, free radicals can also damage complex molecules such as proteins<sup>(33)</sup> and lipids<sup>(34)</sup> or even DNA.<sup>(35)</sup>

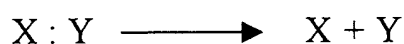
Free radicals have been lately associated with diseases such as cancer, heart disease, rheumatoid arthritis and it is speculated that numerous diseases in both adult and neonatal medicine are associated with free radicals and their damaging effect.<sup>(36)</sup>

**Definition:**

A free radical can be defined as a chemical species possessing one or more unpaired electron(s) in their outer orbital. It can also be considered as a fragment of a molecule.<sup>(37)</sup> The unpaired electron and the radical nature of a species are conventionally indicated by writing it with a heavy superscript dot.

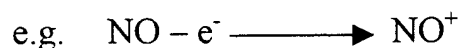
**Mechanism of formation of free radicals:<sup>(38)</sup>**

1- By the homolytic cleavage of a covalent bond of a normal molecule with each fragment retaining one of the paired electrons.

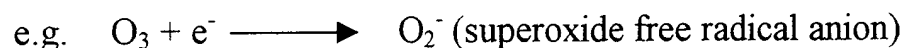


Homolytic fission generally requires high energy input resulting from either high temperature, UV light or ionizing radiation

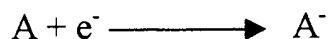
2- By the loss of a single electron from a normal molecule.



3- By the addition of a single electron to a normal molecule.



Electron transfer is a far more common process in biological system than is homolytic fission.

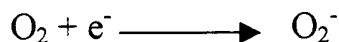


As a consequence of having an unpaired electron in their outer orbital, free radicals have an increased reactivity with other molecules.

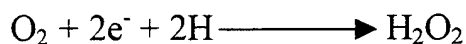
### **Oxygen free radicals and reactive oxygen species:**

One of the most important molecules in free radicals biochemistry is oxygen. Electrons are distributed in oxygen in such a way that two of the electrons are unpaired.

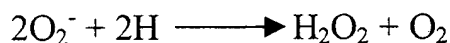
Reduction of oxygen by the transfer to it of a single electron will produce the superoxide free radical anion.



Reduction of oxygen with 2 electrons would yield hydrogen peroxide

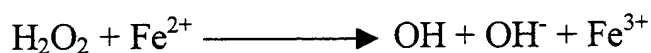


2 superoxide molecules can react together with hydrogen to form hydrogen peroxide and oxygen.



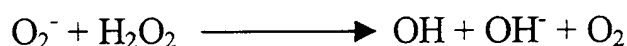
Hydrogen peroxide is not a free radical, but it is an important compound in free radical biochemistry because it can easily breakdown

in the presence of transition metal ions, to produce the most reactive and damaging of the oxygen free radicals, the hydroxyl radical.



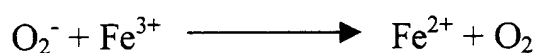
(Fe-catalyzed HaberWeiss reaction or fenton reaction)

The superoxide can react directly with hydrogen peroxide:

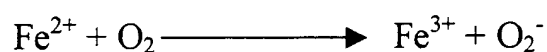


(noncatalysed Haber Weiss reaction)

The iron (or copper) catalysed reaction is considered to be dependant on superoxide.



The auto-oxidation of reduced transition metals can also generate superoxide.



Thus the reactions of the transition metal ions with oxygen can be considered reversible redox reactions and are extremely important in the promotion of the free radical reactions.<sup>(39)</sup>

Thus “the keyplayers” in the biochemistry of oxygen free radicals are oxygen itself, superoxide, hydrogen peroxide, transition metal ions and the hydroxy radical, the first 4 of which conspire by a variety of reactions to generate the last.<sup>(36,38,40)</sup>

**Superoxide:**

It is not a damaging species, its main significance is probably as a source of hydrogen peroxide and as a reductant of transition metal ions.

**Hydrogen peroxide:**

It is a source of hydroxyl radical in the presence of reactive transition metal ions.<sup>(41)</sup>

**The hydroxyl radical:**

It is an extremely reactive oxidising radical that will react with most biomolecules. It has an extremely short half-life, but is capable of causing great damage within a small radius of its site of production.

Oxygen free radicals are not the only important free radicals in biochemistry, although they are often the initial species formed. Carbon centered radicals  $R\cdot$  arise from the attack of an oxidising radical on a biological molecule such as lipid, nucleic acid, carbohydrate and protein.<sup>(37)</sup>

Sulphur atoms can be the centre for free radicals  $RS\cdot$  e.g. in the oxidation of glutathione.<sup>(38)</sup>

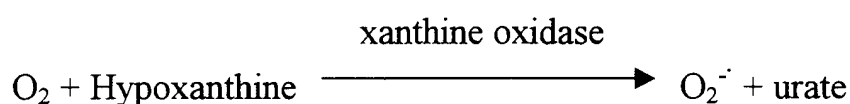
Nitric oxide free radicals NO produced by phagocytes and vascular endothelial cells acts on smooth muscle cells in vessel walls to produce relaxation.<sup>(42,43)</sup>

### **Production of free radicals in cells:**

The major source of free radicals in cells is electron leakage from electron transport chains, such as those in mitochondria and in endoplasmic reticulum, to molecular oxygen generating superoxide.<sup>(42)</sup>

Other enzymes can also produce superoxide or hydrogen peroxide, such as flavin oxidases located in peroxisomes. Another source of superoxide in the cells is the autoxidation of certain compounds including ascorbic acid (vitamin C), thiols (glutathione, cysteine), adrenaline and flavin coenzymes.<sup>(37)</sup>

An important enzymatic oxygen radical generating system is the hypoxanthine-xanthine oxidase system. It is considered as a cytosolic source for free radical formation.<sup>(36)</sup>



Prostaglandin production through the arachidonic acid cascade involves the formation of free radicals.<sup>(44)</sup>

### **Assay for free radicals:**

Thiobarbituric acid reactive substances, (TBARs) assay is the most popular and easiest method used as an indicator of lipid

peroxidation, TBA react with malondialdehyde (MDA), heating is required then after extraction into an organic solvent fluorimetric measurement is applied.<sup>(45)</sup> In uncharacterized systems it is usual to refer to the assay of TBARS (TBA reactive substances) as the test is not specific for MAD. Direct assessment of free MAD is most reliably done by HPLC.<sup>(45)</sup>

### **Low-density lipoprotein and atherogenesis:**

Clinical and experimental studies of various kinds have firmly established that elevated plasma concentrations of low-density lipoprotein (LDL) are associated with accelerated atherogenesis.<sup>(46-48)</sup>

The cholesterol that accumulates in atherosclerotic lesions originates primarily in plasma lipoproteins including LDL.<sup>(49)</sup> Most foam cells arise from circulating monocytes that have taken up residence beneath the vascular endothelium,<sup>(50,51)</sup> although some are derived from medial smooth-muscle cells.<sup>(51,52)</sup> According to response-to-injury theory of atherogenesis, endothelial cell injury is the basic mechanism for initiation and maintenance of atherosclerosis and is followed by atherosclerotic lesions.<sup>(53,54)</sup>

The arterial uptake, giving rise to foam cells and fatty streaks must be by pathways independent of the LDL receptor, as lesions rich in macrophage-derived foam cells develop even in patients and animals deficient in functional LDL receptors.<sup>(51,55,56)</sup>



**Cell-induced oxidation of LDL:**

When LDL was incubated with cultured endothelial cells for 12 to 18 hours, it underwent a striking series of physical and chemical changes and most important, that the resulting endothelial cell-modified form of LDL was taken up by cultured macrophages 3 to 10 times more rapidly than native LDL.<sup>(57,58)</sup>

It has been shown that monocytes and macrophages can themselves effect similar modification in LDL.<sup>(59,60)</sup>

Cell-induced oxidative modification can be mimicked by simply incubating LDL in a serum-free medium in the presence of a sufficiently high concentration of copper or iron.<sup>(61,62)</sup> Whatever the details of the initiating step may be, once the LDL contains fatty acid lipid peroxides, there follows (especially in the presence of metal ions) a rapid propagation that amplifies dramatically the number of free radicals and leads to extensive fragmentation of the fatty acid chains.<sup>(63)</sup>

Some workers suggest that shorter-chain aldehydes (e.g., malondialdehyde and 4-hydroxynonenal) are generated, and some of these can attach covalently to apoprotein B.<sup>(63)</sup>

## **Cytotoxicity of oxidized LDL and its possible role in atherogenesis**

It is shown by some workers that oxidized LDL is highly cytotoxic, but native LDL is not.

Because certain cell types, including endothelial cells, can themselves oxidatively modify LDL, the incubation of native LDL with such cells can lead to cytotoxicity; this can be prevented by the addition of antioxidants.<sup>(64)</sup>

Considerable experimental evidence implicates endothelial injury as an important factor in atherogenesis.<sup>(65)</sup> However, frank denudation of the endothelium does not appear to occur before the development of the fatty-streak lesion. Still, the cytotoxicity of oxidized LDL can conceivably induce functional changes in the endothelial cell that favor the penetration of circulating monocytes or favor the movement of LDL into the subendothelial space and thus accelerate the formation of fatty streak.<sup>(66)</sup>

## **The role of apolipoproteins in the development of atherosclerosis**

### **APO A:**

APO A is the major protein component of HDL. HDL functions as a circulating acceptor of cholesterol from peripheral cells. Under normal conditions there is a net transport of cholesterol from peripheral tissue cells to the site of excretion and metabolism in the liver.<sup>(67)</sup> Most plasma cholesterol is in the esterified form, where as efflux from peripheral cells is in the form of free cholesterol.<sup>(68)</sup>

The major factor catalyzing cholesterol ester synthesis in plasma is lecithin cholesterol acyl transferase (LCAT). LCAT reactivity is dependent upon apolipoprotein A-I and the reaction is inhibited by apolipoprotein A-II.<sup>(69,70)</sup> Efflux of cholesterol is in part catalyzed by a lipoprotein containing apoprotein A-I unassociated with apoprotein A-II. The LCAT is present in plasma in the form of a complex with a minor apoprotein, apoprotein D.<sup>(71)</sup>

### **APO B:**

Apo B-100 forms the main protein mass of LDL. LDL is the principal carrier of cholesterol in plasma and it is the most atherogenic class of cholesterol carrying lipoproteins. The level of plasma LDL is regulated by the LDL receptor, a cell surface glycoprotein, that specifically binds lipoproteins that contain apoprotein B-100.<sup>(72)</sup>

The Apo B containing LDL interacts with the receptors initiating a series of intracellular events including internalization, lysosomal degradation and regulation of cholesterol metabolism.<sup>(73)</sup>

The liver has a wide capacity to Apo B production, which may or may not be associated with hypercholesterolaemia or hypertriglyceridaemia.<sup>(74,75)</sup>

Patients with primary over-production of ApoB appear to be at an increased risk for coronary heart disease.<sup>(76)</sup>

#### **APO C:**

APO C plays an important role in the metabolism of triglyceride rich lipoproteins.<sup>(77)</sup> APO C-II is a specific Co-factor essential for triglyceride hydrolysis by extrahepatic lipoprotein lipase and its deficiency leads to hypertriglyceridemia.<sup>(78)</sup> However, APO C-III acts as specific inhibitor of such an enzyme.<sup>(69)</sup>

#### **APO D:**

The APO D may participate in the activation of the LCAT system and may be a specific carrier of the lysolecithin formed after LCAT has acted on HDL.<sup>(69)</sup>

#### **APO E:**

A subfraction of HDL<sub>2</sub> (unique in containing the E apoprotein) sometimes called HDL with APO E has got a specific antiatherogenic

function. It is produced in response to tissue overload with cholesterol to mediate its removal from tissues.<sup>(73)</sup>

It was found that lipoproteins containing the APO E such as the chylomicron remnants are cleared with much greater efficiency than the lipoproteins containing APO B such as LDL.<sup>(79)</sup>

The APO E seems to play an important role in the binding of lipoproteins to cell surface receptors.<sup>(69,80)</sup>

## Iron Metabolism and transport

Iron participates in numerous metabolic pathways in all cells and organisms. Its ability to exist in  $\text{Fe}^{+3}$  and  $\text{Fe}^{+2}$  states underlines its importance in oxygen and electron transport system concerned with cellular energy production. The liver is intimately involved in many aspects of iron metabolism as a result of the presence of numerous iron dependent metabolic functions.<sup>(81)</sup>

In acidic solutions, iron ions are surrounded by six water molecules, and have the forms of  $\text{Fe}(\text{H}_2\text{O})_6^{+2}$  and  $\text{Fe}(\text{H}_2\text{O})_6^{+3}$ . At physiological pH  $\text{Fe}^{+2}$  is reasonably soluble; whereas the  $\text{Fe}^{+3}$  is almost insoluble.<sup>(82)</sup>

### Iron compounds

The iron content of adult humans is approximately 40 mg/kg in women and 50 mg/kg in men.<sup>(82)</sup> This may be subdivided into:

### Transport iron:

As transferrin which is glycoprotein about 30% saturated with iron and other forms of transport iron as, lactoferrin, haptoglobin and albumin.

Also iron in plasma can be complexed with some chemicals such as citrate, ascorbate and amino acids.

**Storage iron:**

as ferritin and haemosiderin.

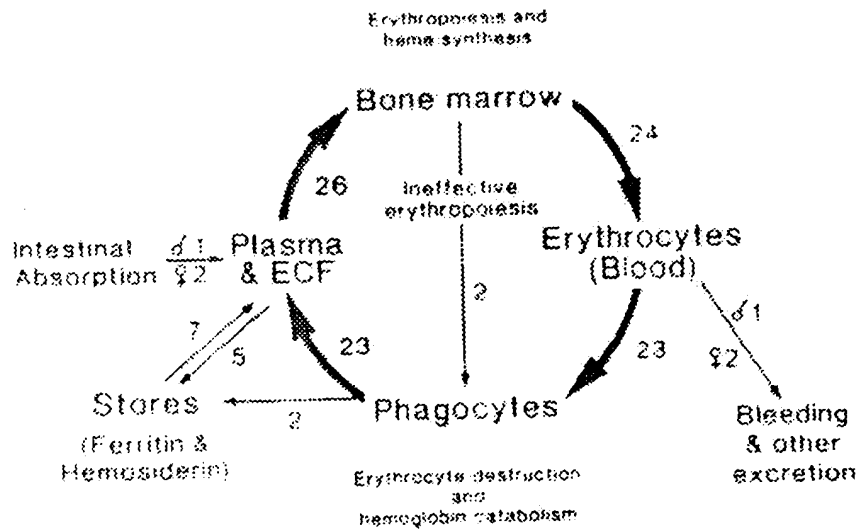
**Functional iron proteins:**

Functional iron containing proteins can be divided into those involved in oxygen transport e.g.: hemoglobin and myoglobin and the heme and nonheme iron containing enzymes as oxidase, peroxidase and catalase.<sup>(83)</sup>

## **Pathways of iron metabolism**

Iron is tightly conserved in a nearly closed system in which each iron atom cycles repeatedly from plasma and extracellular fluid (ECF) to the bone marrow, where it is incorporated into hemoglobin. Then it travels to the peripheral blood, where it remains within erythrocyte in the blood for 4 months. It then travels to phagocytes of the reticuloendothelial system, where senescent erythrocytes are engulfed and destroyed, hemoglobin is digested, and iron is released to plasma, where the cycle continues. With each cycle, a small proportion of iron is transferred to storage sites, where it is incorporated into ferritin or hemosidrin, a small proportion of storage iron is released to plasma, a small proportion is lost in urine, sweat, feces or in blood, and an equivalent small amount of iron is absorbed from the intestinal tract. In addition a small proportion (about 10%) of newly formed erythrocytes normally is destroyed within the bone marrow and its iron released bypassing the circulating blood part of the cycle (ineffective erythropoiesis).





**Pathways of iron metabolism.<sup>(84)</sup>**

The numbers indicate the approximate amount of iron (in mg) that enters and leaves each of these iron compartments every day in healthy adults who do not have bleeding and other blood disorders.<sup>(84)</sup>

#### **Transport iron:**

#### **Transferrin:**

It is a glycoprotein (containing about 6% carbohydrate). It consists of single polypeptide chain of molecular weight 79,570 containing 679 amino acids, organised into two domains of approximately equal size.

Each domain contains one binding site for iron in ferric state.

(C terminal domain and N terminal domain).<sup>(85)</sup>

The iron in the transferrin is in the ferric state. Each molecule of transferrin can bind two ferric atoms with simultaneous binding of one  $\text{HCO}_3^-$  for each ferric atom bound.<sup>(86)</sup>

The stability of the iron-transferrin complex is dependant on the oxidation state of the iron, the presence of the anion, and the confirmation of protein as well as the pH of the solution. Therefore iron can be released from transferrin by reduction of iron, removal of the anion, deformation of the protein (when interacts with receptor) or acidification of the solution.<sup>(87)</sup>

#### **Iron uptake by the cells:**

##### **Transferrin receptor:**

The function of transferrin receptors is to mediate the cellular uptake of iron bound transferrin. In most types of cells, this is achieved by receptor-mediated endocytosis; but in some cells, such as hepatocytes iron possibly can be released from transferrin while it is on the plasma membrane of the cell.<sup>(88)</sup>

##### **Receptor mediated endocytosis of transferrin:**

After binding to its receptor at the cell surface, the transferrin receptor complex is endocytosed into the cell. This is followed by acidification to pH 5.5-6.5. The iron is then released from transferrin and transported to the cytosol.

The apotransferrin remains bound to the receptor and recycled to the cell membrane by exocytosis. At the extracellular pH,

apotransferrin, has poor affinity to its receptor. Apotransferrin is thus displaced from the receptor by iron containing transferrin and the cycle is repeated.<sup>(88)</sup>

Once iron has passed through the plasma and then the endosomal membrane, it is transported to intra cellular sites, to be incorporated into heme, ferritin and other non heme iron protein.<sup>(82)</sup>

#### **Other forms of transport iron:**

Several proteins, that bind iron or iron containing compounds are present in the plasma and play subsidiary roles in the iron transport especially in abnormal conditions such as; iron overload, hemolytic states and inflammatory conditions. Those proteins include, ferritin, lactoferrin, hemopexin, haptoglobin and albumin.<sup>(89)</sup>

#### **Storage iron:**

##### **Ferritin:**

Ferritin is water soluble, iron binding protein with wide distribution in nature. The ferritin molecule is approximately spherical. It consists of an outer protein coat and inner iron rich electron-dense core. The core of the ferritin molecule can contain 4000 iron atoms in the form of ferric hydroxyphosphate.<sup>(90)</sup>

Iron is believed to enter the ferritin core in the ferrous state through hydrophilic channels, it is then oxidised into ferric ions and stored into the molecule. When cells are in the state of positive iron

balance, synthesis of ferritin is stimulated and the iron is deposited as ferritin in a non toxic storage form.<sup>(91)</sup>

When the cell goes into negative iron balance, iron is released from ferritin to enter the metabolic pathways or leave the cell for use elsewhere in the organism.<sup>(92,93)</sup>

### **Hemosidrin:**

Cellular storage iron is also present in water insoluble form called hemosidrin. It consists of iron oxyhydroxy cores and the protein of hemosidrin reacts with antiferritin antibodies but the iron: protein ratio is higher in hemosiderin than in ferritin.<sup>(94)</sup>

Hemosidrin is probably formed by the partial digestion of ferritin by lysosomal enzymes. In normal subjects, storage iron is two thirds in the form of ferritin and one third hemosidrin, but in iron overload the hemosidrin increases considerably.<sup>(94)</sup>

### **Oxidative stress:**

It has been defined as a disturbance in the equilibrium status of pro-oxidant/antioxidant systems in intact cells.

This definition of oxidative stress implies that cells have intact pro-oxidant/antioxidant systems that function continuously to generate and to detoxify oxidants during normal aerobic metabolism. When additional oxidative events occur, the pro-oxidant systems may outbalance the antioxidant, resulting in oxidative damage to lipids,

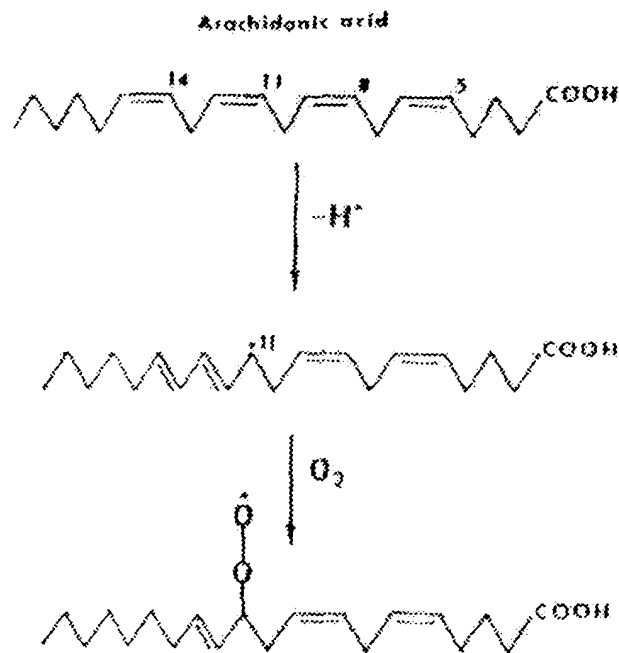
proteins, carbohydrates, and nucleic acids. Oxidative stress may also induce a rapid alteration in the antioxidant systems by depleting cellular stores of antioxidant materials such as glutathione and vitamin E. A disturbance in pro-oxidant/antioxidant systems results from a myriad of different oxidative challenges, including radiation, xenobiotic metabolism of environmental pollutants and other materials, and challenges to the immune system in human disease.<sup>(95)</sup>

### **Effect of oxidants on macromolecules:**

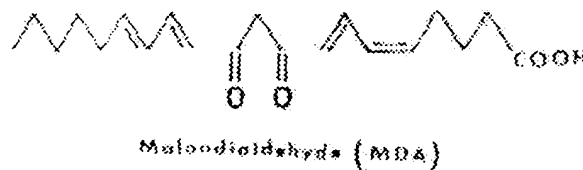
#### **Lipids:**

Peroxidation of cellular lipid results in a variety of deleterious effects on important membrane functions. Most peroxidized lipid occurs as a result of oxidative stress in intact cells, but some peroxidized lipid in the diet may be directly incorporated into cell structures.<sup>(96)</sup>

The reaction of this process shown in figure below.



#### PRODUCTS OF ARACHIDONIC ACID PEROXIDATION



#### Reactions of lipid peroxidation.<sup>(96)</sup>

The products of lipid peroxidation are easily detected in blood plasma and have been used as a measure of oxidative stress. The most commonly measured product is malondialdehyde.<sup>(96)</sup>

In addition the unsaturated aldehydes produced from these reactions have been implicated in the modification of cellular proteins and other

materials. Vitamin E is particularly effective as an antioxidant in lipid peroxidizing systems.<sup>(97)</sup>

### **Cellular antioxidants:**

The most effective antioxidant in oxidative stress depends on the specific molecules causing the stress (e.g., superoxide anion, lipid peroxides, iron generated hydroxyl radical) and the cellular or extracellular location of the source of these molecule. As an example, damage to a cell membrane occurs from both internally and externally generated oxidative stress. This damage is most effectively prevented by vitamin E, which reacts with peroxy and hydroxyl radicals; by carotenoids, which react with singlet oxygen; and possibly by membrane bound proteins. The chain-breaking anti-oxidant function of vitamin E in membranes results from its close association with polyunsaturated components of the membrane.<sup>(98)</sup>

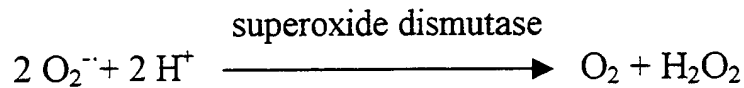
Vitamin E radical can be reduced by cytoplasmic vitamin C and glutathione, or by membrane-bound quinols.

Similarly, when oxidative stress occurs in plasma, a variety of different antioxidants participate in the response. Many plasma proteins are affected by the process, causing either irreversible or reversible loss of functional protein activity.<sup>(99)</sup>

### **Enzymes:**

Superoxide dismutase is one of the most important enzymes that function as cellular antioxidants. It is present in cell cytoplasm (copper-

zinc enzyme) and in mitochondria (manganese enzyme) in order to maintain a low concentration of superoxide anion.<sup>(100)</sup>

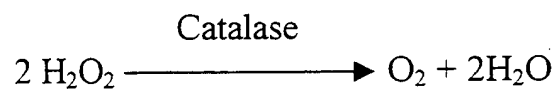


The absence of this enzyme is lethal, but increasing its concentration in cells may not increase antioxidant protection. On the other hand, it may produce hydrogen peroxide at a rate that makes it more toxic to cells in oxidative stress.

Superoxide dismutase is increased by specific redox-sensitive genes in cells under continued stress.<sup>(101)</sup>

An extracellular form of superoxide dismutase, different from the intracellular forms of the enzymes, occurs in plasma, lymph and synovial fluid.<sup>(102)</sup>

The extracellular enzymes may function at cell surfaces. Catalase is a heme protein. It is usually found in peroxisomes. In cells such as erythrocytes, which don't contain these organelles, it is a cytoplasmic enzyme. Catalase provides a protective role that is similar to that of glutathione peroxidase in most cells.<sup>(103)</sup>





Superoxide dismutase and catalase provide a rapid means of equilibrating and detoxifying superoxide anion and hydrogen peroxide in cells.

In addition, both enzymes have considerable use as pharmacologic agents to decrease the effect of oxygen radicals in human disease.

There has been interest in these enzymes for prevention of reperfusion injury.<sup>(104)</sup>

### **Plasma antioxidants:**

Human plasma contains little catalase, superoxide dismutase, glutathione, or glutathione peroxidase; the antioxidant properties of this important fluid reside primarily in a number of small molecules and protein constituents.

In studies using both artificially generated oxidants and natural oxidants such as cigarette smoke, ascorbate is among the first compounds that become oxidized in stress.<sup>(105,106)</sup>

The antioxidant proteins of plasma that are most important include ceruloplasmin, albumin, transferrin, haptoglobin and hemopexin. The first three proteins may sequester iron and copper ions in forms that prevent their participation in reactions that generate the aggressive hydroxyl radical. Haptoglobin and hemopexin bind free heme, a source of iron that can participate in lipid oxidation reactions

(oxidative stress also affects plasma lipid particles such as low-density lipoproteins (LDL)).<sup>(106)</sup> The proteins and lipids in LDLs are good targets for oxidation, and the oxidized forms of LDL are now strongly implicated in formation of fatty lesions (atheromas) in artery walls. The apoprotein B component of these particles is fragmented by oxidation.

A variety of lipid peroxidation products, including adducts of lipid and apoprotein B, are produced. LDL particles contain a significant amount of vitamin E and carotinoids that may serve as primary antioxidants.<sup>(98)</sup>

### **Iron in human atheroma and LDL oxidation by macrophages following erythrophagocytosis**

The oxidative modification of low density lipoprotein (LDL) within vessel walls, as well as its uptake by macrophages, are considered important steps in the development of the early atheromatous lesion.<sup>(107)</sup>

It has been shown that LDL modification is due to free-radical chain reactions and that both copper and iron in reduced forms are able to induce such processes.<sup>(108)</sup>

Macrophages after engulfing aged or damaged erythrocytes, or other cellular material, may contain secondary lysosomes with catalytically active low-molecular weight iron complexes. If such macrophages are located in the arterial wall in the close vicinity of LDL

particles, these cells might exocytose iron with reduction of ferric iron into its catalytically active ferrous form. An iron dependent oxidative process may thus be initiated, leading to oxidation and conversion of the LDL to a form that promotes lipid accumulation in macrophages.<sup>(109)</sup>

Iron could promote the modification of LDL by initiating lipid peroxidation or by promoting the breakdown of lipid hydroperoxides.<sup>(110,111)</sup>

The mechanisms by which iron initiates lipid peroxidation are far from clear. It could catalyse the formation of hydroxyl radicals by a Haber Weiss reaction or Fenton reaction.<sup>(112)</sup>



or could possibly form preferryl ions ( $\text{Fe}^{2+}\text{-O}_2$ ) or a ferrous-dioxygen-ferric complex.

These species could then initiate lipid peroxidation. Both  $\text{Fe}^{2+}$  and soluble  $\text{Fe}^{3+}$  complexes promote the conversion of lipid hydroperoxides to free radicals (lipid alkoxy and peroxy radicals respectively).<sup>(113)</sup>

These radicals can stimulate the chain reaction of lipid peroxidation or can fragment by complex reactions to form a variety of compounds including aldehydes, which may react with apolipoprotein B-100.<sup>(114,115)</sup>

Iron has been shown to bind to phospholipids in liposomes and microsomes and this raises the interesting possibility that iron-mediated free-radical generation may actually occur in the surface of LDL particles.<sup>(116)</sup>

# Aim of THE WORK

## **AIM OF THE WORK**

The aim of the present work is to study serum iron and ferritin in patients with ischemic heart disease in a trial to evaluate their possible role in coronary atherosclerosis.

# MATERIAL & METHODS

## MATERIAL AND METHODS

This study included:

**- Thirty patients with ischemic heart disease.**

They were subdivided into two subgroups:

***A- Fifteen with stable angina.***

***B- Fifteen with unstable angina pectoris.***

Unstable angina pectoris was diagnosed if:

1- Angina worsened in term of frequency, intensity, ease of provocation.

2- Rest pains, nocturnal and prolonged episodes.

3- Recent onset of typical angina within one month, provided that no ECG or enzymatic evidence of infarction.

**- Ten apparently healthy subjects as control group.**

All the subjects were of comparable age and sex, and clinically free.



## METHODS

To all subjects included in the present study, the following were done:

### **1- Thorough history taking:**

With special stress on presenting complaint including:

- Chest pain; its type, duration and radiation.
- Other symptoms as dyspnea and orthopnea.
- Autonomic symptoms as nausea, vomiting and excessive sweating.

### **2- Full clinical examination.**

### **3- Standard twelve lead ECG.**

### **4- Laboratory investigations.**

*Sample collection:*

- All blood samples were taken in the morning after at least 12 hours fasting.
- Subjects were seated for 20 minutes before withdrawal of blood.
- About 10 ml of blood were withdrawn from the anticubital vein with minimal stasis using disposable plastic syringes.
- The ten ml were divided as follows:
  - \* 2ml into a tube containing EDTA for complete blood picture.Then this blood were centrifuged and separated plasma was used for assay of TBARs.

\* A serum sample was obtained. The separated serum was used for assay of fasting blood sugar, kidney function, lipid survey test, tests to assess iron status and superoxide dismutase.

- An early morning specimen of urine was collected for every subject and examined as soon as possible the same morning.

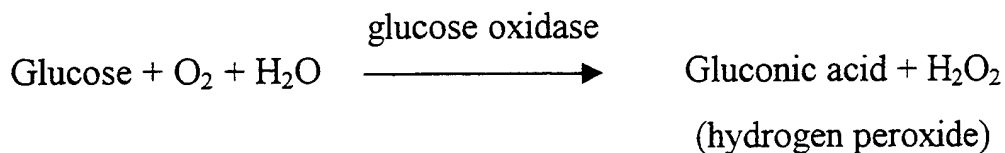
*Laboratory investigations included in the present study:*

- Urine analysis.<sup>(117)</sup>
- Complete blood picture.<sup>(118)</sup>

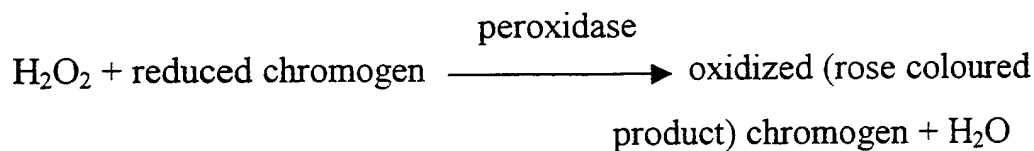
**Fasting blood glucose level: (bioMérieux kits)<sup>(119)</sup>**

It was determined by using an enzymatic method based on the following reactions:

**1st step: (specific for glucose)**



**2nd step: (not specific for glucose)**



The rose coloured product which is proportionate to the concentration of glucose in the sample (T) was measured spectrophotometrically at  $\lambda 505$  nm.

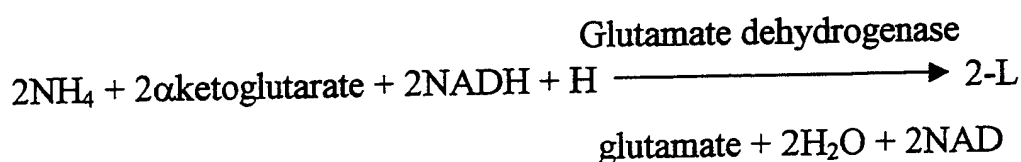
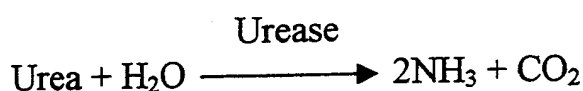
The concentration of glucose was determined after comparison with a standard glucose solution of known concentration (mg/dl) (S) similarly treated and glucose was calculated as follows:

$$\text{mg glucose/dl} = T/S \times C_s \text{ (mg/dl)}$$

$$\text{mmol glucose/L} = \text{mg/dl} \times 0.055$$

**- Serum urea: (bioMerieux kits)<sup>(120)</sup>**

It was determined by an ultraviolet kinetic method (urease/glutamate dehydrogenase coupled system) according to the following reactions.



The sample without deproteinization was taken, mixed well with the reagents and the variation in optical density per minute was read at 30 and at 90 seconds ( $\Delta$ ) at  $\lambda 340$  nm (T).

A standard urea solution of a known concentration (mg/dl) (S) was similarly treated and urea was *calculated as follows*:

$$\text{mg urea/dl} = \Delta T / \Delta S \times C_s \text{ (mg/dl)}$$

$$\text{mmol urea/L} = \text{mg/dl} \times 0.166$$

**Serum creatinine: (bioMerieux kits)<sup>(121)</sup>**

It was determined kinetically without deproteinization. The complex formed by creatinine in the sample and picric acid in an alkaline medium (sodium hydroxide) was measured at an interval of 1 min at  $\lambda$  500 nm. A standard creatinine of a known concentration (mg/dl) was similarly treated.

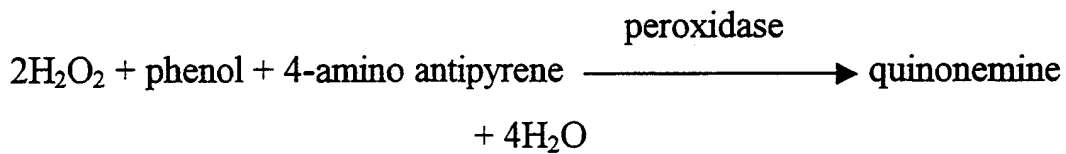
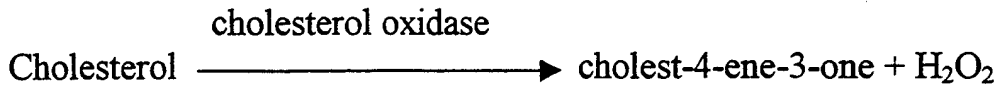
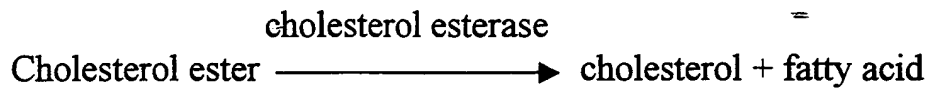
The difference in optical density at 20 and 80 seconds ( $\Delta$ ) was used to determine the creatinine in the sample (T) and standard (S) according to the following equation:

$$\text{mg creatinine/dl} = \Delta T / \Delta S \times C_s \text{ (mg/dl)}$$

$$\text{mmol creatinine/L} = \text{mg/dl} \times 0.0884$$

**Total serum cholesterol:<sup>(122)</sup>**

It was determined enzymatically without deproteinization according to the following reactions:



The rose coloured chromogen which was proportionate to cholesterol concentration in the sample (T) was measured at  $\lambda$  505 nm, and compared to the colour of a standard of a known cholesterol concentration (mg/dl) similarly treated (S).

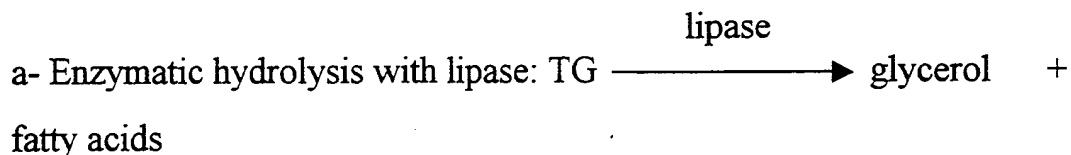
The concentration of cholesterol was obtained by the following equation:

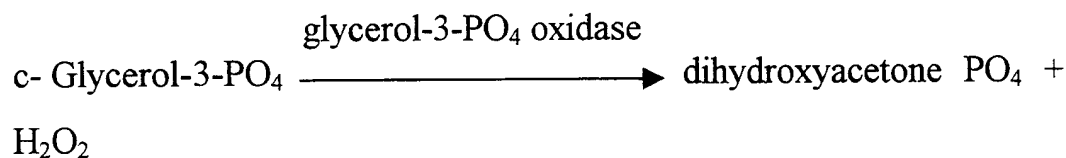
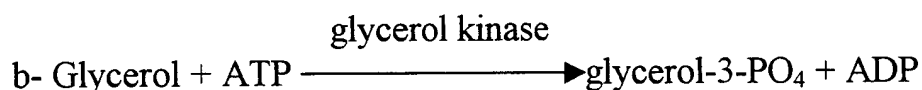
$$\text{mg cholesterol/dl} = T/S \times C_s \text{ (mg/dl)}$$

$$\text{mmol cholesterol/L} = \text{mg/dl} \times 0.026$$

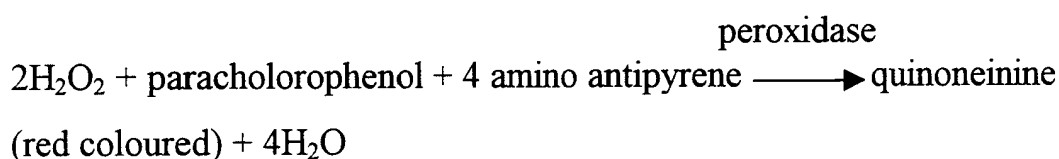
### **Serum triglycerides (triacylglycerol) (TG): (bioMerieux kits)<sup>(123)</sup>**

It was determined enzymatically without deproteinization according to the following reaction:





**d- Colour development:**



The rose coloured chromogen which is proportionate to TG concentration in the sample (T) was measured spectrophotometrically at  $\lambda$  505nm and compared to a standard of known TG concentration (mg/dl) similarly treated (S).

**TG level was obtained by the following equation:**

$$\text{mgTG/dl} = \text{T/S} \times \text{Cs (mg/dl)}$$

$$\text{mmol triglycerides/L} = \text{mg/dl} \times 0.01145$$

**High density lipoprotein (HDL) - cholesterol:<sup>(124)</sup>**

It was determined by mixing the plasma with HDL precipitating mixture (phosphotungstic acid and magnesium chloride) that precipitates both VLDL and LDL together with chylomicrons in the sample.

In the supernatant obtained after centrifugation (containing the HDL), the cholesterol content was determined according to the Liebermann-Burchard (L-B) reaction and read spectrophotometrically at  $\lambda$  620 nm (T), and compared to a standard cholesterol of known concentration (mg/dl) (S).

**HDL cholesterol was obtained according to the following equation:**

$$\text{mg HDL chol/dl} = T/S \times C_s \text{ (mg/dl)}$$

$$\text{mmol HDL-cholesterol/L} = \text{mg/dl} \times 0.026$$

**Low density lipoprotein (LDL)- cholesterol:<sup>(125)</sup>**

It was determined by mixing the plasma with heparin calcium chloride mixture which upon centrifugation left the HDL in the supernatant to be decanted.

Determination of LDL in the precipitate (pellet) was done according to the Liebermann-Burchard reaction (T), read spectrophotometrically at  $\lambda$  620 nm, and compared to a known standard cholesterol solution (mg/dl) (S).

**LDL chol was calculated as follows:**

$$\text{mg LDL - chol/dl} = T/S \times C_s \text{ (mg/dl)}$$

$$\text{mmol LDL - chol/L} = \text{mg/dl} \times 0.026$$

**Thiobarbituric acid reactive substances (TBARs):**<sup>(126,127)</sup>

The thiobarbituric acid reactive substances assay measured the amount of malondialdehyde (MDA) and malondialdehyde like substances formed by the lipid peroxidation process.

It involved heating the sample under acidic condition with thiobarbituric acid (TBA) which reacts with MDA to form a red MDA-TBA complex (T) which was measured spectrophotometrically at  $\lambda$  532 nm and compared to a standard similarly treated (S).

**Reagents:**

a- Thiobarbituric acid:

0.7% in 10% Trichloro acetic acid (0.7 gm TBA was dissolved in 100 ml 10% trichloroacetic acid).

b- Standard used was malondialdehyde-bis-diethyl acetate (1,1,3,3 tetraethoxypropane, (4 molar - 97% pure-density of 0.91).

The stock solution was diluted 1:1000 with 50 mmol/L (0.05 Molar) sulphuric acid ( $H_2SO_4$ ) and from this dilution 0.25 ml was taken and completed to 50 ml in a volumetric flask with 50 mmol/L  $H_2SO_4$ .

With each run 2 standard concentrations were done, the first was equivalent to 20  $\mu\text{mol/L}$  and the second was equivalent to 10  $\mu\text{mol/L}$  as follows:



0.3 ml was taken from the last dilution which was equivalent to 20  $\mu\text{mol/L}$ .

0.15 ml and completed to 0.3 ml with dist  $\text{H}_2\text{O}$  which was equivalent to 10  $\mu\text{mol/L}$ .

**Specimens:**

Two ml heparinized blood were put on 50  $\mu\text{L}$  of 3.8% EDTA (that acted as a preservative for TBARs), centrifuged and plasma was separated.

**Procedure:**

To 300  $\mu\text{L}$  of plasma, 4 ml of thiobarbituric acid were added, in a centrifuge tube and incubated for 30 minutes at  $100^\circ\text{C}$ .

The tubes were left to cool down, centrifuged for 15 minutes at 1500 g and the supernatant was read spectrophotometrically at  $\lambda$  532nm (T) using distilled water as a blank.

A similarly treated standard of a 10  $\mu\text{mol/L}$  concentration was done with each run.

**Calculation:**

$$\mu\text{mol TBARs/L} = T/S \times 10$$

**Serum Ferritin: (ELAgen kits)<sup>(128)</sup>****Principle:**

The ELAgen ferritin is based on the principle of a solid phase enzyme-linked immunosorbent assay.

The assay utilizes rabbit anti-ferritin bound to solid phase microwells and mouse monoclonal antiferritin in the antibody-enzyme (horseradish peroxidase) conjugate solution.

The test sample is allowed to react simultaneously with the antibodies, resulting in the ferritin molecules being sandwiched between the solid phase and enzyme-linked antibodies.

After a 45 minute incubation at room temperature, the wells are washed with water to remove unbound antibody complexes.

A solution of 3,3',5,5' - tetramethylbenzidine (TMB) is added and incubated for 15 minutes, resulting in the development of a blue colour.

The colour development is stopped by the addition of stop solution, and the resulting yellow colour is measured spectrophotometrically at 450 nm.

The concentration of ferritin is directly proportional to the colour intensity of the test sample.

**Reagents:****- EI Agen ferritin microplate:**

The bag contains a microplate of 12 strips X 8 wells.

Each well is coated with rabbit antiferritin antibodies.

**- EI Agen HRP-anti-ferritin conjugate:**

The vial contains 13 ml of mouse monoclonal anti-ferritin conjugated to horseradish peroxidase.

**Preservatives:**

Gentamicin sulphate 0.005%, thimerosal 0.02%, phenol 0.006%, Kathon CG 0.002%. Ready to use.

**- EI Agen ferritin standards:**

The vials contain ferritin in bovine serum with preservatives, the concentrations of the standards are the following: 0,15,80,250,500 and 1000 ng/ml.

**Lyophilized preservatives:**

Gentamicin sulphate, 0.005%, thimerosal 0.02%, phenol 0.006%, Kathon CG 0.002%.

**- EI Agen TMB H<sub>2</sub>O<sub>2</sub> HS:**

The vial contains 13 ml of a stabilized mixture of TMB (3,3',5,5' - tetramethylbenzidine) and H<sub>2</sub>O<sub>2</sub> (hydrogen peroxide) ready to use.

**- ELAgen stop solution (H<sub>2</sub>SO<sub>4</sub>):**

The vial contains 13 ml of 0.3 M H<sub>2</sub>SO<sub>4</sub>. Ready to use.

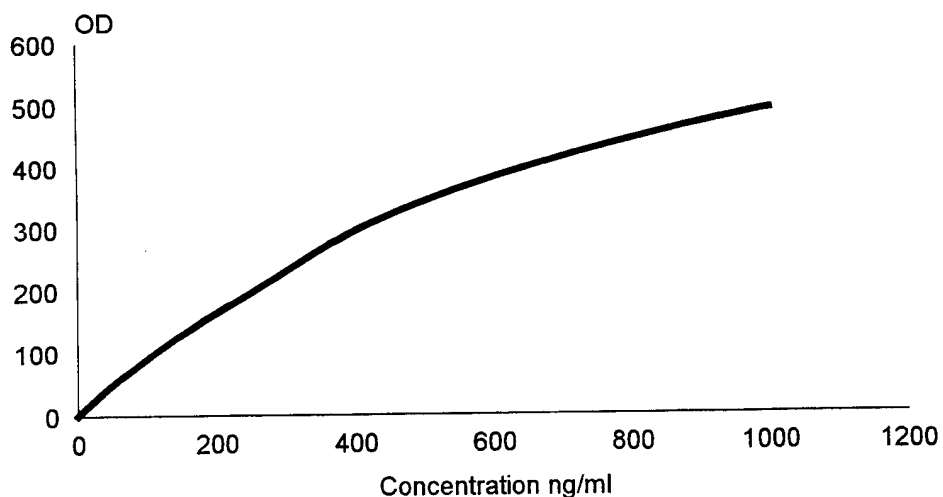
**Procedure:**

- 1- The desired number of coated wells were secured in the holder.
- 2- 20 µL of standards, samples, and controls were dispensed into appropriate wells.
- 3- 20 µL of HRP-anti-ferritin conj. were dispensed into each well.
- 4- The plate was thoroughly shaken for 10 seconds.
- 5- Then the plate was incubated at room temperature for 45 minutes.
- 6- Incubation mixture was removed by flicking the plate contents into a suitable waste container.
- 7- The microwells were rinsed and flicked 5 times with distilled water.
- 8- The inverted wells were struck sharply on absorbent paper to remove residual water droplets.
- 9- 100 µL TMB substrate reagent were dispensed into each well.
- 10- The plate was incubated at room temperature in the dark for 15 minutes.
- 11- The reaction was stopped by adding 100 µL of stop solution to each well.
- 12- The absorbance was read at 450 nm.

**Calculation:**

- 1- The mean absorbance values for the duplicate reference standards, controls and patient samples were calculated.
- 2- A standard curve was constructed on graph paper by plotting the mean absorbance obtained for each reference standard against its concentration in ng/ml. with OD on the vertical axis (y) and concentration on the horizontal axis (X).
- 3- The mean OD value for each sample was used to determine the corresponding concentration of ferritin in ng/ml from the standard curve.
- 4- Any diluted samples was multiplied by the appropriate dilution factor.

## Ferritin standard curve.



**Serum iron: (Ferrimat-kit)<sup>(129)</sup>****Principle:**

In an acid medium and in the presence of guanidine, ferric ions are released from their protein bounds.

Using hydroxylamine, the iron is reduced from the ferric to the ferrous state which forms a coloured complex with ferrozine. The colour was read spectrophotometrically at  $\lambda$  562 nm.

**Calculation:**

$$\mu\text{g/dl} = \frac{\text{A sample} - \text{A sample blank}}{\text{A standard}} \times n$$

$$n = 200$$

$$\mu\text{mol iron /L} = \mu\text{g/dL} \times 0.179$$

**Serum TIBC:<sup>(129)</sup>****Principle:**

Serum total iron binding capacity was measured after saturation of the transferrin with iron solution.

The excess iron was adsorbed on magnesium hydroxycarbonate, then determination of iron was performed.

**Calculation:**

$$\mu\text{g/dl} = \frac{\text{A sample} - \text{A sample blank}}{\text{A standard}} \times n$$

$$n = 600$$

$$\mu\text{mol TIBC/L} = \mu\text{g/dL} \times 0.179$$

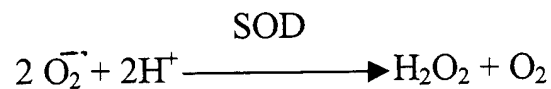
$$\text{Transferrin \% saturation} = \frac{\text{serum iron concentration}}{\text{TIBC}} \times 100$$

### Superoxide dismutase (SOD):<sup>(130)</sup>

#### Principle:

The method depend on the spontaneous autoxidation of pyrogallol at alkaline pH, resulting in the production of superoxide anion radical  $O_2^-$  which turn enhances the autoxidation of pyrogallol. Autoxidation is manifested as an increase in absorbance at 420 nm.

The presence of superoxide dismutase in the medium leads to removal of superoxide anion.



Thereby inhibiting the autoxidation of pyrogallol.

#### Reagents:

1- 50 mM tris Hcl buffer containing 1 mM DTPA (diethylene-triamino penta acetic acid).

2- 20 mM pyrogallol (1,2,3' benzentriol) in 10 mM Hcl. Stock solutions in 10 mM Hcl are stable for weeks.

3- Bovine Cu-Zn superoxide dismutase (erythrocyuprein) was prepared from erythrocytes. A standard solution was prepared in double distilled  $H_2O$ .

**Procedure:**

Blank: in a spectrophotometer cuvette maintained at 30°C, 1ml of the tris Hcl buffer was added.

+ 10  $\mu$ L pyrogallol.

The absorbance was followed for 1 min at 420 nm.

The rate of increase in absorbance under these condition was 0.02/min.

**Standard:**

The same procedure for the blank was repeated except the addition of SOD before recording absorbance.

- 1 ml tris buffer.

- 10  $\mu$ L pyrogallol.

- 30  $\mu$ L standard solution.

**Sample:** The procedure processed as standard.

**SOD unit:**

1 unit SOD activity is the amount of enzyme which inhibit the rate of autoxidation of pyrogallol by 50% from standard curve.



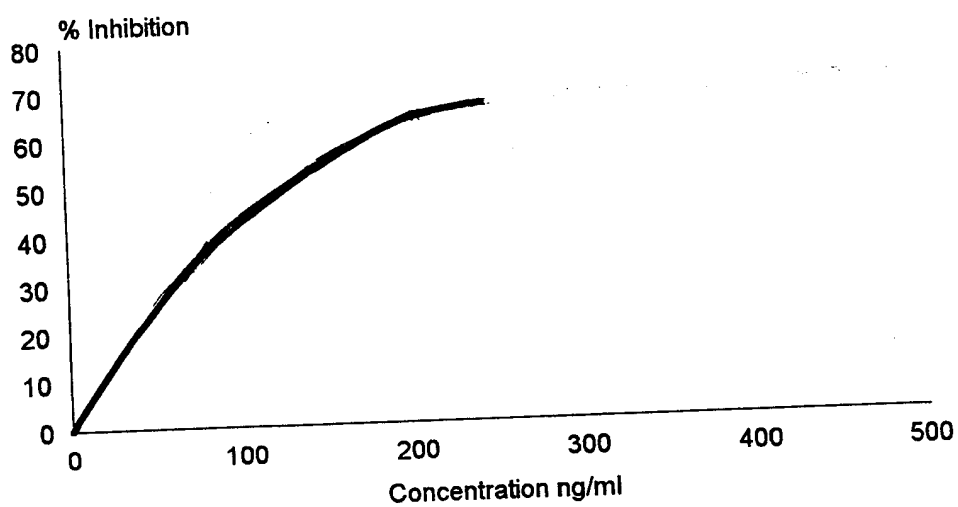
**Calculation:**

% of inhibition of pyrogallol autoxidation =

$$100 - \frac{\text{absorbance/min (standard or sample)}}{\text{absorbance/min (blank)}} \times 100$$

Standard curve was made by blotting the amount of SOD in (ng) versus % of inhibition of autoxidation. The amount of SOD was determined from standard curve and expressed in (ng/ml).

## Standard curve of SOD.



## Statistical analysis<sup>(131)</sup>

The row data were coded and transmitted into coding sheets. The coded data were typed onto a computer file using the D base program. The coded data were checked for transmission error. The data were analyzed using the SPSS statistical software. The personal computer IPM compatible was employed for statistical analysis, which include:

### 1- Arithmetic mean (X): It was calculated as follows:

$$X = \frac{\sum X}{n}$$

Where  $\sum X$  = sum of observations  
 $n$  = number of observations.

### 2- Standard deviation (S.D.): It was calculated as follows:

$$S.D = \sqrt{\frac{\sum X^2 - (\sum X)^2 / n}{n - 1}}$$

Where  $\sum X^2$  = sum of squared observations  
 $(\sum X)^2$  = square of the sum of observations.

### 3- Standard error (S.E):

$$S.E. = \frac{S.D}{\sqrt{n}}$$

The results were considered significant if the probability “P” value was less than 0.05. The following tests of significance were used to study the association between different variables in the different groups:

#### 4- Student “t” test:

It is a test of significance for the difference between two sample means.

It can be calculated from the following equation:

$$t = \frac{X_1 - X_2}{\sqrt{(S.E_1)^2 + (S.E_2)^2}}$$

Where  $X_1$ = mean of the first sample  
 $X_2$ = mean of the second sample  
 $S.E_1$ = the standard error of the first sample  
 $S.E_2$ = the standard error of the second sample

#### 5- Pearson’s correlation:

This measure reports the strength of the relationship between two variables.

The “r” can have any value from -1.00 to +1.00. It is obtained by:

$$r = \frac{n(\sum XY) - (\sum X)(\sum Y)}{\sqrt{(n(\sum X^2) - (\sum X)^2)(n(\sum Y^2) - (\sum Y)^2)}}$$

Where  $n$  = the number of paired observations  
 $\Sigma X$  = the sum of the X-variable  
 $\Sigma Y$  = the sum of the Y variable  
 $\Sigma X^2$  = the sum of the squared X-variable  
 $(\Sigma X)^2$  = the square of the summed X-variable  
 $\Sigma Y^2$  = the sum of the squared Y-variable  
 $(\Sigma Y)^2$  = the square of the summed Y-variable.

# RESULTS

## RESULTS

### **Age and blood picture in the studied groups: (Table Ia,b)**

#### ***Age:***

In the present study control group consists of 10 subjects, their mean age value was  $54\pm 6.8$  years, in the group of total patients with angina (30 subjects) their mean age value was  $51.6\pm 5$  years, in the group of patients with stable angina the mean age value was  $51.4\pm 5.6$  years and in the group of patients with unstable angina the mean age value was  $51.7\pm 4.6$  years.

No significant difference between the studied groups was found.

#### ***Haemoglobin:***

In the present study, control group showed a mean haemoglobin value of  $13.7\pm 0.88$  G/dl while the group of whole patients with angina showed mean Hb value of  $12.6\pm 1.2$  G/dl. The group of patients with stable angina showed mean Hb value of  $12.5\pm 1.1$  G/dl. and in the group of patients with unstable angina the mean Hb value was  $12.5\pm 1.2$  G/dl.

A significant difference was found when the total patients with angina group, patients with stable angina group and patients with unstable angina group were compared with the control group ( $P < 0.01$ ).

No significant difference was found between the group of patients with stable angina and the group of patients with unstable angina.

***Haematocrit:***

In the present study control group showed mean Ht value of  $41\pm 2.3\%$ .

In group of total patients with angina mean Ht value was  $37.4\pm 4.6\%$ . This was significantly lower than that of the control group ( $P<0.01$ ).

In group of patients with stable angina mean Ht value was  $36.9\pm 4.9\%$ . This was significantly lower than that of the control group ( $P<0.01$ ).

In group of patient with unstable angina mean Ht value was  $37.9\pm 4.5\%$ . This was significantly lower than that of the control group ( $P<0.05$ ).

No significant difference was found between the group of patients with stable angina when compared with the group of patients with unstable angina.

**Table I a: Age, Blood haemoglobin (Hb) g/dl and Haematocrit (Ht) % in controls (group I) and total patients with angina (group II).**

Case No.	Controls			Total		
	Age	Hb G/dl	Ht %	Age	Hb G/dl	Ht %
1	55	13.3	40	59	14.5	44
2	45	12.8	39	56	13.8	41
3	50	15	45	43	13	40
4	60	13	40	56	13.5	40
5	65	12.9	39	59	14	42
6	57	12.8	38	46	13	40
7	58	14.4	41	49	11.1	32
8	60	14	41	55	12	32
9	46	14.2	43	59	11.1	30
10	47	15	44	46	11.7	32
11				50	12.8	39
12				49	12.2	39
13				47	13.3	40
14				53	11.1	30
15				45	11.8	32
16				54	11.1	31
17				55	12	39
18				56	14.5	43
19				57	13	40
20				45	12	39
21				46	13.2	40
22				53	12.8	40
23				51	12.8	39
24				52	11	30
25				47	11	31
26				54	14	41
27				55	11	32
28				47	13	40
29				59	14.6	44
30				45	12.2	39
X	54	13.7	41	51.6	12.6	37.4
SD	6.8	0.88	2.3	5	1.2	4.6
P1				Ns	<0.01	<0.01

**P1:** Statistical significance between the control group and the group of total patients with angina.



**Table Ib: Age, Blood haemoglobin (Hb) g/dl and Haemātocrit (Ht) % in patients with stable angina (group III) and patients with unstable angina (group IV).**

Case No.	Patients with stable angina			Patients with unstable angina		
	Age	Hb G/dl	Ht %	Age	Hb G/dl	Ht %
1	59	14.5	44	54	11.1	31
2	56	13.8	41	55	12	39
3	43	13	40	56	14.5	43
4	56	13.5	40	57	13	40
5	59	14	42	45	12	39
6	46	13	40	46	13.2	40
7	49	11.1	32	53	12.8	40
8	55	12	32	51	12.8	39
9	59	11.1	30	52	11	30
10	46	11.7	32	47	11	31
11	50	12.8	39	54	14	41
12	49	12.2	39	55	11	32
13	47	13.3	40	47	13	40
14	53	11.1	30	59	14.6	44
15	45	11.8	32	45	12.2	39
X	51.4	12.5	36.9	51.7	12.5	37.9
SD	5.6	1.1	4.9	4.6	1.2	4.5
P2	Ns	<0.01	<0.01			
P3				Ns	<0.01	<0.05
P4				Ns	Ns	Ns

**P2:** Statistical significance between the control group and the group of patients with stable angina.

**P3:** Statistical significance between the control group and the group of patients with unstable angina.

**P4:** Statistical significance between the group of patients with stable angina and the group of patients with unstable angina.

**Fasting blood glucose, urea, creatinine levels (mmol/L) in the studied groups (Table IIa,b):**

***Fasting blood glucose:***

In the present study the control group showed mean value of FBG  $4.7\pm 0.53$  mmol/L. The group of total patients with angina showed mean value of  $4.8\pm 0.58$  mmol/L, the group of patients with stable angina showed mean value of  $4.95\pm 0.6$  mmol/L and in the group of patients with unstable angina the mean value of FBG was  $4.7\pm 0.55$  mmol/L.

No significant difference was found between the studied groups.

***Urea:***

In the present study the mean value of urea was  $4.8\pm 0.46$  mmol/L in the control group, was  $5.14\pm 0.96$  mmol/L in the group of total patients with angina, was  $5.4\pm 0.86$  mmol/L in the group of patient with stable angina, and  $4.9\pm 1$  mmol/L in the group of patients with unstable angina.

No significant difference was found between the studied groups.

***Creatinine:***

In the present study the control group showed mean value of creatinine of  $0.079\pm 0.0166$  mmol/L, the group of total patients with angina showed mean value of  $0.072\pm 0.0179$  mmol/L, the group of patients with stable angina showed mean value of  $0.076\pm 0.0164$

mmol/L and the mean value of creatinine in the group of patients with unstable angina was  $0.068 \pm 0.019$  mmol/L

No significant difference was found between the studied groups.

**Table II a: Fasting blood glucose, urea and creatinine levels (mmol/L) in controls (group I) and total patients with angina (group II).**

Case No.	Controls			Total		
	FBG mmol/L	Urea mmol/L	Cr mmol/L	FBG mmol/L	Urea mmol/L	Cr mmol/L
1	4.24	5.16	0.10	4.18	6.49	0.10
2	4.84	4.83	0.08	5.50	5.66	0.08
3	5.28	5.66	0.09	4.13	4.16	0.07
4	5.34	4.50	0.06	4.45	5.33	0.06
5	5.34	5.33	0.05	4.73	5.99	0.07
6	4.46	5.00	0.10	6.05	4.50	0.06
7	3.85	4.33	0.09	5.28	5.99	0.07
8	4.57	4.83	0.07	5.45	6.49	0.10
9	4.18	4.66	0.07	4.56	4.50	0.10
10	4.40	4.16	0.08	4.70	4.83	0.08
11				5.56	5.66	0.08
12				4.68	5.00	0.07
13				4.84	6.66	0.08
14				4.40	5.49	0.08
15				5.78	4.00	0.04
16				4.95	6.16	0.08
17				4.51	4.50	0.08
18				4.35	4.66	0.08
19				4.07	6.16	0.05
20				4.79	5.99	0.08
21				4.07	5.33	0.10
22				5.61	5.83	0.04
23				5.39	3.50	0.04
24				4.24	5.33	0.08
25				4.46	3.33	0.05
26				4.68	5.66	0.04
27				5.34	3.83	0.07
28				4.68	3.50	0.08
29				3.96	5.33	0.08
30				5.50	4.33	0.07
X	4.70	4.80	0.079	4.80	5.14	0.072
SD	0.53	0.46	0.0166	0.58	0.96	0.0179
P1				Ns	Ns	Ns

**P1:** Statistical significance between the control group and the group of total patients with angina.

**Table II b: Fasting blood glucose, urea and creatinine levels (mmol/L) in patients with stable angina (group III) and patients with unstable angina (group IV).**

Case No.	Patients with stable angina			Patients with unstable angina		
	FBG mmol/L	Urea mmol/L	Cr mmol/L	FBG mmol/L	Urea mmol/L	Cr mmol/L
1	4.18	6.49	0.10	4.95	6.16	0.08
2	5.50	5.66	0.08	4.51	4.50	0.08
3	4.13	4.16	0.07	4.35	4.66	0.08
4	4.45	5.33	0.06	4.07	6.16	0.05
5	4.73	5.99	0.07	4.79	5.99	0.08
6	6.05	4.50	0.06	4.07	5.33	0.10
7	5.28	5.99	0.07	5.61	5.83	0.04
8	5.45	6.49	0.10	5.39	3.50	0.04
9	4.56	4.50	0.10	4.24	5.33	0.08
10	4.70	4.83	0.08	4.46	3.33	0.05
11	5.56	5.66	0.08	4.68	5.66	0.04
12	4.68	5.00	0.07	5.34	3.83	0.07
13	4.84	6.66	0.08	4.68	3.50	0.08
14	4.40	5.49	0.08	3.96	5.33	0.08
15	5.78	4.00	0.04	5.50	4.33	0.07
X	4.95	5.4	0.076	4.7	4.9	0.068
SD	0.6	0.86	0.0164	0.55	1.0	0.019
P2	Ns	Ns	Ns			
P3				Ns	Ns	Ns
P4				Ns	Ns	Ns

**P2:** Statistical significance between the control group and the group of patients with stable angina.

**P3:** Statistical significance between the control group and the group of patients with unstable angina.

**P4:** Statistical significance between the group of patients with stable angina and the group of patients with unstable angina.

**Blood lipid (cholesterol – HDL-C, LDL-C, TG) levels mmol/L in the studied groups Table III (a,b):**

***Total cholesterol:***

Mean total cholesterol value was  $4.3\pm 0.6$  mmol/L in the control group,  $6.3\pm 1.5$  mmol/L,  $6.3\pm 1.7$  mmol/L and  $6.3\pm 1.3$  mmol/L in the group of total patients with angina, stable angina group and unstable angina group respectively.

Significantly higher values was found in these three groups when compared with control group ( $P < 0.001$ ).

***High density lipoprotein cholesterol:***

Mean HDLC value was  $1.3\pm 0.2$  mmol/L in the control group,  $1\pm 0.2$  mmol/L and  $1\pm 0.2$  mmol/L (in group of total patients with angina and group of patients with stable angina). Significantly lower values was found between these two groups when compared with the control group ( $P < 0.01$ ), and  $1.03\pm 0.27$  mmol/L in group of patients with unstable angina and this was significantly lower than that of the control group ( $P < 0.02$ ).

***Low density lipoprotein cholesterol:***

Mean LDLC value was  $2.4\pm 0.5$  mmol/L in the control group,  $4.7\pm 1.3$  mmol/L,  $4.66\pm 1.5$  mmol/L and  $4.7\pm 1.1$  mmol/L in (group of total patients with angina, stable angina group and unstable angina group) respectively.

Significantly higher values was found in these three groups when compared with the control group ( $P<0.001$ ).

***Triglycerides:***

Mean TG value was  $1.1\pm 0.3$  mmol/L in the control group,  $1.4\pm 0.6$  mmol/L in the group of total patients with angina and this was significantly higher than that of control group ( $P<0.05$ ).

Mean TG level was  $1.47\pm 0.6$  mmol/L in the group of patients with stable angina and  $1.4\pm 0.7$  mmol/L in the group of patients with unstable angina.

No significant difference was found in these two groups when compared with the control group.

**Table III a: Blood lipid levels (mmol/L) in controls (group I), and total patients with angina (group II).**

Case No.	Controls				Total			
	TCh mmol/L	HDL-C mmol/L	LDL-C mmol/L	TG Mmol/L	TCh mmol/L	HDL-C mmol/L	LDL-C mmol/L	TG mmol/L
1	4.18	1.03	2.63	1.20	6.81	1.03	5.29	1.14
2	4.08	1.50	2.12	1.02	5.16	1.19	3.56	0.96
3	3.48	1.03	2.06	0.90	4.88	1.03	3.20	1.44
4	3.95	1.37	1.86	1.56	8.67	1.03	6.97	1.50
5	4.52	1.47	2.45	1.32	6.35	0.98	4.80	1.30
6	4.08	1.06	2.50	1.19	3.87	0.95	2.45	1.08
7	3.64	1.60	1.73	0.72	4.90	0.90	3.59	0.96
8	4.21	1.03	2.58	1.32	9.29	1.08	7.48	1.68
9	5.31	1.34	3.35	1.44	7.22	1.06	5.55	1.44
10	5.21	1.86	3.04	0.72	3.56	0.67	2.37	1.18
11					5.42	1.01	4.00	0.96
12					7.74	1.08	5.83	1.92
13					6.19	1.55	3.72	2.16
14					6.66	1.03	5.06	1.30
15					8.05	0.65	6.06	3.12
16					7.22	1.44	5.16	1.44
17					5.16	0.70	4.00	1.08
18					5.16	1.19	3.56	0.96
19					5.93	1.03	4.23	1.56
20					6.97	0.93	5.42	1.44
21					7.61	0.85	5.83	2.16
22					5.73	1.60	3.61	1.20
23					5.42	0.70	4.41	0.72
24					5.16	1.01	3.74	0.84
25					4.64	0.77	3.61	0.60
26					6.71	1.39	4.90	0.96
27					7.74	0.83	5.47	3.36
28					7.53	0.93	5.68	2.16
29					5.06	1.03	3.61	0.96
30					9.29	1.08	7.48	1.68
X	4.3	1.3	2.4	1.1	6.3	1.0	4.7	1.4
SD	0.6	0.2	0.5	0.3	1.5	0.2	1.3	0.6
P1					<0.001	<0.01	<0.001	<0.05

**P1:** Statistical significance between the control group and the group of total patients with angina.



**Table III b: Blood lipid levels (mmol/L) in patients with stable angina (group III) and patients with unstable angina (group IV).**

Case No.	Patients with stable angina				Patients with unstable angina			
	TCh mmol/L	HDL-C mmol/L	LDL-C mmol/L	TG mmol/L	TCh mmol/L	HDL-C mmol/L	LDL-C mmol/L	TG mmol/L
1	6.81	1.03	5.29	1.14	7.22	1.44	5.16	1.44
2	5.16	1.19	3.56	0.96	5.16	0.70	4.00	1.08
3	4.88	1.03	3.20	1.44	5.16	1.19	3.56	0.96
4	8.67	1.03	6.97	1.50	5.93	1.03	4.23	1.56
5	6.35	0.98	4.80	1.30	6.97	0.93	5.42	1.44
6	3.87	0.95	2.45	1.08	7.61	0.85	5.83	2.16
7	4.90	0.90	3.59	0.96	5.73	1.60	3.61	1.20
8	9.29	1.08	7.48	1.68	5.42	0.70	4.41	0.72
9	7.22	1.06	5.55	1.44	5.16	1.01	3.79	0.84
10	3.56	0.67	2.37	1.18	4.64	0.77	3.61	0.60
11	5.42	1.01	4.00	0.96	6.71	1.39	4.90	0.96
12	7.74	1.08	5.83	1.92	7.74	0.83	5.47	3.36
13	6.19	1.55	3.72	2.16	7.53	0.93	5.68	2.16
14	6.66	1.03	5.06	1.30	5.06	1.03	3.61	0.96
15	8.05	0.65	6.06	3.12	9.29	1.08	7.48	1.68
X	6.3	1.0	4.66	1.47	6.3	1.03	4.7	1.4
SD	1.7	0.2	1.5	0.6	1.3	0.27	1.1	0.7
P2	<0.001	<0.01	<0.001	Ns				
P3					<0.001	<0.02	<0.001	Ns
P4					Ns	Ns	Ns	Ns

**P2:** Statistical significance between the control group and the group of patients with stable angina.

**P3:** Statistical significance between the control group and the group of patients with unstable angina.

**P4:** Statistical significance between the group of patients with stable angina and the group of patients with unstable angina.

**Iron status in the studied groups (Table IV a,b):*****Serum iron:***

In the present study, control group showed a mean serum iron value of  $16.3 \pm 2.2$   $\mu\text{mol/L}$ , total patients with angina group showed a mean value of  $17.35 \pm 4.8$   $\mu\text{mol/L}$ , patients with stable angina group showed a mean value of  $17.7 \pm 4.6$   $\mu\text{mol/L}$  and patients with unstable angina showed a mean value of  $16.9 \pm 5.1$   $\mu\text{mol/L}$ .

No significant difference was found was found between the studied groups.

***TIBC:***

In the present study, control group showed a mean TIBC value of  $56.9 \pm 12.2$   $\mu\text{mol/L}$ , group of total patients with angina showed a mean value of  $63.1 \pm 7.6$   $\mu\text{mol/L}$ , patients with stable angina group showed a mean value of  $62.1 \pm 8.2$   $\mu\text{mol/L}$  and the group of patients with unstable angina showed a mean value of  $64 \pm 7$   $\mu\text{mol/L}$ .

No significant difference was found between the studied groups.

***Ferritin:***

In the present study, control group showed a mean ferritin level of  $62.5 \pm 13.8$   $\mu\text{g/L}$ . Mean ferritin levels were  $138.1 \pm 60$   $\mu\text{g/L}$ ,  $142.9 \pm 69.75$   $\mu\text{g/L}$  and  $133 \pm 50.9$   $\mu\text{g/L}$  in the total angina group patients, stable angina group and unstable angina group respectively. The levels were significantly higher in these 3 groups when compared to the control group ( $p < 0.001$ ).

**Table IV a: Iron status in controls (group I) and total patients with angina (group II).**

Case No.	Controls			Total		
	Fe $\mu\text{mol/L}$	TIBC $\mu\text{mol/L}$	Ferritin $\mu\text{g/L}$	Fe $\mu\text{mol/L}$	TIBC $\mu\text{mol/L}$	Ferritin $\mu\text{g/L}$
1	21.48	72.50	59	25.42	72.32	270
2	14.32	62.65	60	24.34	71.06	210
3	15.93	75.18	50	19.51	67.48	105
4	16.83	63.00	37	22.38	68.92	97
5	15.93	40.63	80	22.38	68.92	199
6	17.90	62.47	55	17.90	63.36	99
7	15.75	44.03	59	10.74	47.44	85
8	14.68	42.07	70	12.89	49.23	224
9	16.29	52.98	80	10.74	47.44	100
10	13.43	54.24	75	15.75	61.22	90
11				16.83	62.47	110
12				15.75	60.86	100
13				19.51	67.66	85
14				17.72	62.83	100
15				13.78	59.97	270
16				11.28	48.87	90
17				12.53	59.97	130
18				25.06	71.60	115
19				20.94	64.44	225
20				12.89	49.23	250
21				20.94	64.44	95
22				16.29	63.55	95
23				16.11	63.55	105
24				11.64	66.23	110
25				11.64	65.34	145
26				23.27	70.71	190
27				11.99	70.88	100
28				18.80	64.44	120
29				25.06	72.50	80
30				16.29	64.44	150
X	16.3	56.9	62.5	17.35	63.1	138.1
SD	2.2	12.2	13.8	4.8	7.6	60
P1				Ns	Ns	<0.001

P1: Statistical significance between the control group and the group of total patients with angina.

**Table IV b: Iron status in patients with stable angina (group III)  
and patients with unstable angina (group IV).**

Case No.	Patients with stable angina			Patients with unstable angina		
	Fe μmol/L	TIBC μmol/L	Ferritin μg/L	Fe μmol/L	TIBC μmol/L	Ferritin μg/L
1	25.42	72.32	270	11.28	48.87	90
2	24.34	71.06	210	12.53	59.97	130
3	19.51	67.48	105	25.06	71.60	115
4	22.38	68.92	97	20.94	64.44	225
5	22.38	68.92	199	12.89	49.23	250
6	17.90	63.36	99	20.94	64.44	95
7	10.74	47.44	85	16.29	63.55	95
8	12.89	49.23	224	16.11	63.55	105
9	10.74	47.44	100	11.64	66.23	110
10	15.75	61.22	90	11.64	65.34	145
11	16.83	62.47	110	23.27	70.71	190
12	15.75	60.86	100	11.99	70.88	100
13	19.51	67.66	85	18.80	64.44	120
14	17.72	62.83	100	25.06	72.50	80
15	13.78	59.97	270	16.29	64.44	150
X	17.7	62.1	142.9	16.9	64	133
SD	4.6	8.2	69.75	5.1	7	50.9
P2	Ns	Ns	<0.001			
P3				Ns	Ns	<0.001
P4				Ns	Ns	Ns

**P2:** Statistical significance between the control group and the group of patients with stable angina.

**P3:** Statistical significance between the control group and the group of patients with unstable angina.

**P4:** Statistical significance between the group of patients with stable angina and the group of patients with unstable angina.

**Plasma thiobarbituric acid reactive substances levels ( $\mu\text{mol/L}$ ) in the studied groups (Table Va,b):**

Mean plasma TBARs value was  $2.3\pm 0.78 \mu\text{mol/L}$  in control group,  $3.6\pm 1.4 \mu\text{mol/L}$  in the group of total patients with angina and it was significantly higher than that of the control group ( $p < 0.01$ ). It was  $3.4\pm 1.5 \mu\text{mol/L}$  in stable angina group, and it was significantly higher than that of control group ( $p < 0.05$ ) and was  $3.7\pm 1.2 \mu\text{mol/L}$  in unstable angina group and it was significantly higher than that of control group ( $P < 0.01$ ).

**Table V a: Plasma thiobarbituric acid reactive substances (TBARs) levels ( $\mu\text{mol/L}$ ) in controls (group I) and total patients with angina (group II).**

Case No.	Controls	Total
	TBARs $\mu\text{mol/L}$	TBARs $\mu\text{mol/L}$
1	1.2	3.4
2	1.8	5.4
3	2.4	2.0
4	3.5	3.5
5	2.5	2.2
6	2.7	3.3
7	1.4	3.8
8	3.0	5.6
9	3.3	2.5
10	1.9	6.2
11		2.0
12		2.0
13		5.4
14		2.2
15		2.3
16		3.3
17		3.9
18		6.0
19		4.1
20		2.3
21		2.9
22		2.1
23		2.5
24		5.3
25		3.8
26		3.9
27		3.0
28		3.4
29		6.2
30		3.5
X	2.3	3.6
SD	0.78	1.4
P1		<0.01

**P1:** Statistical significance between the control group and the group of total patients with angina.

**Table V b: Plasma thiobarbituric acid reactive substances (TBARs) levels ( $\mu\text{mol/L}$ ) in patients with stable angina (group III) and patients with unstable angina (group IV).**

Case No.	Patients with stable angina	Patients with unstable angina
	TBARs $\mu\text{mol/L}$	TBARs $\mu\text{mol/L}$
1	3.4	3.3
2	5.4	3.9
3	2.0	6.0
4	3.5	4.1
5	2.2	2.3
6	3.3	2.9
7	3.8	2.1
8	5.6	2.5
9	2.5	5.3
10	6.2	3.8
11	2.0	3.9
12	2.0	3.0
13	5.4	3.4
14	2.2	6.2
15	2.3	3.5
X	3.4	3.7
SD	1.5	1.2
P2	<0.05	
P3		<0.01
P4		Ns

**P2:** Statistical significance between the control group and the group of patients with stable angina.

**P3:** Statistical significance between the control group and the group of patients with unstable angina.

**P4:** Statistical significance between the group of patients with stable angina and the group of patients with unstable angina.

**Serum superoxide dismutase (SOD) level (ng/ml) in the studied groups (Table VI a,b):**

In the present study mean serum SOD value was  $30 \pm 8.6$  ng/ml in the control group,  $61.8 \pm 13.3$  ng/ml,  $60.8 \pm 14.6$  ng/ml and  $62.7 \pm 12.4$  ng/ml in the group of total patients with angina, stable angina group, unstable angina group respectively.

Significantly higher values were found in these three groups when compared with the control group ( $P < 0.001$ ).



**Table VI a: Serum superoxide dismutase (SOD) levels (ng/ml) in controls (group I) and total patients with angina (group II).**

Case No.	Controls	Total
	SOD ng/ml	SOD ng/ml
1	35	73
2	30	60
3	26	56
4	40	73
5	17	75
6	21	40
7	40	45
8	30	90
9	21	71
10	40	65
11		45
12		50
13		40
14		65
15		65
16		62
17		58
18		73
19		59
20		58
21		45
22		73
23		56
24		71
25		56
26		40
27		65
28		60
29		90
30		75
X	30	61.8
SD	8.6	13.3
P1		<0.001

**P1:** Statistical significance between the control group and the group of total patients with angina.

**Table VI b: Serum superoxide dismutase (SOD) levels<sup>±</sup>(ng/ml) in patients with stable angina (group III) and patients with unstable angina (group IV).**

Case No.	Patients with stable angina	Patients with unstable angina
	SOD ng/ml	SOD ng/ml
1	73	62
2	60	58
3	56	73
4	73	59
5	75	58
6	40	45
7	45	73
8	90	56
9	71	71
10	65	56
11	45	40
12	50	65
13	40	60
14	65	90
15	65	75
X	60.8	62.7
SD	14.6	12.4
P2	<0.001	
P3		<0.001
P4		Ns

**P2:** Statistical significance between the control group and the group of patients with stable angina.

**P3:** Statistical significance between the control group and the group of patients with unstable angina.

**P4:** Statistical significance between the group of patients with stable angina and the group of patients with unstable angina.

# DISCUSSION

## DISCUSSION

Extensive reviews have been published concerning the role of iron in free radical reactions, such as lipid peroxidation. Briefly, except in states of iron overload, all iron in human serum is bound to proteins. (39, 132-134)

About two thirds of body iron is found in hemoglobin, with smaller amounts in myoglobin, various enzymes, and the transport protein transferrin. Iron not required for these is largely stored in ferritin.<sup>(39)</sup>

To promote free radical production, iron must be liberated from proteins. It is believed that oxidant stress itself can provide the iron necessary for formation of reactive oxygen species, for example, by mobilizing iron from ferritin or by degrading heme proteins to release iron.<sup>(39)</sup> For instance, superoxide radicals have been observed to liberate iron from ferritin, promoting lipid peroxidation.<sup>(39,135)</sup>

Halliwell and Gutteridge have proposed that the general effect of iron catalysts is to convert poorly reactive free radicals into highly reactive ones, such as the hydroxyl radical.<sup>(39)</sup>

Balla and Coworkers<sup>(136)</sup> demonstrated that the combination of physiological concentrations of hydrogen peroxide and hemin induce a rapid peroxidation of LDL in vitro and that free iron is released from the degraded heme ring. They also observed a simultaneous loss of reactive lysine amino groups of LDL, which has

simultaneous loss of reactive lysine amino groups of LDL, which has been shown to promote LDL uptake by scavenger receptors of macrophages leading to foam cell formation.<sup>(137)</sup>

Also transition metal ions are probably required for the peroxidation of LDL by monocyte/macrophages, smooth muscle cells, and endothelial cells.<sup>(138,139)</sup>

There is little doubt that an excessive amount of stored iron is cardiotoxic. Myocardial failure is a prominent feature of haemochromatosis and in diseases associated with large exogenous iron loads, such as thalassaemia major.<sup>(140-142)</sup>

Patients with these diseases usually have iron stores many times higher than normal. The mechanism of myocardial damage by iron is not clearly understood. It is dependent on poorly defined effects within the myocardium, where iron either damages the muscle itself or excites fibrosis.<sup>(143)</sup>

A role for iron in the development of coronary lesions is not indispensable to the hypothesis that iron depletion protects against IHD. Iron depletion could exert a protective effect by mechanisms that don't involve effects on atherogenesis for example, by decreasing the vulnerability of the myocardium to ischemia.<sup>(144)</sup>

Iron depletion could inhibit atherogenesis through a variety of mechanisms including, for example, protection of endothelial cells from

oxygen radical injury<sup>(145)</sup> or inhibition of low density lipoprotein modification.<sup>(146,147)</sup>

Salonen et al,<sup>(148)</sup> reported that dietary iron intake to be positively associated with an increased risk of myocardial infarction. In contrast, Rimm et al,<sup>(149)</sup> Cooper and Liao<sup>(150)</sup> reported no association between dietary iron intake and the risk of coronary heart disease.

In the present study, when comparing the serum iron level in CHD patients with the corresponding values of control group, it was noticed that the serum iron level was ( $17.7 \pm 4.6$  mmol/L in stable group -  $16.9 \pm 5.1$  mmol/L in unstable group -  $17.35 \pm 4.8$  mmol/L in total group) and  $16.3 \pm 2.2$  mmol/L in control group and the difference was not statistically significant.

Also the values of serum TIBC in CHD patients ( $62.1 \pm 8.2$  mmol/L,  $64 \pm 7$  mmol/L,  $63.1 \pm 7.6$  mmol/L in stable, unstable and total groups respectively) was mildly higher than that of the control group ( $56.9 \pm 12.2$  mmol/L), but not to a significant degree.

These results are in agreement with study of Christopher et al,<sup>(151)</sup> several studies, which suggested that neither serum iron nor TIBC are related to the risk of CHD.

Normal serum iron, and TIBC levels noticed in the patient group of this study with elevated serum ferritin level could be due to the fact

that the effective iron in the process of lipid peroxidation is either the free iron or the ferritin molecule itself.

This can be further evidenced by the absence of any correlation between both serum iron, TIBC on one hand and the rest of risk factors (TCh, LDL-C, TG) on the other hand.

The concentration of serum ferritin in healthy individuals is positively correlated with the body iron stores.<sup>(152)</sup> In acute inflammation the serum ferritin level increases concomitantly with a decrease in serum iron acting as an acute phase reactant.<sup>(153)</sup>

Finlayson and Fraser had measured the ferritin level during the 1st 48 hours post AMI and found that the serum ferritin level gradually increases following AMI.<sup>(154)</sup>

The lack of acidic type ferritin in the serum (derived from myocardial cells) of AMI patients as tested by specific anti-heart and antiplacental ferritin McABS argues against the possibility that ferritin is released from damaged myocardial cells following infarction. This assumption is supported by the study of Cavanna et al, who reported a lack of heart ferritin in sera of patients examined 48 hours Post AMI.<sup>(155)</sup>

It has been reported that inflammation causes accumulation of iron in the reticuloendothelial cells, which may serve as an intracellular stimulus for ferritin synthesis.<sup>(156)</sup>

Inflammatory processes are characterized by an increase of mononuclear cells in the damaged area. In this sense, the iron binding neutrophilic glycoprotein, lactoferrin, plays an important role in the initial interaction with monocytes/macrophages during inflammation and is responsible for the inflammatory-reduced hypsideremia and accumulation of iron in these cells.<sup>(157)</sup>

Theoretically, ferritin as an iron binding protein may be considered as one of the primary antioxidants,<sup>(158)</sup> but many evidences contrast this hypothesis.<sup>(148,159,160)</sup>

Ferritin may be considered as a dangerous protein, as under certain conditions, the active oxygen species (superoxides) enter the ferritin-cores through the hydrophilic channels, followed by reduction of Fe(III) to Fe(II). This enables the release of free iron with its damaging effects.<sup>(148)</sup>

Nonenzymatic reduction of Fe(III) to Fe(II) occurs as iron is released from ferritin, thus ferritin is both a very efficient iron trap and a readily available source of iron for metabolic requirements and for formation of hemoglobin and other haemproteins.<sup>(93)</sup>

Salonen et al.,<sup>(148)</sup> reported an increased risk of acute myocardial infarction among men with serum ferritin level greater than 200  $\mu\text{g/L}$ , where as preliminary results reported by Stampfer et al. did not support an increase in this risk.<sup>(161)</sup>



In the present work serum ferritin level was significantly higher in the angina patient group either stable ( $142.9 \pm 69.75 \mu\text{g/L}$ ) or unstable ( $133 \pm 50.9 \mu\text{g/L}$ ) or the whole group ( $138.1 \pm 60 \mu\text{g/L}$ ) compared to the corresponding control value ( $62.5 \pm 13.8 \mu\text{g/L}$ ). This goes in hand with the elevated atherosclerotic risk lipid factors (cholesterol, LDL-C) and the diminished antirisk factor (HDL-C) as will be shown after words. It was noticed that the lipid profile of controls shows normal distribution and normal levels of different components. Their iron and ferritin levels where within the expected normal range.

There is a positive correlation between ferritin level and each of total cholesterol  $r$  (0.756) and LDL-C  $r$  (0.732). This would indicate the protective role of ferritin acting as an iron trap which would prevent its role in lipid peroxidation, hence guard against their atherogenic effects.

The present study revealed the following:

- A significant increase in serum total cholesterol level in CHD patients ( $6.3 \pm 1.7 \text{ mmol/L}$  in stable group -  $6.3 \pm 1.3 \text{ mmol/L}$  in unstable group -  $6.3 \pm 1.5 \text{ mmol/L}$  in total group) more than controls ( $4.23 \pm 0.6 \text{ mmol/L}$ ).
- A significant increase in serum LDL-C level in CHD patients either stable group ( $4.66 \pm 1.5 \text{ mmol/L}$ ) or unstable group ( $4.7 \pm 1.1 \text{ mmol/L}$ ) or

the whole group ( $4.7 \pm 1.3$  mmol/L) more than controls ( $2.4 \pm 0.5$  mmol/L).

In the control group TBARs mean level was ( $2.3 \pm 0.78$   $\mu\text{mol/L}$ ) and in the CHD patients, the mean level was ( $3.4 \pm 1.5$   $\mu\text{mol/L}$ ) in stable group, ( $3.7 \pm 1.2$   $\mu\text{mol/L}$ ) in unstable group ( $3.6 \pm 1.4$   $\mu\text{mol/L}$ ) in the whole group, which were significantly higher than that of the control group.

It has been proposed that alteration of LDL-C produces clearance by subendothelial macrophages and result in intracellular deposition of lipoprotein-derived cholesterol.<sup>(162)</sup>

A study in vitro had implicated modification of LDL-C by lipid peroxide products as one potential mechanism.<sup>(163)</sup> MDA is an end product of lipid peroxidation. Reaction of MDA with a critical number of lysine residues of the apolipoprotein B-100 of LDL-C produce internalization by the scavenger receptor of human monocyte-macrophages and the subsequent intracellular accumulation of lipoprotein-derived cholesterol ester in vitro.<sup>(164)</sup>

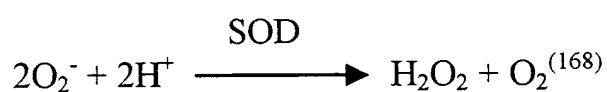
Other study revealed that higher serum MDA, LDL-C levels and correlation of serum MDA with the extent of CA lesions exist in patients with CHD during both the early and late stages of coronary atherosclerosis, suggesting the existence in vivo of protein modified by a physiological product of lipid peroxidation within arterial lesion.<sup>(165)</sup>

In the present study, HDL-C was significantly lower in CHD patients ( $1.0 \pm 0.2$  mmol/L) in stable group, ( $1.03 \pm 0.27$  mmol/L) in unstable group, ( $1.0 \pm 0.2$  mmol/L) in total group than controls ( $1.3 \pm 0.2$  mmol/L).

Serum TG level was higher in CHD patient either stable ( $1.47 \pm 0.6$  mmol/L) or unstable ( $1.4 \pm 0.7$  mmol/L) than controls ( $1.1 \pm 0.3$  mmol/L) but not to a significant degree. In the whole group the level was ( $1.4 \pm 0.6$  mmol/L) which is significantly higher than controls.

It has been shown that triglycerides are not an independent cardiovascular risk factor.<sup>(166)</sup> In addition, there is some limited evidence suggesting that the risk of CHD increases as triglyceride level increases in patients exhibiting high levels of total cholesterol or LDL-C and low levels of HDL-C.<sup>(167)</sup>

Superoxide dismutase (SOD) catalyzes the dismutation of the superoxide anion radical to hydrogen peroxide and oxygen.



Removal of excess  $\text{O}_2^-$  by SOD is an important antioxidant defense mechanism in aerobic organisms.<sup>(42,169)</sup> SOD is ubiquitous in oxygen-metabolizing cells, protecting against oxygen toxicity.<sup>(170-173)</sup>

Mn-SOD can be considered as an enzyme specific for mitochondria that could be used as a blood marker for cardiac mitochondrial damage as it is located predominantly in the mitochondria and is present in the heart in high concentrations.<sup>(174-178)</sup>

The rise in SOD observed in the present study in all patient groups denotes an active antioxidant mechanism that tries to counteract the risky lipid and ferritin adverse effects.

In the group of stable angina, they showed a significant correlation between SOD and TCh - LDL-C and ferritin indicating that SOD activity keeps pace with these risk factors, hence counteract their adverse effects.

Such correlations disappeared in the group of unstable angina patients, it could be postulated that failure of SOD protective action may play a role in the development of unstable condition.

Levels of SOD in CHD patient groups was (60.8±14.6 ng/ml) in stable group, (62.7±12.4 ng/ml) in unstable group, (61.8±13.3 ng/ml) in total group than controls (30±8.6 ng/ml).

SOD had been reported to rise in serum after AMI, remains high for 2-7 days then returns to basal level. The cause for this rise is thought to be due to its release from myocardial tissue (mitochondria), that takes longer time to pass to blood stream than CK-MB.<sup>(179)</sup>

This transient rise in SOD level following AMI, differs from the rise noticed in its level in the patient groups (stable, unstable or total) reported in the present work which is not produced as a result of cell injury (which actually does not exist in such patients) but rather to a process of over production, over activity to counteract the adverse oxidative stress evidenced by the higher cholesterol, LDL-C, TG, ferritin and TBARs.

# SUMMARY & CONCLUSION

## SUMMARY AND CONCLUSION

The oxidative modification of low density lipoprotein, as well as, its uptake by macrophages in vessel wall via scavenger receptors are considered important steps in the development of the atheromatous lesion. This uptake is not down regulated by the internalized cholesterol and cholesteryl esters and lead to their transformation to cells with the characteristic properties of lipid laden foam cells.

It has been shown that low density lipoprotein modification is due to free radical reaction and both copper and iron in reduced forms are able to induce such process.

To promote free radical production, iron must be liberated from proteins. It is believed that oxidant stress itself can provide the iron necessary for formation of reactive oxygen species, for example, by mobilizing iron from ferritin or by degrading heme proteins to release iron.

Theoretically, ferritin as an iron binding protein may be considered as one of the primary antioxidants that inhibits lipid peroxidation but many evidences contrast this hypothesis.

A reductive mechanism is probably responsible for the release of iron from ferritin into the low molecular weight pool, where it could induce lipid peroxidation and subsequent tissue injury by Fenton type reaction.

The aim of the present work is to study serum iron and ferritin in serum of patients with ischemic heart disease in a trial to evaluate their possible role in coronary atherosclerosis.

The present study was conducted on thirty patients with ischemic heart disease admitted to cardiology unit in the medical research institute as well as ten apparently normal healthy control subjects.

The selected patients included 15 with unstable angina pectoris and 15 with stable angina.

To all studied subjects included in the present study the following was done: full clinical examination, electrocardiogram, laboratory investigation including complete blood picture, complete urine analysis, serum fasting glucose, urea, creatinine, total cholesterol, high density lipoprotein cholesterol, low density lipoprotein cholesterol, triglycerides, iron, ferritin, superoxide dismutase and plasma thiobarbituric acid reactive substances.

*In the present study:*

Serum ferritin level was found to be significantly higher in the diseased groups when compared with control group.

On the other hand no differences were found among the diseased groups when compared with control group regarding both serum iron and serum total iron binding capacity levels.



Both serum superoxide dismutase and plasma thiobarbituric acid reactive substances were significantly higher in the diseased groups when compared with control group.

The result of this work showed a significant increase in serum total cholesterol, low density lipoprotein cholesterol in the diseased groups more than the control group. Also high density lipoprotein cholesterol was significantly lower in the diseased groups of patients when compared with the control.

Serum triglycerides level was significantly higher in the total group only when compared with control group in this study.

#### **Conclusions:**

- 1- Serum iron level is not critical in the process of lipid peroxidation. But serum ferritin is the determining factor being the source of iron.
- 2- The lipid peroxidation (as indicated by the thiobarbituric acid reactive substances) together with the increased atherogenic cholesterol both total, low density lipoprotein cholesterol and triglycerides all showed higher values in the diseased group.
- 3- As a result of the increased oxidative stress reported in the group of patients, Superoxide dismutase a component of antioxidant counterparts showed a reactive increase in the same groups.

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# PROTOCOL

# STUDY OF SERUM IRON AND FERRITIN IN PATIENTS WITH ISCHEMIC HEART DISEASES

دراسة مستوى الحديد والفيريتين في مصلى مرضى قصور الشرايين التاجية

Protocol of thesis submitted  
to Medical Research Institute  
Alexandria University  
In partial fulfillment of  
*Master Degree*

In Chemical Pathology

خطة بحث مقدمة  
لمعهد البحوث الطبية  
جامعة الإسكندرية  
إيفاءً جزئياً للحصول على  
درجة الماجستير

في الباثولوجيا الكيميائية

By

**Samah Saad Abd El-Aleem**

M.B.B.Ch  
(Alex.) 1992

من الطيبية

**سماح سعد عبد العليم**

بكالوريوس الطب والجراحة  
الإسكندرية ١٩٩٢

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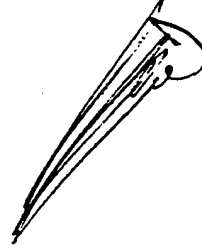
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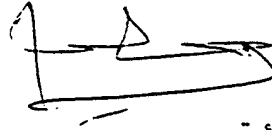


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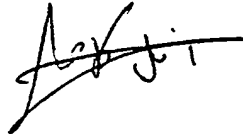


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## INTRODUCTION

The first step in atheroma formation is LDL oxidation,<sup>(1)</sup> the oxidized LDL is taken by phagocytic cells to form what is termed as “foam cells”: the cholesterol loaded macrophage-derived cells of the atherosclerotic lesion.<sup>(2)</sup>

Oxidized LDL is taken up by macrophages via scavenger receptors, leading to cholesterol accumulation.<sup>(3)</sup>

Native LDL, however, is not recognized by scavenger receptors and thus does not cause cholesterol accumulation in macrophages.<sup>(4)</sup>

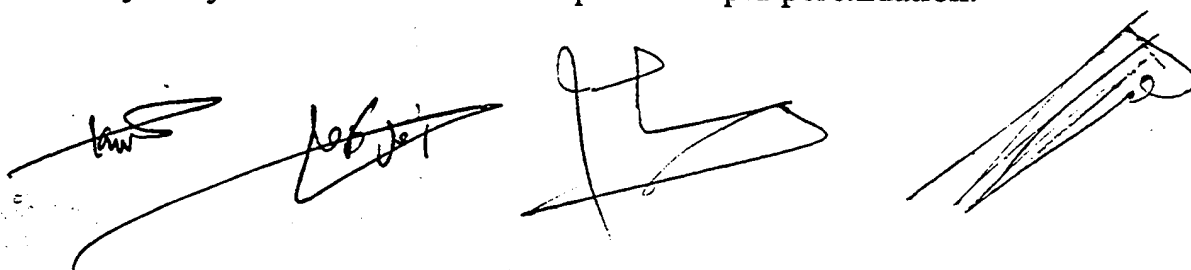
Monocytes, the precursors of macrophages are attracted towards oxidized LDL by chemotaxis and therefore oxidized LDL may attract monocytes into subendothelial space.<sup>(5)</sup>

It has been shown that LDL modification is due to “free radical reaction” and that both copper and iron in reduced forms are able to induce such process.<sup>(6)</sup>

To promote free radical reaction, iron is required promoting lipid peroxidation according to Fenton reaction.



This reaction liberates highly reactive oxygen species which is “hydroxyl radical”  $\text{OH}^\bullet$  that is capable of lipid peroxidation.<sup>(8)</sup>



It has been reported by some workers that increased level of plasma ferritin is associated with increase incidence of myocardial atherosclerosis due to increased LDL oxidation.<sup>(9)</sup>

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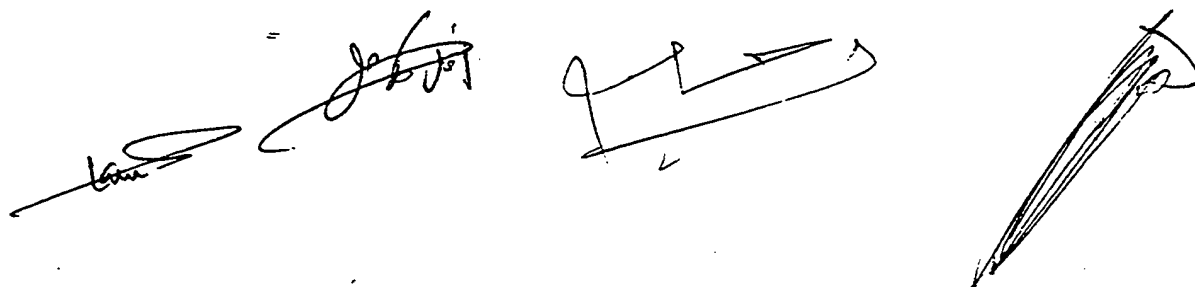
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## AIM OF THE WORK

The aim of the present work is to study serum iron and ferritin in a trial to evaluate their possible role in coronary atherosclerosis.

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# MATERIAL AND METHODS

## Material

Thirty subjects will be included in the study. They will be divided as follows:-

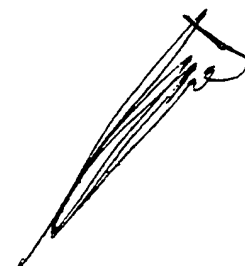
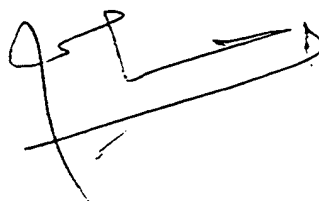
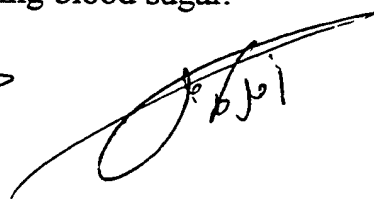
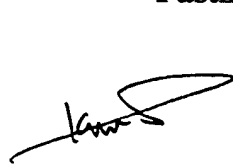
- 1- Fifteen patients with stable angina.
- 2- Fifteen patients with unstable angina.
- 2- Ten apparently normal healthy control subjects.

All subjects will be of comparable age, sex and socio-economic status.

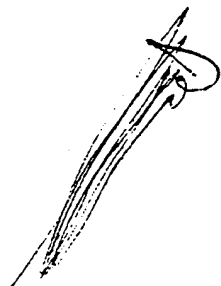
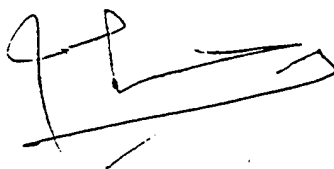
## Methods

To all studied subjects:

- 1- Thorough history taking and assessment of risk factors profile.
- 2- Full clinical examination.
- 3- Standard twelve lead ECG.
- 4- The following laboratory investigations will be done:
  - Urine analysis.<sup>(10)</sup>
  - From fasting patients (at least 12 hours) blood sample will be drawn for estimation of the following:-
    - Complete blood picture.<sup>(11)</sup>
    - Fasting blood sugar.<sup>(10)</sup>

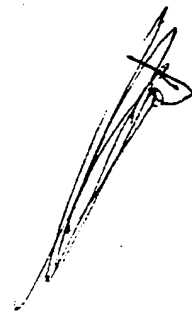
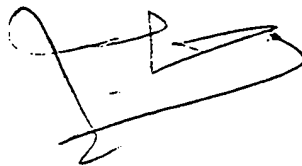
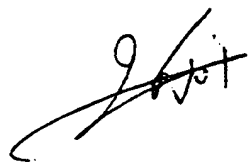


- Serum urea.<sup>(10)</sup>
- Serum creatinine.<sup>(10)</sup>
- Total serum cholesterol.<sup>(10)</sup>
- High density lipoprotein cholesterol.<sup>(12)</sup>
- Low density lipoprotein cholesterol.<sup>(13)</sup>
- Serum triglycerides.<sup>(10)</sup>
- Thiobarbituric acid reactive substances (TBARS).<sup>(14)</sup>
- Serum iron, ferritin, total iron binding capacity.<sup>(15)</sup>
- Serum superoxide dismutase (SOD).<sup>(16,17)</sup>



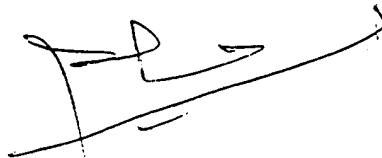
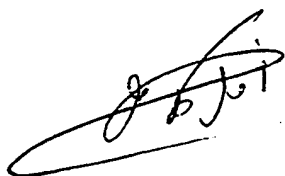
## RESULTS AND DISCUSSION

The results obtained from the study will be tabulated statistically, analyzed, compared with other available works and discussed.

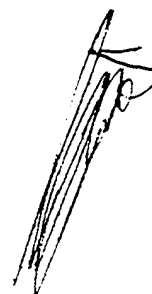
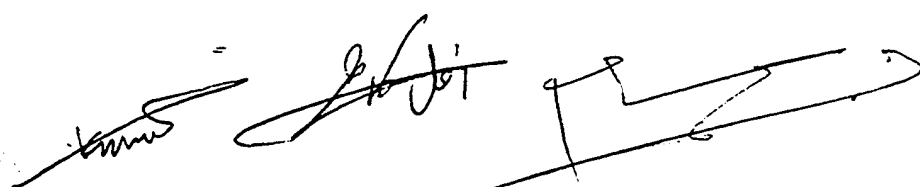


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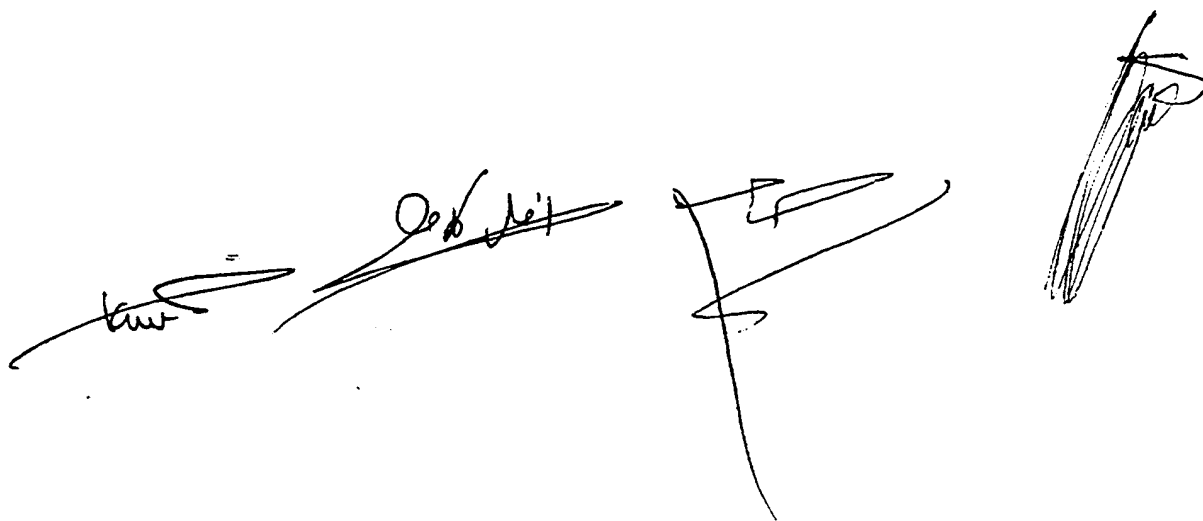
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The bottom of the page contains several handwritten marks. On the left, there is a signature that appears to be 'Liu'. In the center, there is a signature that appears to be 'J. Li'. To the right of these, there are several large, overlapping scribbles and lines, possibly representing a signature or a set of initials.

# ARABIC SUMMARY



## الملخص العربي

### دراسة مستوى الحديد والفيريتين في مصل مرضى

#### قصور الشرايين التاجية

يمثل التأكسد التحویری للدهنيات منخفضة الكثافة وكذلك مدى استيعابها بواسطة الخلايا الملتهمة خطوات مهمة في حدوث تصلب الشرايين خاصة وأن هذه العملية لا تقل بدخول الكوليستيرول وأملاحه داخل الخلايا وبهذا يتم تكوين الخلايا الرغوية المشبعة بالدهون.

كما ثبت أن تحور الدهنيات المنخفضة الكثافة يعود إلى وجود الشوارد الحرة والتي قد تحدث نتيجة تواجد عنصرى الحديد والنحاس والأمر يتطلب تحرير الحديد من بروتيناته الحاملة له وهذا قد يحدث نتيجة توافر الحمل التأكسدى ذاته. وعلى المستوى النظرى فإن مادة الفيريتين كمادة ضامة للحديد وعليه فإنه يمكن اعتبارها كمادة مضادة للأكسدة. ولكن الواقع يثبت العكس حيث أنه يمكن اعتبارها مخزناً ينبعث منه الحديد.

والغرض من هذه الرسالة هو دراسة مستوى الحديد والفيريتين في مصل مرضى قصور الشرايين التاجية في محاولة لتقييم أثرهما المحتمل في حدوث تصلب شرايين القلب.

وقد أجريت هذه الدراسة على ثلاثين مريض بمرض قصور الشرايين التاجية ممن دخلوا في قسم الأمراض الباطنة بمعهد البحوث الطبية وقد تم تقسيم مجموعة المرضى إلى نصفين يتكون كل منهما من خمسة عشر مريضاً ممن يعانون من مرض قصور الشرايين التاجية المستقر والغير مستقر هذا بالإضافة إلى مجموعة ضابطة مكونة من عشرة أشخاص طبيعيين.

وقد أجرى لكل شخص شملته هذه الدراسة فحص أكلينيكى كامل ورسم قلب كهربائى بالإضافة إلى الأبحاث المعملية الآتية: بول كامل - عد دم - سكر صائم - بولينا وكرياتينين - كوليستيرول كلى ومرتفع ومنخفض الكثافة ودهون ثلاثية وحديد المصل

والفيريئين وكلا من انزيم فوق الأوكسيد المانع للتحرور SOD ومستوى المركبات المتفاعلة مع حمض الثيوباربيتيوريك TBARS.

وقد وجد أن:

١- مستوى الفيريئين أظهر ارتفاعا ملحوظا في مجموعات المرضى عن المجموعة الضابطة.

٢- لم تظهر فروق في مستويات الحديد والقدرة الكلية لضم الحديد في مصل جميع من شملتهم الرسالة.

٣- أظهر كل من ال SOD وال TBARS ارتفاعا ملحوظا في المجموعات المرضية عن المجموعة الضابطة.

٤- أظهرت الدراسة ارتفاعا ملحوظا في تركيز كل من الكولستيرول الكلى ومنخفض الكثافة والدهون الثلاثية في المجموعات المرضية عن المجموعة الضابطة بينما انخفض مستوى الدهون المرتفعة الكثافة في مجموعة المرضى.

وبهذا يمكن الوصول للنتائج الآتية:

١- إن مستوى حديد المصل لا يمثل أهمية كبرى في عملية أكسدة الدهون بينما الفيريئين هو العامل المؤثر.

٢- إن عملية أكسدة الدهون كما يستدل عليها من زيادة ال TBARS بالإضافة إلى الزيادة في مستوى كل من الكولستيرول والدهنيات منخفضة الكثافة والدهون الثلاثية وكلها مؤشرات على تصلب الشرايين قد أظهرت ارتفاعا في المجموعة المرضية.

٣- نتيجة لإرتفاع هذه المواد فإن مستوى ال SOD قد ازداد كرد فعل لمعادلة التأثيرات الضارة للمواد المؤكسدة.

## المشرفون

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معهد البحوث الطبية

الدراسات العليا

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موافقة مجلس المعهد على منح الدرجة ٢٠١١ / ٦ / ١٢

يعتمد \*\*

وكيل المعهد

لدراسات العليا والبحوث

  
(أ.د. عزت محمد حش) ١٢

دراسة مستوى الحديد والفيريتين في صل  
مرضى قصور الشرايين التاجية

رسالة مقدمة

لمعهد البحوث الطبية

ايفاء جزئيا للحصول على

درجة الماجستير في

الباثولوجيا الكيميائية

من

الطبيبة/ **سماح سمك عبد العلي الشبي**

بكالوريوس الطب والجراحة

الإسكندرية (١٩٩٢)

معهد البحوث الطبية

جامعة الإسكندرية

٢٠٠١