Detection of Serum Biochemical Markers in Elderly Patients with Type 2 Diabetes Mellitus and Its Relationship with Coronary Heart Disease

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Received 2020 June 27; Revised 2020 July 11; Accepted 2020 July 29.

Abstract

Background: Coronary heart disease (CHD) is myocardial ischemia caused by coronary atherosclerosis and stenosis.

Objectives: This study aimed to investigate the relationship of CHD with glycated albumin (GA), glycosylated hemoglobin (HbA 1c), glucose (Glu), homocysteine (Hcy), triglyceride (TG), and apolipoprotein A (apoA) levels in elderly patients with type 2 diabetes mellitus (T2DM). Moreover, it was attempted to predict which changes had a significant correlation with the occurrence of CHD in these biochemical indicators.

Methods: This clinical cohort study included a total of 472 patients admitted to our hospital from June 2018 to June 2019. They were then divided into three groups of concurrent (n=168, T2DM complicated with CHD), DM (n=148, T2DM alone), and control (n=156, healthy individuals).

Results: There were significant differences between the disease group (i.e., DM and concurrent groups) and control group in terms of GA (P=0.013, 28.58±8.01 vs. 15.77±1.44), HbA 1c (P=0.022, 9.5±1.5 vs. 5.5±0.5), Glu (P=0.012, 8.54±2.23 vs. 4.12±0.39), Hcy (P=0.031, 11.16±3.28 vs. 5.03±2.87), TG (P=0.021, 1.83±0.49 vs. 0.84±0.18), and apoA (P=0.031, 1.10±0.12 vs. 1.30±0.18). Moreover, GA (P=0.025, 27.14±6.34 vs. 28.58±6.01; concurrent group vs. DM group), HbA 1c (P=0.033, 8.3±1.2 vs. 9.5±1.5; concurrent group vs. DM group), Glu (P=0.019, 8.62±3.56 vs. 8.54±2.23; concurrent group vs. DM group), Hcy (P=0.031, 17.56±6.36 vs. 11.16±3.28; concurrent group vs. DM group), total cholesterol (TTC) (P=0.022, 3.06±0.20 vs. 3.69±0.29; concurrent group vs. DM group), and low-density lipoprotein (LDL) (P=0.037, 2.57±1.02 vs. 3.40±1.17; concurrent group vs. DM group) were higher than those in the control group; however, apoA (P=0.023, 0.95±0.12 vs. 1.10±0.12; concurrent group vs. DM group) in disease group was lower than that in the control group. Furthermore, there were significant differences between the disease and control groups in terms of GA, HbA 1c, Glu, Hcy, TC, LDL, and apoA (P<0.05). Pearson correlation analysis between DM and concurrent groups was performed on the clinical parameters with statistical differences, and GA was highly correlated with HbA 1c and Glu (P<0.01).

Conclusion: High-risk patient screening with high levels of GA, Hcy, and apoA in elderly patients with T2DM can not only improve the symptoms of patients with targeted treatment but also reduce the incidence of CHD by timely intervention, which is of great significance to improve the quality of life of patients.

Keywords: Apolipoprotein A, Coronary heart disease, Glycated albumin, Homocysteine, Type 2 diabetes mellitus

1. Background

Coronary heart disease (CHD) is myocardial ischemia caused by coronary atherosclerosis and stenosis. It is a cardiovascular disease that seriously threatens the health of human life (1). Its incidence has increased year by year and is developing in younger adults. About 70% of patients with CHD have abnormal blood glucose. Type 2 diabetes mellitus (T2DM) is considered an independent risk factor for cardiovascular disease. In patients with T2DM, coronary atherosclerotic heart disease has an early onset, high disease rate, and severe coronary lesions. The risk of stroke and cardiovascular death in the T2DM population is nearly 4 times higher than that in the non-T2DM population. In addition to the high-risk factors (i.e., dyslipidemia, hypertension, smoking, age, obesity, hyperhomocysteinemia) of the disease, T2DM is also a risk factor for the development of CHD (2). The incidence of T2DM patients complicated with CHD can be as high as 72.3%, which is 2 to 4 times that of non-diabetic patients (3). Long-term effective blood glucose monitoring has an important clinical guiding value for reducing T2DM complications (4). According to the Chinese Guidelines for Clinical Application of Blood Glucose Monitoring, the determination of glycated albumin (GA) has become one of the effective methods for blood glucose monitoring in patients with T2DM, and its evaluation of diabetic vascular complications has also important clinical value (5). However, there is little data on controlling blood glucose in elderly patients with T2DM. Furthermore, there is no conclusion about the GA risk factors associated with T2DM in China (6).

2. Objectives

This study aimed to investigate the relationship of CHD with GA, glycosylated hemoglobin (HbA 1c), glucose (Glu), homocysteine (Hcy), triglyceride (TG), and apolipoprotein A (apoA) levels in elderly patients with T2DM. Moreover, it was attempted to predict which changes in these biochemical
indicators had a significant correlation with the occurrence of CHD.

3. Methods

This clinical cohort study included a total of 472 patients admitted to our hospital from June 2018 to June 2019. They were then divided into three groups of concurrent (n=168, T2DM complicated with CHD), DM (n=148, T2DM alone), and control (n=156, healthy individuals). Concurrent and DM groups collectively referred to as the disease group, and patients in the control group were selected from the physical examination center of our hospital.

Subsequently, the blood samples of the patients were taken on an empty stomach, and the serum was taken after centrifugation. The inclusion criteria were: 1) fulfillment of the diagnostic criteria for T2DM in the 2007 edition of Chinese Guidelines for Diabetes Prevention and Control, 2) normal cognition and clear mind, and 3) lack of serious external trauma recently. It should be mentioned that the patients in the control group met the diagnostic criteria for CHD after clinical diagnosis.

On the other hand, the patients who were allergic to the test equipment and drugs; those with hearing, comprehension, vision, and reading disorders; the cases who exited the research procedure during diagnosis and treatment; and the individuals with the combination of the severe immune blood system, nervous system or malignant tumor were excluded from the study.

The study protocol was approved by the Ethics Committee of Linyi Central Hospital, Shandong, China. It is worth mentioning that informed consent was obtained from the patients.

The control group included 156 healthy cases (78 males and 78 females) who referred to our Health Checkup Center for physical examinations with a mean age of 76.6±5.50 years (age range: 70-81 years). Physical and internal surgery routine examinations were required, and the inclusion criteria were lack of malignant tumors, chronic infectious diseases, organ failure, and immune dysfunction were excluded from the study.

The study protocol was approved by the Ethics Committee of Linyi Central Hospital, Shandong, China. It is worth mentioning that informed consent was obtained from the patients.

The data were analyzed in SPSS software (version 17.0), and the results were expressed as mean±SD (x±s). Moreover, one-way analysis of variance (ANOVA) was used to make a comparison among the three groups. Additionally, Pearson correlation analysis between DM and concurrent groups was performed on clinical parameters with statistical
differences, and the correlation between parameters was also investigated in this study. Dual regression analysis between DM and concurrent groups was performed at the same time. Continuous variables were tested for normality, and one-way ANOVA was used to make multiple comparisons among the groups. In this study, the normal assumptions were made, and the non-parametric method was used for the variables that failed to meet the normal distribution. A p-value less than 0.05 was considered statistically significant.

4. Results

This study included 472 patients admitted to our hospital from June 2018 to June 2019. They were then divided into three groups of concurrent (n=168, T2DM complicated with CHD, DM (n=148, T2DM alone), and control (n=156, healthy controls). The concurrent and DM groups collectively referred to as the disease group, and the patients in the control group were selected from those who referred for a physical examination to our center in the hospital. There were no significant differences among the groups in terms of the demographic characteristics, such as height, weight, body mass index, as well as systolic and diastolic blood pressure (P>0.05).

Furthermore, no significant differences were observed between the disease and control groups regarding GA, HbA 1c, Glu, Hcy, TG, and apoA (P>0.05). The GA (P=0.013), HbA 1c (P=0.022), Glu (P=0.012), Hcy (P=0.031), and TG (P=0.021) were higher in the disease group, compared to the control group. However, the apoA (P=0.031) in the disease group was lower than that in the control group.

According to the results, there were significant differences between the concurrent and DM groups (P<0.05) in terms of GA (P=0.025), HbA 1c (P=0.033), Glu (P=0.019), Hcy (P=0.031), TC (P=0.022), LDL (P=0.037), and apoA (P=0.023). Table 1 tabulates the differences among the three groups in terms of GA, HbA 1c, Glu, Hcy, and apoA.

There was a significant difference between DM and concurrent groups in terms of HbA 1c (P=0.007), Glu (P=0.004), and GA (P=0.009). However, no significant difference was observed between these two groups regarding Hcy, apoA, and GA (P>0.05).

The correlation coefficients between HbA 1c and GA as well as Glu and GA were 0.680, and 0.465, respectively. It should be noted that the correlation between HbA 1c and GA was more significant than that between Glu and GA (Table 2). The 95% confidence intervals of GA, Hcy, and apoA were 0.765, 0.962; 1.015, 1.201; and 0.004, 0.403, respectively. A p-value of HbA 1c and Glu was over 0.05 which showed no significant differences (Table 3).

Table 1. Comparison between the disease and control groups in terms of clinical parameters

<table>
<thead>
<tr>
<th>Items</th>
<th>Control group (n=156)</th>
<th>DM group (n=168)</th>
<th>Concurrent group (n=148)</th>
</tr>
</thead>
<tbody>
<tr>
<td>GA (%)</td>
<td>15.77±1.44</td>
<td>28.58±8.01*a</td>
<td>27.14±6.34ab</td>
</tr>
<tr>
<td>HbA 1c (%)</td>
<td>5.5±0.5</td>
<td>9.5±1.5</td>
<td>8.3±1.2</td>
</tr>
<tr>
<td>Glu (mmol/L)</td>
<td>4.12±0.39</td>
<td>8.54±2.23*</td>
<td>8.62±3.56*</td>
</tr>
<tr>
<td>Hcy (mmol/L)</td>
<td>5.03±2.87</td>
<td>11.16±3.28*</td>
<td>17.56±6.36*</td>
</tr>
<tr>
<td>TG (mmol/L)</td>
<td>0.84±0.18</td>
<td>1.83±0.49*</td>
<td>1.25±0.61*</td>
</tr>
<tr>
<td>TC (mmol/L)</td>
<td>3.23±0.66</td>
<td>3.69±0.29</td>
<td>3.06±0.20</td>
</tr>
<tr>
<td>HDL (mmol/L)</td>
<td>1.35±0.15</td>
<td>1.01±0.30*</td>
<td>1.11±0.28*</td>
</tr>
<tr>
<td>LDL (mmol/L)</td>
<td>2.58±0.45</td>
<td>3.40±1.17*</td>
<td>2.57±0.12*</td>
</tr>
<tr>
<td>apoA (g/L)</td>
<td>1.30±0.18</td>
<td>1.10±0.12</td>
<td>0.95±0.12</td>
</tr>
<tr>
<td>apoB (g/L)</td>
<td>0.91±0.51</td>
<td>1.03±0.32</td>
<td>1.03±0.12</td>
</tr>
</tbody>
</table>

Note: GA: glycated albumin; HbA 1c: glyosylated hemoglobin; Glu: glucose; Hcy: homocysteine; TG: triglyceride; TC: total cholesterol; HDL: high-density lipoprotein; LDL: low-density lipoprotein. Compared with the control group, *P<0.05; compared with the DM group, #P<0.05.

Table 2. Correlation analysis between DM and concurrent groups in terms of GA and other clinical parameters

<table>
<thead>
<tr>
<th>Items</th>
<th>HbA 1c</th>
<th>Glu</th>
<th>Hcy</th>
<th>apoA</th>
</tr>
</thead>
<tbody>
<tr>
<td>r</td>
<td>0.680*</td>
<td>0.485*</td>
<td>0.023</td>
<td>0.014</td>
</tr>
<tr>
<td>P</td>
<td>0.001</td>
<td>0.000</td>
<td>0.642</td>
<td>0.703</td>
</tr>
</tbody>
</table>

Note: GA: glycated albumin; HbA 1c: glyosylated hemoglobin; Glu: glucose; Hcy: homocysteine. *P<0.01; #P<0.05, r is the correlation coefficient.

Table 3. Dual regression analysis between DM and concurrent groups

<table>
<thead>
<tr>
<th>Items</th>
<th>b</th>
<th>Standard error</th>
<th>P</th>
<th>OR</th>
<th>Confidence interval</th>
</tr>
</thead>
<tbody>
<tr>
<td>GA</td>
<td>-0.123</td>
<td>0.048</td>
<td>0.009*</td>
<td>0.846</td>
<td>0.765-0.962</td>
</tr>
<tr>
<td>HbA 1c</td>
<td>-0.098</td>
<td>0.171</td>
<td>0.562</td>
<td>0.925</td>
<td>0.648-1.265</td>
</tr>
<tr>
<td>Glu</td>
<td>-0.063</td>
<td>0.006</td>
<td>0.539</td>
<td>0.948</td>
<td>0.799-1.126</td>
</tr>
<tr>
<td>Hcy</td>
<td>-0.079</td>
<td>0.043</td>
<td>0.021*</td>
<td>1.156</td>
<td>1.015-1.201</td>
</tr>
<tr>
<td>apoA</td>
<td>-3.140</td>
<td>1.189</td>
<td>0.006*</td>
<td>0.039</td>
<td>0.004-0.406</td>
</tr>
</tbody>
</table>

Note: GA: glycated albumin; HbA 1c: glyosylated hemoglobin; Glu: glucose; Hcy: homocysteine. *P<0.01; #P<0.05; b is the partial regression coefficient; OR is the odds ratio.
5. Discussion

The GA is mainly used to monitor the mean blood glucose level in the past 2-4 weeks in T2DM patients (7). According to a study, it was found that GA ≥17.1% could screen out most undiagnosed diabetic patients (8). In addition to being able to monitor the patient’s blood glucose status, GA is more advantageous in determining the speed of treatment and adjusting the timeliness of administration (9). At the same time, for patients with reduced HbA 1c detection accuracy due to hemoglobin variation, GA can also be used as a necessary supplement that is not affected by red blood cell changes (10). As an early glycosylation product, GA is transformed and accumulated in the blood vessels causing vascular damage (especially to the microvessels) and leading to vascular sclerosis, stenosis, and obstruction, thereby promoting the occurrence of diabetic microvascular complications (11, 12).

Diabetes as one of the high-risk risks of CHD is one of the risk factors for T2DM patients complicated with CHD (13). This is an early and effective warning of high-risk CHD in patients with diabetes and also a timely screening of high-risk patients. It is especially important to reduce the incidence of adverse cardiovascular events and mortality (14). According to the Chinese Guidelines for the Prevention and Treatment of Type 2 Diabetes, most patients with T2DM have metabolic syndrome, which is accompanied by hypertension, hyperglycemia, and dyslipidemia (15, 16).

In this study, GA, HbA 1c, Glu, Hcy, and TG in the disease group were higher than those in the control group; however, the disease group had lower levels of HDL and apoA , compared to the control group, indicating that the results of the disease group were consistent with the Chinese Guidelines for the Prevention and Treatment of Type 2 Diabetes. There were significant differences between DM and concurrent groups in terms of GA, HbA 1c, Glu, Hcy, TC, LDL, and apoA (P<0.05). Moreover, the three groups were significantly different regarding GA, HbA 1c, Glu, Hcy, and apoA. Individual lipid profiles, including LDL, TC, and apoA abnormalities play an important role in the formation of coronary atherosclerosis; in addition, they have important clinical significance in the diagnosis of CAD, which is consistent with the results of the literature and fully confirmed the representativeness of the relevant clinical parameters included in the disease group (17). However, GA, HbA 1c, and Glu are the most representative indicators of blood glucose control in T2DM patients (18). The HbA 1c is the international gold standard for monitoring long-term glycemic control and is an indicator of glycemic control status in the past 1-2 months (19). A prospective study of diabetes in the United Kingdom found that for every 1% increase in HbA 1c, the risk of cardiovascular disease increased by 1% (20). The Glu value is the diagnostic cut-off point of T2DM. It is easy to measure at the bedside and grasp the static instantaneous blood glucose level of the patient; however, it is very susceptible to various factors (21). The HbA 1c and Glu play an important role in the development of CHD (3). Table 2 summarizes the correlation of GA with HbA 1c and Glu.

In the development of CHD, the changes of HbA 1c and Glu will inevitably lead to the change of GA; accordingly, GA has a certain effect on the progress of CHD. However, it is found in Table 3 that HbA 1c and Glu are highly correlated with GA in the concurrent group, which has no significance in the regression analysis (P>0.05) suggesting that HbA 1c and Glu can promote the development of CHD and have differences based on diabetes. Nonetheless, it cannot be a risk factor for T2DM complicated with CHD. However, GA showed a significant correlation in this regard (P<0.01). It can be seen that GA level monitoring is more predictive than HbA 1c and Glu for patients with potential CHD complicated with T2DM.

In addition to abnormally elevated blood glucose in patients with T2DM, this study also showed that T2DM was associated with high Hcy and low apoA. The Hcy is a vascular injury-causing amino acid, and hyperhomocysteinemia is associated with vascular endothelial cell injury and atherosclerosis (22). The apoA is an apolipoprotein closely related to HDL. According to the Framingham Heart Study, the risk of CHD decreased by 2% to 3% for every 0.03 mmol/L increase in HDL (23). As a protective factor, apoA reduces the corresponding protective effect of T2DM and accelerates the development of CHD (24). This study verified the above literature using the results of regression analysis. Table 3 displays that Hcy and apoA in patients with T2DM have statistical significance for complicated CHD (P<0.05). The apoA is reduced, and the trend of T2DM complicated with CHD is revealed with an increase in Hcy.

For diabetic patients, cardiovascular complications must be also prevented in addition to control blood glucose. Therefore, it is of great significance to search for early cardiovascular disease monitoring indicators of T2DM to guide the direction of clinical treatment. This study mainly discussed the relevance of GA to T2DM complicated with CHD. There is a significant correlation between T2DM complicated with CHD. Therefore, in the clinical stage, GA levels are highly correlated with HbA 1c and Glu that should be monitored early in T2DM, and potential CHD patients should be screened for early treatment. In addition, this study found that apoA was a protective factor for CHD complicated with T2DM, and Hcy was a risk factor. Combined with changes in Hcy and apoA, it was helpful for clinicians to understand the development and outcome of T2DM patients and improve the accuracy of screening. This study...
discussed the detection of serum biochemical indicators in elderly patients with T2DM and its relationship with CHD for the first time. Although previous studies have found a correlation between T2DM and the occurrence of CHD, there is no clear discussion from the perspective of blood biochemical indicators, and it cannot be used in clinical practice.

This study explored the relevance of CHD from the perspective of clinical biochemical indicators and guided the diagnosis, treatment, and prognosis of clinical diseases from a certain level. Moreover, this study set the research object to the elderly, a high-risk group of CHD and diabetes, which had more clinical significance.

6. Conclusion

High-level monitoring of the of GA, Hcy, and apoA in elderly patients with T2DM can not only improve the symptoms of patients with targeted treatment but also reduce the incidence of CHD by timely intervention, which is of great significance to improve the quality of life of high-risk patients. The GA, Hcy, and apoA may be important predictors of CHD in elderly diabetic patients. This study first proposed the indicators of cardiovascular disease in elderly diabetic patients, which were easily available in clinical testing.

Regarding the limitations of this study, one can name the focus on clinical patient data from one of our hospitals and a lack of multi-center research data. Due to the limited number of cases collected, it is still not enough to discuss whether GA is a factor in CHD complicated with T2DM. It is necessary to collect a large number of cases later for further discussion. The future investigations are recommended to focus on GA’s trend research on T2DM complicated with CHD, and conduct a large number of prospective experiments on the subject selection group.

Acknowledgements

None to declare.

References


