# Separation of Lipoxygenase and Estimation of its Level in Blood of Males with Cardiovascular Disease

# Ameera A. Mahmood

Rawya N. Rashed

Department of Chemistry College of Education University of Mosul

Email: ameeraaziz 2000@yahoo.com

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#### **ABSTRACT**

The lipoxygenase (LOX) can influence the risk of cardiovascular disease (and diabetes) by virtue of its effect on lipid metabolism. Therefore, the present study deals with the separation and estimation of the level of LOX in blood of males with cardiovascular disease (only /or in addition to diabetes).

The enzyme was separated and purified from the blood-cells homogenate by precipitation (using 0-55% ammonium sulfate), dialysing against the buffer solution and then by applying ion exchange chromatography, when three bands were distinguished. The enzyme showed different specific activities in the elution of these bands and the highest activity was for the second band. Concerning the characterization of enzyme, it has been found that the optimum condition of the enzyme was at pH 8, 40°C and the enzyme has km=1.99 mM at 12mM of the substrate (linoleate) concentration.

The results showed that the incidence of cardiovascular disease and diabetes led to a significant increase in the level of LOX in both the blood-cells homogenate and the serum.

The study also focused on assessing the serum total cholesterol (TC), high density lipoprotein–cholesterol (HDL-C), glutathione (GSH) and malondialdehyde (MDA). The results indicated that there was a significant increase in the level of TC and MDA and there was a significant decrease in the level of HDL-C and GSH in addition to an increase in the ratio of TC/HDL-C.

On the basis of the results, it may be concluded that the LOX in blood cells has more than one isoenzyme with different activities. In addition, the high level of LOX is considered as a negative indicator that promotes cardiovascular disease and diabetes.

**Keywords**: lipoxygenase, cardiovascular disease, diabetes.

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### INTRODUCTION

Metabolic syndrome, which includes a number of systemic clinical and biochemical processes (insulin resistance, abdominal obesity, hypertension and dyslipidemia) attracts the attention of endocrinologists and cardiologists. This is due to the large spread of this syndrome in the population and its effect on the development of cardiovascular disease and diabetes mellitus type 2 (Novgorodtseva *et al.*, 2011).

Leukotrienes (LTs) are potent lipid mediators of inflammatory responses and have been implicated in the pathophysiology of acute and chronic inflammatory diseases including asthma, arthritis, psoriasis and inflammatory bowel disease (Maskrey and O'Donell, 2008). LTs are one type of eicosanoids that are produced from lipid peroxidation of 20-carbon (eicosa-) polyunsaturated fatty acids. They are generated in biological systems by lipoxygenases, LOXs (Murphy and Gljon, 2007).

LOXs are a family member of non-heme iron-containing dioxygenase that catalyze the step of conversion of arachidonic acid to 12 and 15-hydroxy eicosatetraenoic acid (12-HETE and 15-HETE) (Radmark *et al.*, 2007). The 12/15 LOX was originally isolated from porcine leukocytes, but its tissue distribution is now known to be relatively wide, including blood vessels, brain, skin and kidneys. Several lines of evidence have suggested that 12/15-LOX may play an important role in the development of atherosclerosis, diabetes and neurodegenerative disease (Spanbrock *et al.*, 2003).

The LTs family is divided into two classes, the classical chemoattractant LTB<sub>4</sub> and spasmogenic cysteinyl LTs such as LTC<sub>4</sub>, LTD<sub>4</sub> and LTE<sub>4</sub>. The LTs are involved in chronic inflammatory disease, including atherosclerosis, myocardial infarction and stroke (Chabane *et al.*, 2009). The peptidyl LT and LTB<sub>4</sub> increase the adhesion of leukocytes to endothelial cell (Maskrey and O'Donell, 2008). The LTs are predominantly synthesized by inflammatory cells, such as polymorphonuclear leukocytes, macrophages and mast cells (Huo *et al.*, 2004).

In addition to lipoxygenase, other variables have been found to be useful for detecting cardiovascular disease and diabetes. Both produce disturbances of lipid profiles, especially an elevation in the level of LDL-cholesterol and a decrease in the level of HDL-cholesterol, the major risk factors of coronary artery disease (Riendeau *et al.*, 1991).

Diabetes and cardiovascular disease increase susceptibility to lipid peroxidation. An increased oxidative stress has been observed as indicated by high free radical production. The measuring of serum antioxidants and malondialdehyde (the end product of lipid peroxidation) would provide an index of oxidative stress (Willams and Harwood, 2000). One of the most important endogenous antioxidants that counter the effect of free radicals is glutathione. It plays the role of a sulfhydryl group (SH) provider for direct scavenging reactions (Kayama *et al.*, 2009).

The present work aims to separate the lipoxygenase from the blood cells-homogenate. It also aims to investigate the relation between the level of lipoxygenase, total cholesterol (TC), high density lipoprotein-cholesterol (HDL-C), malondialdehyde (MDA) and glutathione (GSH) in the serum (in addition to the level of lipoxygenase in blood cells-homogenate) and the

probability of causing vascular disease and diabetes taking into consideration the age of the subjects and whether or not they are smokers.

### MATERIALS AND METHODS

The subjects were divided into two main groups: a healthy group (72 males) and a patient group (83 males). Of the patient group, 55 subjects had cardiovascular disease only and 28 subjects had diabetes millitus in addition to cardiovascular disease. All patients were residents in the Care Unit in Ibn Seena Teaching Hospital, Mosul. Then, each group is divided into three age categories (20-34 35-49 and  $\geq$  50 y ) taking in to consideration whether/ or not the subjects were smokers.

# **Blood sampling**

At the approval of Hospital management, blood samples were drawn from all subjects into plain plastic tubes and the serum was separated 2h after venipuncture by centrifugation.

# Separation of lipoxygenase from blood-cells homogenate

The precipitated blood-cells produced after centrifugation of blood samples were washed three times with isotonic saline and haemolysed in twice the volume of distilled water. The haemolysate was cautiously adjusted to pH6 by adding 1M HCl. The stroma was removed by centrifugation at 2000×g for 20 min. In the preparative experiments, 5mM mercaptoethanol was added in order to stabilize the enzyme. Ammonium sulfate was added to the supernatant and the precipitate, produced at 55% saturation, dissolved in water and then dialyzed against 0.003M Tris-HCl at pH7.2 using the semipermeable membrane, cellophane (Rapoport *et al.*, 1979).

Further enrichment was achieved by anion-exchange chromatography on DEAE-cellulose, which removed the residul haemogllobin (Lowenstein, 1981). A column with 40×2.5 cm diameters, supplied from Pharmacia Fine Chemicals, was used.

# Characterization of lipoxygenase

In order to find the optimum pH, the activity of the purified lipoxygenase was determined by using Tris-HCl buffers with a range of pH value (5-10), and in order to find the optimum temperature, the activity of the purified enzyme was determined by using a range of different temperatures  $(4-90^{\circ}\,\text{C})$ .

Michael's constant, km, of lipoxygenase was determined by carrying out the reaction using substrate (linoleate) solutions with different concentrations ranging between 0-14 mM.

### **Assay method**

The activity of lipoxygenase to convert linoleate to its hydroperoxide was assayed by Holman method (Rapoport *et al.*, 1979).

The protein level in the studied solutions (accept the serum ) was assayed by a modified Lowary method (Scharcterle and Pollack, 1973).

# **Estimation of biochemical parameters**

The serum total cholesterol (TC) and high density lipoprotein—cholesterol (HDL-C) were analyzed enzymatically using a commercial reagent (Kits) obtained from Fortress, U.K.

Al-Zamely *et al*'s modified method (2001) was used to estimate the concentration of serum glutathione which depends on the reduction of disulfide bond of 5, 5'-dithio-bis-2-nitrobenzoic acid (DTNB) by the sulfhydryl group of glutathione.

The formation of malondialdehyde (MDA) was determined by the thiobarbituric acid method (Akande and Akinyinka, 2005).

## **Statistical analysis**

For comparison between more than two groups, Duncan multiple range test was used. All the data were expressed as mean  $\pm$  S.D. of the mean. Different letters indicate a significant difference at p  $\leq$  0.05 level.

### **RESULTS AND DISCUSSION**

# Separation of lipoxygenase from blood-cells homogenate

As stated before, several experiments have been applied to separate lipoxygenase from blood-cells homogenate. The separation methods used are based on the differences in the molecular size, charge and hydrophobic affinity. After each experiment, the total protein, total activity and specific activity were determined. The specific activity was taken into consideration.

As shown in table 1, the specific activity of lipoxygenase in the initial homogenate was 0.854 U/mg protein. After precipitation by ammonium sulfate, it was raised to 1.253 U/mg protein, i.e. it was multiplied by 1.468 folds compared with that of initial homogenate. Dialysis increased the specific activity of the enzyme up to 1.973 U/mg protein or it was multiplied by 2.31 folds.

The ion exchange chromatography was applied to the residue obtained from dialysis. During the elution with 0.03M Tris-HCl at pH 7.2, three well defined-lipoxygenase bands have been appeared (Fig.1) and they showed different specific activities; 2.949, 5.733 and 3.435 U/mg protein for band I, II and III respectively.

Table 1: Separation of lipoxygenase from blood-cells homogenate

Fraction	Volume (ml)	Total protein (mg)	Total activity (units)	Specific activity (units/mg protein)	Purification fold	Yields %
Blood-cells Homogenate	15	9.435	8.055	0.854	1	100
Precipitation by (0-55%) ammonium sulfate	12	7.2	9.024	1.253	1.468	112.0298
Dialysis	13.5	5.4	10.6515	1.973	2.310	132.2346
Ion-exchange (DEAE-Cellulose)						
band I	60	2.865	8.448	2.949	3.454	104.879
band II	40	0.72	4.128	5.733	6.716	71.95531
band III	45	1.6875	5.796	3.435	4.023	51.24767

<sup>\*</sup> U, the amount of enzyme that produce 1  $\mu M$  of product per minute.

<sup>\*\*</sup> U/mg, the number of enzyme units per milligram of protein.

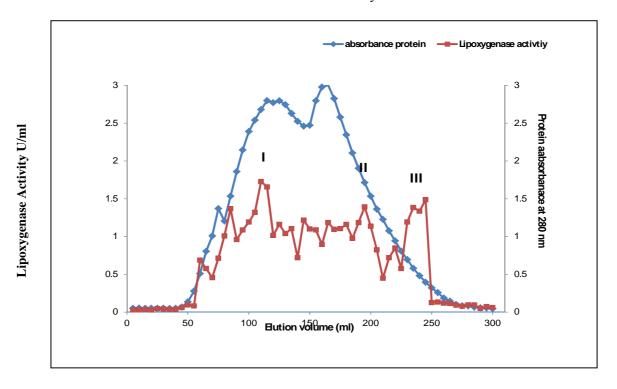


Fig. 1: Elution profile of protein and lipoxygenase activity from the purification of the enzyme on DEAE-cellulose by applying ion-exchange chromatography.

It was stated that the lipoxygenase of human blood-cells was soluble enzyme. However, when subjected to centrifugation at  $100000 \times g$  for 1 h, only 30-60% of the initial activity could be recovered in the supernatant. In addition, 40-60% of the enzymatic activity in the purified preparation was lost upon standing in the chromatography buffer at 0°C for 30 min (Rouzer and Samuelsson, 1985). This indicates that the purification of the human lipoxygenase would be extremely difficult.

On the other hand, maximal enzymatic activity is clearly dependent on the presence of stimulating factors such as Ca<sup>2+</sup> and ATP. Also, glutathione and mercaptoethanol protect lipoxygenase activity and the later is much more effective (Rapoport *et al.*, 1979), hence it was added to the blood-cells homogenate to achieve this purpose.

As shown in Table (1), the specific activity of lipoxygenase in the blood-cells homogenate has been increased by 1.468 and 2.310 folds after separation by ammonium sulphate precipitation and dialysis, respectively comparing with its activity in the initial homogenate.

More than one band has been obtained in case of applying ion-exchange chromatography (Fig. 1) with different activities suggesting microheterogeneity (Rouzer and Samuelsson, 1985). The second band exhibited the higher activity, it increased by 6.716 folds. This indicates the presence of more than one isoenzyme and the more active form, toward linoleate at pH9 and 37°C, has been presented in the second band-elution produced by applying ion exchange chromatography using DEAE-cellulose.

## Characterization of lipoxygenase

This study has dealt with the characterization of lipoxygenase appearing in the elution of band II produced from ion-exchange chromatography.

The effect of pH (using Tris-HCl buffer) on the activity of lipoxygenase is shown in Fig 2. The top of the curve shows that the optimum pH of the enzyme activity was to be at pH8, with the activity falling off symmetrically on both sides.

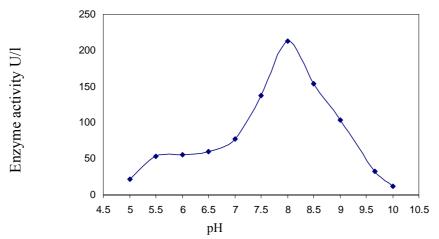


Fig.2: Effect of pH on lipoxygenase activity using Tris-HCl buffer.

The catalytic mechanism of the enzymes is largely due to the fact that its active sites possess structures which are uniquely suited to promote catalysis. Several factors contribute to enzyme catalysis and one of the most important of these factors is the pH which plays on important role in the formation, maintenance or disturbance of the three-dimensional structure of the enzyme molecule or its active site (McKee and McKee, 1996). Each enzyme has its own optimum pH to reach the maximum velocity and the optimum pH is not necessarily the best pH for physiological control *in vivo* (Christenson *et al.*, 2001).

Figure 3 exhibited the effect of temperature on the activity of lipoxygenase. The top of the curve shows that the optimum temperature of the enzyme activity was to be at 40°C, with the activity falling off symmetrically on both sides.

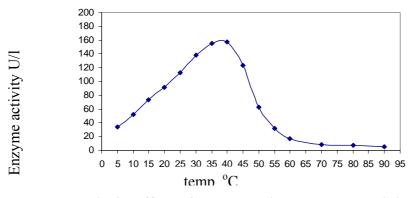


Fig.3: Effect of temp. on lipoxygenase activity.

Increasing the temperature results in an increase in enzyme activity until the optimum temperature is reached, above which any increase in temperature results in a decrease in enzyme activity. The sudden loss of enzyme activity that occurs at temperatures above the optimum temperature is due to thermal denaturation. Most enzymes are inactivated at 55-60 °C (Christenson *et al.*, 2001).

This study has also dealt with the estimation of lipoxygenase affinity toward linoleate through the determination of km of its reaction. As shown in Fig. 4, the highest activity was encountered at 12mM substrate (linoleate) concentration, so Km was 1.8mM (applying Michael-Menten equation) and 1.99 mM (applying Lineweaver-Berk equation).

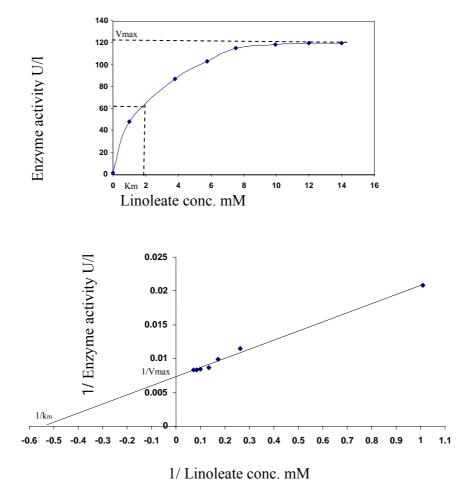


Fig. 4: Effect of substrate conc. on lipoxygenase activity.

This result indicated that lipoxygenase has a high affinity toward linoleate, since the affinity is inversely proportional to km-value and it is worth-noting that the km-values of most of the enzymes are ranged between  $10^{-1} - 10^{-6}$  M (Al-Flayeh, 2000).

# Estimation of lipoxygenase level in serum and blood-cell homogenate

The level of lipoxygenase in the serum and in the blood-cells homogenate of studied subjects are presented in Table (2) (I-VI). Compared with the healthy subjects, all subjects with cardiovascular disease, whether smokers or nonsmokers, exhibited a significant increase in the level of lipoxygenase in the serum and in the blood-cells homogenate. This result agrees with what Imig (2006) and Jawien and Korbut (2010) conclude; they state that the LOX is abundantly expressed in atherosclerosis lesions. The high expression of this enzyme in the lesions of animal atherosclerosis and humans, and its absence in normal vessel wall suggest its involvement in atherosclerosis (Imig, 2006; Kuhn *et al.*, 1992).

There are numerous potential targets that could be useful in the intervention in LT-metabolism in atherosclerosis. The microsomal protein, five lipoxygenase activating protein (FLAP), was found to be critical for regulation of 5-LOX activity and biosynthesis of LTs within certain compartments of plasma membrane. The genetic polymorphisms of FLAP are associated with myocardial infarction and stroke by increasing LT production and inflammation in the arterial wall (Helgadottir *et al.*, 2004). In advanced human atherosclerotic lesions, the hydroxy linoleic acid/ linoleic acid ratio is much higher and may be used as a reliable parameter to quantify the degree of oxidative modification which reflects the level of LOXs (Kuhn *et al.*, 1992).

Table 2: The level of the lipoxygenase and some biochemical variables in male blood.

I Nonsmokers aged 20-34 y				
	Mean ± S.D.			
Parameters	Control n=11	Cardiovascular Patients n=6	Diabetic Patients n=0(not obtained)	
LOX level in the serum***	188.5 ± 13.03	286.50 ± 8.66		
U/l	B	A		
LOX level in the blood-	300 ± 21.5	516.0±28.3		
cells homogenate*** U/l	B	A		
TC level***	140.0±21.8	161.77±6.85		
mg/100 ml	B	A		
HDL-C level*	58.159 ± 4.524	50.6±7.1		
mg/100 ml	A	B		
TC/HDL-C*	3.06±0.85 B	3.90±0.62 A		
GSH level***	16.93 ± 7.34	7.88 ± 2.46		
μ Μ/Ι	A	B		
MDA level*	2.31 ± 0.71	3.13±0.60		
μ Μ/Ι	B	A		

II Nonsmokers aged 35-49 y					
	Mean ± S.D.				
Parameters	Control n=10	Cardiovascular Patients n=6	Diabetic Patients n=8		
LOX level in the serum*** U/l	$231.3 \pm 57.8$	367.2 ± 78.9 B	533.5 ± 79.3 A		
LOX level in the blood-cells homogenate*** U/l	541±137	975.2±19.4	1272 ± 257		
	C	B	A		
TC level***	166.99 ± 70.05	213.53 ± 18.47	228.75 ± 15.5		
mg/100 ml	C	B	A		
HDL-C level***	51.657±4.990	40.58 ± 6.79	34.47 ± 7.07		
mg/100 ml	A	B	C		
TC/HDL-C*	3.82±1.01	4.55 ± 1.30	4.93 ± 1.66		
	B	A	A		
GSH level***	14.71 ± 3.68	6.33 ± 2.76	5.7300±1.83		
μ M/I	A	B	B		
MDA level*	2.88 ± 1.061	3.71 ± 2.39	4.1±1.29		
μ M/l	B	A	A		

III Nonsmokers aged ≥ 50 y					
	Mean ± S.D.				
Parameters	Control n=14	Cardiovascular Patients n=11	Diabetic Patients n=8		
LOX level in the serum** U/l	268.5 ± 73.9	477±107	706±154		
	C	B	A		
LOX level in the blood-cells homogenate*** U/l	580.40±170 C	1304±218 B	1756±280.4 A		
TC level**	179.3 ± 24.91	221.5±33.23	248.01 ± 45.3		
mg/100 ml	C	B	A		
HDL-C level**	46.080±3.779	38.1 ± 2.20	33.8±2.8		
mg/100 ml	A	B	B		
TC/HDL-C*	$4.073 \pm 0.911$ B	5.15 ± 1.89 A	5.62±1.48 A		
GSH level**	13.40 ± 2.42	5.63 ± 1.47	4.646±0.719		
µ M/l	A	B	B		
MDA level*	3.66±0.55	3.89±0.36	4.37±0.73		
μ M/l	B	B	A		

IV smokers aged 20-34 y				
	Mean ± S.D.			
Parameters	Control n=9	Cardiovascular Patients n=8	Diabetic Patients n=0(not obtained)	
LOX level in the serum** U/l	$226.3 \pm 64.4$ B	423 ± 49.8 A	<del></del>	
LOX level in the blood-	480 ± 126.7	844.0 ± 139.6		
cells homogenate*** U/l	B	A		
TC level**	151.7±20.5	180.4±15.9		
mg/100 ml	B	A		
HDL-C level**	50.12 ± 6.72	40.6±8.1		
mg/100 ml	A	B		
TC/HDL-C*	3.60 ± 1.25 B	4.93 ± 1.124 A	_	
GSH level***	16.35 ± 2.07	6.490 ± 0.910	_	
µ M/l	A	B		
MDA level*	2.88 ± 0.23	3.26±0.42		
μ M/l	B	A		

	V smokers aged 34-49 y				
	Mean ± S.D.				
Parameters	Control n=11	Cardiovascular Patients n=8	Diabetic Patients n=0(not obtained)		
LOX level in the	$270 \pm 25.6$	$543 \pm 39.5$			
serum** U/l	В	A			
LOX level in the					
blood-cells	$567 \pm 140.9$	1452 ± 191			
homogenate***	В	A			
U/l					
TC level**	$183.3 \pm 17.9$	$230 \pm 22$			
mg/100 ml	В	A			
HDL-C level*	$39.349 \pm 4.075$	$32.5 \pm 7.8$			
mg/100 ml	A	В			
TC/HDL-C*	$4.28 \pm 0.45$	$5.26 \pm 0.95$			
TC/HDL-C	В	A			
GSH level**	$14.300 \pm 2.141$	$5.70 \pm 1.81$			
μ M/l	A	В			
MDA level*	$3.4650 \pm 0.466$	$3.97 \pm 0.495$			
μ Μ/Ι	В	A			

VI smokers aged $\geq 50~\mathrm{y}$					
	Mean ± S.D.				
Parameters	Control n=17	Cardiovascular Patients n=16	Diabetic Patients n=12		
LOX level in the serum** U/l	494±168.3	711 ± 201	907±342		
	C	B	A		
LOX level in the blood-cells homogenate* U/l	1107.0±375.6 B	1722 ± 503 A	1976±559 A		
TC level***	187.589 ± 44	247.6±68	257±72		
mg/100 ml	B	A	A		
HDL-C level*	36.47 ± 6.03	31.6±7.5	30.3 ± 8.6		
mg/100 ml	A	B	B		
TC/HDL-C*	4.86±0.40	5.52±2.25	5.81 ± 2.88		
	B	A	A		
GSH level***	13.270 ± 2.07	5.48±1.06	4.40±1.08		
µ M/l	A	B	B		
MDA level*	3.8505 ± 1.05	4.09 ± 1.13	4.74±1.27		
μ M/l	B	A	A		

- Abbreviations: LOX, lipoxygenase; TC, total cholesterol; HDL-C, high density lipoprotein-cholesterol; GSH, glutathaione; MDA, malondialdehyde.
- Different letters horizontally mean a significant difference at 0.05 level according to Duncan test.
- \*,\*\* and \*\*\* means that P-values are > 0.05, >0.01 and >0.001, respectively.

As shown in Table 2, the same pattern was obtained for the patients with diabetes, as well as cardiovascular disease, for all of the studied groups accept the smokers and nonsmokers groups aged 20-34 y and the smoker group aged 35-49 y because samples for these groups of patients could not be obtained. They exhibited a significant increase in the level of LOX in the serum and in the blood-cells homogenate. This result agrees with what was stated earlier; i.e. the increased expression of LOXs and then LTE<sub>4</sub>-generation were associated with diabetes. Streptozotocin-induced diabetic rats had increased urinary excretion of LTE<sub>4</sub>, then its level was decreased by insulin treatment (Kang *et al.*, 2001). High glucose level also increased the LOX enzymes in cultured endothelial cells, VSMC, mesangial cells and glomeruli of diabetic rats (Natarajan and Nadler, 2004; Kang *et al.*, 2001). Interestingly, one of a few genes that are increased in pancreatic  $\beta$ -cells of the insulin-resistant diabetic rates is that for LOXs (Natarajan and Nadler, 2004). In addition, Kang *et al.* (2001) state that the 12/15 LOX product, 12-HETE, is increased in the urine of diabetic patients, which reflects the relation between the level of LOXs and diabetic mellitus.

#### The Biochemical Parameters

#### **Total cholesterol**

As shown in Table 2, there has been a significant increase in the level of total cholesterol, TC, in the serum of patients compared with healthy subjects. This result agrees with many other previous studies (WHO report, 1994). Purushothaman  $et\ al.\ (2011)$  state that peroxisome proliferator-activated receptor alpha (PPAR  $\alpha$ ) regulates genes involved in lipid metabolism, homeostasis and inflammation, in response to fatty acids and fibrates, making it a candidate gene for risk of dyslipidaemia, atherosclerosis and coronary artery disease. Various polymorphic markers that act as candidate genes are related to lipoprotein metabolism and account for variations in the level of serum TC and LDL-C (Ozturk and Killen, 1999).

#### **HDL-Cholesterol**

Table 2 exhibited a significant decrease in HDL-C in the serum of males with cardiovascular disease (with or without diabetes), whether smokers or nonsmokers.

Although the relationship between plasma HDL-C and the development of atherosclerosis is complex, it has been established that the level of plasma HDL-C is inversely proportional to the risk factor of this disease (Chapman, 2004). This establishment agrees with that obtained in the present study. It has been stated that the risk of coronary heart disease, CHD, is elevated by 3% in women and 2% in men for each 1 mg/dL decrement in the level of HDL-C than its normal level which is 44 mg/dL (Gordon *et al.*, 1989). Nevertheless, high levels of HDL-C (> 65 mg/dL) were shown to offer protection against CHD events even when LDL-C level was elevated above 160 mg/dL (Castelli *et al.*, 2011).

Pirillo *et al.* (2008) stated that HDL<sub>3</sub> can be modified in vitro by 15-LOX generating a dysfunctional lipoprotein that exhibits a decrease ability to promote cholesterol efflux from lipid-laden macrophages. This modification significantly impairs the ability of HDL<sub>3</sub> to function as an efficient cholesterol acceptor or leads to decreasing HDL-C. The modification of HDL<sub>3</sub> (the major antiatherogenic HDL subfraction) by 15-LOX, an enzyme overexpressed in the atherosclerotic lesions, impairs the anti-inflammatory activity of this lipoprotein, and the modified HDL<sub>3</sub> failed to inhibit induction of adhesion molecules and chemoattractants which act as mediators of monocyte adhesion to endothelial cells. This establishment demonstrates that the protective role of HDL<sub>3</sub> against the initiation and progression of atherosclerosis of atherosclerosis is impaired (Pirillo *et al.*, 2008).

On the other hand, it was stated that the ratio of TC/HDL-C (but not their absolute value) reflects the real risk factor related to patients with cardiovascular disease; the higher the level of TC/HDL-C, the higher the susceptibility of cardiovascular events (Kumar *et al.*, 2009; Wang *et al.*, 2009; Stevens *et al.*, 2001). Therefore, the high level of this ratio for subjects with cardiovascular disease shown in the table seems to contribute to the development of this disease.

#### Glutathione

As shown in Table (2), the levels of glutathione in serum of patient subjects were significantly decreased. This decrease may be attributed to the increased oxidative stress and the generation of the free oxygen radicals (ROS). Many experimental reports demonstrate that ischemia followed by reperfusion elicit a cascade of proinflammatory reactions that are known to lead to the production of ROS (Pasupathi *et al.*, 2009; Raghuvanshi *et al.*, 2005). It is well recognized that diabetes induces oxidative stress and then the accumulation of ROS which, inturn, reduce the activities of antioxidant enzymes such as catalase and superoxide dismutase. The decrease in the activities of these enzymes result in insufficient availability of GSH (Raghavan and Kumari, 2006; Powell *et al.*, 2004). In addition, it was stated that long term exposure to high glucose decreases the expression of glutamyl cysteine ligase (the regulating enzyme for GSH synthesis) resulting in a decrease in GSH levels (Tran *et al.*, 2004).

# Malondialdehyde

Table (2) exhibited that the levels of serum MDA of patients were elevated significantly. The MDA is an important end product of lipid peroxidation, so, the increase in its level reflects the increase in free radicals (Janero, 1990). Free radicals are considered to have a great role in the pathogenesis of many diseases such as diabetes mellitus, atherosclerosis, cancer, myocardial infarction and immune diseases (Pasaoglu *et al.*, 2004). An increase in free radical formation enhances oxidation of LDL, an important mediator of vascular injury in cardiovascular pathologlogies including hypertension, atherosclerosis and ischemia-reperfusion (Khanna *et al.*, 2008). On the other hand, high level of glucose can increase lipid peroxidation. Hyperglycemia may stimulate autooxidtion glucose leading to free radical production in diabetics (Pasaoglu *et al.*, 2004).

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