Correlation between Lipid parameters and gamma glutamyl transferase in type 2 diabetes mellitus

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Abstract

Background: Serum gamma glutamyl transferase (GGT) which is a simple and reliable marker of liver fat deposition leads to hepatic insulin resistance and long term hepatic insulin resistance may lead to type 2 diabetes mellitus (DM). In this context, the present study was under taken to study the serum GGT levels and its possible associations with serum triglyceride (TG), total cholesterol (TC), low density lipoprotein cholesterol (LDL-C), very low density lipoprotein cholesterol (VLDL-C) and high density lipoprotein cholesterol (HDL-D) in type 2 DM.

Material and methods: The study was conducted at HSK hospital, Bagalkot from Jan 2016 to Jun 2016. The diagnosis of DM was based on WHO criteria. Sixty subjects participated in each group (DM and Controls). Fasting sample was collected and fasting blood sugar (FBS), HbA1c, creatinine, GGT, and lipid parameters were estimated. Data was analysed using SPSS software. Student’s t test and Pearson correlation was applied for quantitative data.

Results: The serum levels of FBS, PPBS, HbA1c, creatinine, urea, TG and VLDL were raised significantly in cases compared to controls, whereas serum HDL was significantly decreased in cases compared to controls (p=0.000). There was statistically significant (p=0.001) positive correlation between the TGL and GGT. HDL-C was significantly, negatively correlated with GGT.

Conclusion: There was positive correlation between GTG and TG, TC, LDL-C and VLDL-C, only TGL was significant, and there was significant negative correlation between HDL and GGT. Hence serum GGT can be used as marker of cardiovascular diseases in type 2 DM patients.

Key words: Diabetes mellitus, Gamma Glutamyl transferase, Lipid profile

Introduction

Diabetes mellitus (DM) is a major metabolic and non-communicable disease, its prevalence is increasing exponentially. World-wide, type 2 DM is constitutes for about 90% of all the cases and is more in men than women[1-2]. The prevalence of DM worldwide was 2.8% in 2000 and is estimated to rise to 4.4% in 2030. The total number of DM is projected to 592 million by 2035. India presently has more than 50 million diabetic patients, making India “the diabetes capital of the world”[3,4].

Dyslipidemia is one of the major risk factor for coronary heart disease[5]. The variations of serum lipid profile from their normal levels can predict the coronary artery disease[6,7]. Hyperglycaemia and atherosclerosis are related in type-2 DM[8]. Patients with type-2 DM have increased risk of cardiovascular disease associated with atherogenic dyslipidaemia[9]. Persistent hyperglycaemia causes glycosylation of proteins, mainly collagen cross linking and matrix proteins of arterial wall. This eventually leads to endothelial cell dysfunction and further to atherosclerosis[10]. The cardiovascular disease is a cause of morbidity and mortality, commonly presents as an abnormally high level of triglycerides (TG), a high proportion of small dense low lipoprotein cholesterol (LDL-C), low high density lipoprotein cholesterol (HDL-C), and postprandial lipemia[11,12,13]. This pattern of lipid profile in DM type 2 is termed diabetic dyslipidemia[14].

Gamma glutamyltransferase (GGT) catalobises extracellular glutathione[15]. It is present, in the liver, bile duct, lungs, pancreas, brain, gall bladder, kidneys and the heart muscles. In the diseases of all these organs, there will be significant increase in serum GGT levels[16,17]. The GGT levels can also be raised in patients who are for a long duration on drugs like phenytoin, barbiturates, amiodarone, tamoxifen and steroids and also who have undergone biliopancreatic surgeries[18]. Although GGT is produced in all tissues, differences in the sugar moieties allow that only the liver GGT is detectable in serum[19]. The studies have demonstrated that abnormal hepatocellular function is associated with obesity, insulin resistance, and type 2 diabetes[20-22]. The loss of a direct effect of insulin to
suppress hepatic glucose production and glycogenolysis in the liver causes an increase in hepatic glucose production\cite{20,23}.

Raised liver enzymes reflect chronic ectopic fat deposition\cite{24}. Serum GGT may be a simple and reliable marker of hepatic fat deposition and hepatic steatosis which can lead to hepatic insulin resistance and long term hepatic insulin resistance may lead to type 2 DM\cite{25}. Only few studies have compared the serum GGT with lipid profile, hence, in this context, the present study was under taken to study the serum GGT levels and its possible associations with TG, TC, LDL-C, and HDL-C in type 2 DM.

**Material and Methods**

The study was conducted at HSK hospital, Bagalkot from Jan 2016 to Jun 2016. The study was approved by institutional ethics committee. Informed consent was obtained from all the participants. The diagnosis of DM was based on WHO criteria. Sixty subjects participated in each group (DM and controls). Alcoholics, smokers, patients with diabetic complications, chronic liver diseases, HIV, severe anemia, CRF, other systemic conditions and patients on iron preparations, lipid lowering drugs were excluded. Pregnant women were also excluded from the study. Under aseptic precautions 5 ml of fasting sample was collected and following biochemical parameters were estimated. FBS and PPBS were estimated by GOD-POD method, serum urea by DAM method, creatinine by Jaffe’s method. TG was measured by CHOD-PAP method\cite{25-29}. Kits were supplied by Biosystems Pvt Ltd, GGT estimated by IFCC method kit supplied by Transasia (Normal 7-50 IU/L), Biosystems A 25 fully automated instrument was used for estimation of above mentioned parameters. LDL-C and VLDL-C were calculated using Fried Wald’s formula\cite{30,31}. HbA1c was measured by using NycoCard reader. Post prandial blood sugar (PPBS) was also estimated.

**Statistical analysis**

Data was tabulated in excel, analysed using SPSS window version; SPSS, 11.5 Inc, Chicago IL. Student’s t test and Pearson correlation was applied for quantitative data. All the values were expressed in mean±SD.

**Results**

In the present study, the mean age of cases and controls was 46.3±10.0 and 49.4±12.7 years respectively, there was no statistically difference between two groups (p=0.429).

The serum levels of FBS, PPBS, HbA1c, creatinine, urea, TG and VLDL-C were raised significantly in cases compared to controls, whereas serum HDL-C was significantly decreased in cases compared to controls (p=0.000). The serum concentration of TC and LDL-C was increased in cases compared to controls, but it was not statistically significant (p value was 0.386 and 0.294 respectively) (Table 1).

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Controls</th>
<th>Cases</th>
<th>p value</th>
</tr>
</thead>
<tbody>
<tr>
<td>FBS mg/dl</td>
<td>86.1±10.3</td>
<td>171.7±38.2</td>
<td>0.000</td>
</tr>
<tr>
<td>PPBS mg/dl</td>
<td>115.0±17.6</td>
<td>274.7±78.4</td>
<td>0.000</td>
</tr>
<tr>
<td>GGT mg/dl</td>
<td>19.0±5.1</td>
<td>42.1±15.4</td>
<td>0.000</td>
</tr>
<tr>
<td>HbA1c %</td>
<td>5.1±0.7</td>
<td>8.0±1.2</td>
<td>0.000</td>
</tr>
<tr>
<td>Urea mg/dl</td>
<td>18.8±5.7</td>
<td>32.7±9.3</td>
<td>0.000</td>
</tr>
<tr>
<td>Creatinine mg/dl</td>
<td>0.9±0.1</td>
<td>1.2±0.2</td>
<td>0.000</td>
</tr>
<tr>
<td>TG mg/dl</td>
<td>116.3±31.7</td>
<td>153.1±33.3</td>
<td>0.002</td>
</tr>
<tr>
<td>TC mg/dl</td>
<td>183.8±36.0</td>
<td>194.5±37.5</td>
<td>0.386</td>
</tr>
<tr>
<td>HDL-C mg/dl</td>
<td>49.7±4.7</td>
<td>33.4±3.7</td>
<td>0.000</td>
</tr>
<tr>
<td>VLDL-C mg/dl</td>
<td>23.2±6.3</td>
<td>30.2±6.5</td>
<td>0.002</td>
</tr>
<tr>
<td>LDL-C mg/dl</td>
<td>110.8±32.6</td>
<td>122.4±33.6</td>
<td>0.294</td>
</tr>
</tbody>
</table>

FBS: Fasting blood sugar  
PPBS: Post prandial blood sugar  
GGT: Gamma glutamyl transferase  
TG: Triglyceride  
TC: Total cholesterol  
HDL-C: High density lipoprotein cholesterol  
VLDL-C: Very low density lipoprotein cholesterol  
LDL-C: Low density lipoprotein cholesterol
There was statistically significant (p=0.001) positive correlation between TG and GGT the correlation coefficient was 0.56. There was positive correlation between TC, VLDL-C and LDL-C, but it was not statistically significant. HDL-C was significantly, negatively correlated with GGT.

Discussion

In the present study, serum GGT was significantly higher in DM patients compared to age and sex matched healthy control group, this finding is in accordance with previous studies[32, 33]. Studies in humans and animal models have showed that serum GGT levels were significantly elevated in diabetes[33].

As we searched the literature, we could find only one original research article on DM that compared lipoproteins and GGT. But studies in other conditions like coronary artery disease, metabolic syndrome and obesity are available.

Study by Emirogulu MY et al.[34] on acute coronary syndrome found no correlation between GGT and Lipid profiles (r=0.166 p=0.03 for GGT and TC, r=0.12 p=0.043 for GGT and LDL-C) but Desai G M et al[6] showed TG, TC and LDL were positively correlated with GGT and only TG showed significant correlation(r=0.58, p=0.04). HDL was negatively correlated with GGT (r=-0.44, p=-0.30). The current study also showed similar results in type 2 DM.

The study by Khan S N et al[35] on metabolic syndrome subjects showed TG was significantly correlated with GGT (r=0.91, p= 0.02) and HDL showed significant negative correlation with GGT(r=-0.192, p=0.018), but study on cardiac syndrome X in Turkey by Demir B et al[36] showed that TC and LDL were negatively correlated with GGT and were not statistically significant, HDL and TG were positively correlated with GGT but only TGL was statistically significant correlation(r=0.293, p=0.039). P. Josephine Latha et al[37] in their study on obese individuals showed that there was positive correlation between GGT and TC (r=0.72), TGL (r=0.662), LDL (r=0.669), VLDL (r=0.662) which were highly significant positively. HDL was negatively correlated(r = -0.773) with GGT and was found to be highly significant.

Rajarajeswari D et al[20] observed a significant correlation between GGT and lipoprotein levels. Positive association between serum GGT and triglycerides (r = 0.112), serum GGT and LDL lipoprotein (r = 0.05), serum GGT and cholesterol(r=0.027) were observed. In the current study, it was found that TGL was positively correlated with GGT, but other lipoproteins were not significantly correlated with GGT. We also observed a significant negative correlation between serum GGT and HDL lipoprotein (r = -0.298), which was similar to the results of the study by Rajarajeswari D et al.

Lipid triad[38] in DM comprises of hypertriglyceridemia, low levels of HDL-C and the appearance of small, dense, LDL (sdLDL)[39, 40]. The present study also revealed similar results. Many studies demonstrated that in type 2 DM, there is increased reactive oxygen species production and induce higher oxidative damage in the circulation and also have reduced antioxidant defenses mechanisms[41].

Association of GGT with lipids and lipoproteins could be explained by the antioxidant property of GGT[21, 42]. Elevated GGT could reflect subclinical inflammation, which represents the underlying oxidative stress. GGT levels are closely related to oxidative stress because cellular GGT has a central role in glutathione homeostasis by initiating the breakdown of extracellular glutathione, a critical antioxidant defence for the cell[43, 44].

The limitations of the present study were small sample size and diet history was not considered in detail. Hence further large sample size with detailed diet history is required.

In conclusion in the current study there was a positive correlation between GGT and TG, TC, LDL and VLDL, however only the correlation between GGT and TG was significant, and there was significant negative correlation between HDL and GGT. Hence, serum GGT can be used as a marker of cardiovascular disease in type 2 DM patients.

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