Role of inflammatory mediators (TNF-α, IL-6, CRP), biochemical and hematological parameters in type 2 diabetes mellitus patients of Kashmir, India

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Abstract

**Background:** Type II Diabetes mellitus (T2DM) is a multifactorial disease and a leading cause of premature deaths. Inflammatory cytokines are reported to have potential to enhance insulin resistance and hence T2DM. Assessment of immunological profile in T2DM patients of Kashmir valley is unclear. So, detection of cytokines is relevant to determine the extent and direction of immune responses. The current research was taken to study the role of inflammatory mediators in T2DM along with insulin sensitivity, biochemical and hematological parameters in mountainous valley of Kashmiri population.

**Methods:** A total of 340 subjects were selected in this study among them 160 were T2DM cases and 180 were healthy controls. Serum expression of inflammatory mediators (TNF-α and IL-6) were quantified by ELISA technique, WBC count was measured on Sysmex (Germany) hematology analyzer, biochemical and Immunoassay parameters were done on Abbott c4000 (USA) and Abbott C1000 (USA) fully automatic analyzer. Data was analyzed using statistical ‘software SPSS 16.1’ (Chicago, IL). For all assessments, \( p<0.05 \) were considered statistically significant.

**Results:** The expressions of candidate cytokines (TNF-α, IL-6, CRP, and WBC) were highly significant (\( p<0.001 \)) in T2DM. Among inflammatory mediators, TNF-α shows a positive correlation (\( p<0.001 \)) with glycemic profile and insulin sensitivity in T2DM cases in comparison with healthy normal. Biochemical (fasting sugar, HbA1c, insulin resistance, lipid profile) and anthropometric (BMI) parameters were highly significant (\( p<0.001 \)) in T2DM cases as compared to non-diabetic normal.

**Conclusion:** Low grade inflammation and up regulation of inflammatory mediators has been purported to play a significant role in pathogenesis of T2DM. Our findings confirm that positive correlation of TNF-α and IL-6 with T2DM and insulin sensitivity. These can act as early prediction biomarkers of T2DM. Further studies on wider range of pro and anti-inflammatory cytokines i.e. mediators, in association with other biochemical, immunoassay and hematological parameters are needed to help clinicians manage and treat T2DM effectively.

**Keywords:** Inflammation, Biomarkers, Cytokines, Kashmir, Mediators, Adult onset type 2 diabetes mellitus, SMHS.

**Conflicts of Interest:** None declared

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\[What is “already known” in this topic:\]
Effect of the pro and anti-inflammatory cytokines like (TNF-α, IL-6, CRP) has been reported in insulin signaling pathways, cross-linking and ultimately developing insulin resistance in β-cells of pancreas which further risks to T2DM. Stability among these Pro and anti-inflammatory cytokines is necessary to make β-cells immune to any infection which may lead to T2DM.

\[What this article adds:\]
Our experimental findings provide evidence that the pattern and variation of these cytokines (TNF-α, IL-6, CRP, and WBC) are important in the pathogenesis of T2DM. The significant correlation of these inflammatory mediators in T2DM cases with glycemic profile and insulin sensitivity leads to the pathogenesis of diseases in this ethnic population. These findings confirms that TNF-α, IL-6, plays a positive role in the pathogenesis of T2DM in this ethnic population of Kashmir and can act as an early prediction biomarkers.
Introduction

Type II diabetes mellitus (T2DM) is a non-communicable, chronic disorder and progresses slowly because of multifactorial etiology. T2DM is a leading cause of premature deaths worldwide, and its exceptional upsurge poses a severe threat and imposes a huge economic burden worldwide (825 billion dollars per year) (1). According to a recent estimation of the World Health Organization (WHO), 422 million people globally are affected from diabetes with a prevalence rate of 8.5% and 46.3% still remains undiagnosed and the number is projected to rise to 552 million in 2030. Furthermore, the maximum percentage lives in developing countries and comprises of 40–60 age group. In 2017 it was reported that India alone has 72 million people affected with T2DM and is projected to rise to 101.2 million in 2030 (2, 3). Most of the Indian populations are unaware of that disease. The risk factors of T2DM are suggestedly increased with changing lifestyle, blood pressure, central obesity, inadequate physical activity and unhealthy diet (4).

Fasting Blood glucose (FBG), 2-hour postprandial blood glucose (2hr-PP) and glycated hemoglobin (HbA1c) levels are most widely used as glycemic control markers which indicates the progression of the disease and development of its complications. T2DM is reported to be linked with lipid and lipoprotein irregularities, including reduced HDL cholesterol and raised triglycerides. (5-8). Trend of T2DM is changed from metabolic disorder to inflammation as effects of the pro and anti-inflammatory cytokines like tumor necrosis factor alpha (TNF-α), interleukin-6 (IL-6) and C-reactive protein (CRP) has been reported in insulin signaling pathways, cross-linking and ultimately developing insulin resistance in β-cells of pancreas which further risks to T2DM (9-11). Stability among these pro and anti-inflammatory cytokines is necessary to make β-cells immune to any infection which may lead to T2DM (12, 13). Some studies had reported high levels of CRP in T2DM as an independent biomarker for T2DM (12-16). The cytokines like TNF-α and IL-6 are expressed higher levels in activated macrophages and pro-inflammatory cytokine-like TNF-α induces insulin resistance by inhibiting phosphorylation of IRS-1 and Akt substrate 160 on insulin signaling cascade pathways (17). Ibfelt et al., also reported that TNF-α is decreasing in insulin production from β-cells of the pancreas, and suggests it as a possible marker which mediates the balance between insulin resistance and diabetes mellitus (18). Interleukin-6 (IL-6) is an inflammatory cytokine with multifunctional biologic effects. Few studies reported that it has an association with the pathogenesis of T2DM (19, 20). On the other hand, many studies suggest that IL-6 has a dual role of anti-inflammatory and improves glucose metabolism in T2DM patients (20, 21). IL-6 has a great impact on glucose homeostasis and metabolism by acting indirectly on the β-cells of the pancreas and on adipocytes (21-24). However, its role as an early biomarker for T2DM is still doubted and more such studies are needed in different populations. Many studies were conducted to assess the correlation of inflammatory markers with T2DM in other populations of the world; they reported significant association of these inflammatory markers (IL-6, TNF-α) (25-28). Kashmir valley is a division of Jammu and Kashmir State of India, which is mostly hilly and surrounded by snow-clad mountains. People of the valley are different in culture and food habits as compared to other states of India. Rice (Oryza sativa; carbohydrate diet) is a staple food of this region. Rapid urbanization and sedentary lifestyle of people are believed to be risk factors which leads to the pathogenesis of T2DM in this region (29, 4). A recent study done by the Indian Council of Medical Research (ICMR) reported that there was a prevalence of 6.1% of T2DM patients in Kashmir (30). Presently, no such study has been carried out on these inflammatory mediators (IL-6, TNF-α, CRP and WBC) and their association with T2DM in Kashmiri ethnic population. Hence, we hypothesized to study the role of these inflammatory mediators (IL-6 and TNF-α, CRP, WBC), biochemical and hemato logical parameters in T2DM cases and controls in association with glycemic profile and Insulin sensitivity in the people of the mountainous valley of Kashmir, India.

Methods

This study was conducted in Government Medical College (GMC) Srinagar at the Department of Biochemistry (Research centre- University of Kashmir) from September 2017 to January 2019.

Subjects

A total of 340 subjects were included in the study. The T2DM cases and healthy controls were enrolled for the study from OPD/IPD of the Sheri-Maharaja Hari Singh (SMHS) Hospital, and Associated Hospitals GMC Srinagar. Among 340 subjects, 160 T2DM patients were finally enrolled in the study. They were diagnosed as per standard of the American Diabetes Association (ADA) criteria 2016 (Table 1) and 180 healthy individuals (age-sex matched) were taken as sample controls. The study procedure and informed consent were reviewed by the ethics committee of GMC, Srinagar under Ref no. 89/ETH/GMC/ICMR as per Helsinki declaration. Written informed consent in local (Urdu/Kashmiri) and working language (English) with questionnaire response from patients and healthy controls was documented and the records were maintained as per hospital protocol.

Inclusion criteria: Patients of Kashmiri ethnicity, confirmed diagnosis of T2DM over two years.

Exclusion criteria: Non-Kashmiri origin, pregnant women, patients suffering from thyroiditis, rheumatoid

| Table 1. American Diabetes Association (ADA) 2016 Criteria for Diagnosing T2DM |
|---------------------------------|-----------------|
| FBG ≥126 mg/dl                 | Fasting means no food ingestion for ≥8 hours |
| 2-hr BG ≥200 mg/dl             | HbA1C ≥6.5%      |
| Random BG ≥200 mg/dl           |                               |
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Weight (kg) was measured by a digital weighing machine. The body mass index (BMI) of subjects was calculated by the formula  $\text{BMI} = \frac{\text{weight (Kg)}}{\text{height (m)}^2}$. Participants with a BMI of ≥30.0 kg/m² were considered obese as per NCEP ATPIII criteria. “Waist circumference” (WC) was evaluated in the middle, between the lower rib margin and the iliac crest with subjects in the upright position.

Criteria for Sample size

Sample size was calculated to detect an effect size of 0.10 at type 1 error of 50% and power of 80% using “G Power version 3.1.9.2”. Given the above parameters, a total size of 340 (160 in T2DM and 180 in controls in each group) will be sufficient for this study.

Sample collection and separation

Five ml blood was collected by phlebotomists by venipuncture from patients and healthy controls at the common collection centre facility of Government Shri-Maharaja Hari Singh Hospital (SMHS), Srinagar. The patients and healthy controls were recruited from the Out-Patient Department (OPD)/ In-Patient Department (IPD) of Post Graduate Department of Medicine, GMC, Srinagar. Blood was immediately transferred into 3ml green top heparin vial and 2ml blood into Purple top EDTA vial. Heparinized 3ml blood was centrifuged at 4000 RPM for 5 minutes and plasma was aliquoted into Eppendorf tubes for ELISA and other biochemical analysis stored at -20°C till further analysis.

Biochemical and Immunoassay Analysis

Glycated hemoglobin (HbA1c) levels and Biochemistry of subjects was documented by calculating the index of Insulin resistance (IR) of subjects was calculated by using the formula: “HOMA-IR = fasting insulin (μU/ml) × fasting glucose (mg/dl)/405” (24). Following calculation, HOMA-IR score was used as a reference range for classification of IR.

a) < 3 = Normal IR  
b) Between 3 and 5 = Moderate IR  
c) >5 =Severe IR

Hematological analysis: Whole blood samples taken in EDTA vials were analyzed for WBC count on Sysmax hematology analyzer (Germany).

Enzyme-linked Immunoassay (ELISA) Analysis

TNF-α assay: Quantitative measurement of TNF-α was done by the ‘Diaclone Human TNF-α ELISA kit’ (Catalogue No. 950.090.09, France). The minimum measurable concentration was estimated to be 8pg/ml. The coefficient of variation for TNF-α was 3.3 % and 9.0 % (intra and inter-assay) respectively. The subsequent washing and absorbance reading were done on microplate reader Lablifeer ew 2007(Germany). Samples were tested in triplicate and a calibration curve was added to each plate.

IL-6 assay: Quantitative measurement of IL-6 was done by use of a commercial ELISA kit; ‘Diaclone Human IL-6 ELISA kit’ (Catalogue No. 950.030.096, France). The minimum measurable concentration was estimated to be 2pg/ml. The coefficient of variation for IL-6 was 3.6% and 7.7% (intra and inter-assay), respectively. The subsequent washing and absorbance reading were done on microplate reader Lablifeer ew 2007(Germany). Samples were tested in triplicate and a calibration curve was added to each plate.

Statistical analysis

Data was inserted on Microsoft excel 2011 spreadsheet. All the data were expressed as a mean ± standard deviation and significance value (p) were calculated. Data analysis was performed by using statistical ‘software SPSS 16.1’ (Chicago, IL). Students’ T-test was done on biochemical, immunoassay and inflammatory mediators. Chi-square test was done on socio-demographic characters. Correlation analysis was performed for determining the association between serum IL-6, TNF-α, CRP and WBC levels and Pearson correlation coefficient (r) was obtained. For all assessments, p<0.05 were considered statistically significant.

Results

Anthropometric analysis of the study group

Table 2 summarizes the Anthropometric characteristics of 340 subjects, including 160 prevalent T2DM cases and 180 healthy controls. The mean± SD age of T2DM patients was 48.9 ±8.4 years, and that of controls was 44.9±9.8 years which is statistically significant (p=0.002). It was observed that in T2DM patients, mean±SD of BMI was 41.2±7.1 kg/m2 and in healthy controls was 20.2±2.2 kg/m2 which is statistically significant (p=0.001). There was no significance (p=0.472) found in waist circumference (WC) within T2DM patients and healthy controls. Among 160 cases, 81 were males and 79 females and in healthy controls, 94 were males and 86 were females, on gender-wise comparison difference in patients and controls are non-significant (p=0.829).

Table 2: Anthropometric analysis of T2DM cases and Controls

<table>
<thead>
<tr>
<th>Variables</th>
<th>T2DM Cases (n=160)</th>
<th>Controls (n=180)</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age (Years)</td>
<td>48.9±8.4</td>
<td>44.9±9.8</td>
<td>0.002</td>
</tr>
<tr>
<td>Sex (M/F)</td>
<td>81/79</td>
<td>94/86</td>
<td>0.829</td>
</tr>
<tr>
<td>BMI (kg/m2)</td>
<td>41.2±7.1</td>
<td>20.2±2.2</td>
<td>0.001</td>
</tr>
<tr>
<td>WC(inches)</td>
<td>47.4±4.6</td>
<td>29.2±4.5</td>
<td>0.472</td>
</tr>
</tbody>
</table>

Data are presented as means ± SD. The significance of the difference is based on the one-way T-test. “BMI: Body mass index; WC: Waist Circumference.”
Socio-demographic profile of the study group

Social class scale is based on Residence, Smoking, Lifestyle and Education items. 43.1% of the T2DM patients were urban and 56.8% belongs to the rural population, while among healthy controls, 41.6% were urban, and 58.3% belongs to the rural population. 31.2% of the T2DM patients were smokers and 68.7% were non-smokers, while among healthy controls, 2.7% were smokers and 97.2% were non-smokers. 31.2% of the T2DM patients were active and 68.7% were sedentary, while among healthy controls, 58.3% were active and 41.6% were sedentary. 36.8% of the T2DM patients were literate and 63.1% were illiterate, while among healthy controls, 55.0% were literate, and 45.0% were illiterates.

Biochemical profile of the study group

The biochemical profile of T2DM cases and healthy controls was summarized in Table 3 and showed that there was an increasing trend in parameters of lipid profile like Serum Triglycerides (TG), total cholesterol (TC), Low Density Lipoprotein (LDL) and High Density Lipoprotein (HDL) among T2DM cases as compared to healthy controls and the trend were significantly high (p<0.05). The glycemic profile (Glucose Fasting and HbA1c) in T2DM cases was higher as compared to healthy controls and are found to be statistically significant (p<0.05).

Table 3. Biochemical analysis of the study group

<table>
<thead>
<tr>
<th>Variables</th>
<th>Diabetes mellitus (n=160)</th>
<th>Controls (n=180)</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Glucose Fasting (mg/dl)</td>
<td>166.4±31.9</td>
<td>80.9±6.8</td>
<td>0.111</td>
</tr>
<tr>
<td>Post-parandial (mg/dl)</td>
<td>314.2±50.9</td>
<td>121.1±8.2</td>
<td>0.001</td>
</tr>
<tr>
<td>Cholesterol (mg/dl)</td>
<td>296.5±53.3</td>
<td>107.1±26.8</td>
<td>0.001</td>
</tr>
<tr>
<td>Triglycerides (mg/dl)</td>
<td>318.5±56.3</td>
<td>145.1±29.4</td>
<td>0.001</td>
</tr>
<tr>
<td>HDL (mg/dl)</td>
<td>91.4±21.7</td>
<td>51.9±11.0</td>
<td>0.021</td>
</tr>
<tr>
<td>LDL (mg/dl)</td>
<td>147.3±8.0</td>
<td>68.8±28.9</td>
<td>0.001</td>
</tr>
<tr>
<td>HbA1c (%)</td>
<td>9.8±2.8</td>
<td>4.7±0.8</td>
<td>0.011</td>
</tr>
</tbody>
</table>

Data are presented as means ± SD. The significance of the difference is based on the one-way T-test.

Immunological profile of study group

Table 5 describes the levels of serum inflammatory mediators (IL-6, TNF-α, CRP and WBC) in T2DM cases and healthy controls; the mean ± SD value of inflammatory markers in T2DM cases was TLC=8395±1843 IL-6=19.2±7.2, TNF-α=34.5±8.8 and CRP=4.2±0.9 while in healthy controls it was TLC =7289±1404, IL-6 =3.0±1.4, TNF-α =12.7±3.4 and CRP=1.4±0.6; and it was found that in T2DM patients the levels of inflammatory mediators were highly significant (p<0.05) in comparison with healthy controls.

Comparison of Inflammatory mediators in the study group

Table 6 shows the comparison of inflammatory mediators within gender groups and it was found that in female

Table 4. Immunoassay analysis of study group

<table>
<thead>
<tr>
<th>Parameters</th>
<th>Diabetes mellitus (n=160)</th>
<th>Controls (n=180)</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Insulin (μU/ml)</td>
<td>32.6±7.5</td>
<td>7.8±2.1</td>
<td>0.001</td>
</tr>
<tr>
<td>HOMA-IR</td>
<td>13.5±4.7</td>
<td>1.5±0.5</td>
<td>0.001</td>
</tr>
</tbody>
</table>

Data are presented as means ± SD. The significance of the difference is based on the one-way T-test. p value <0.05 are statistically significant.

Table 5. Inflammatory mediators in the study group

<table>
<thead>
<tr>
<th>Variables</th>
<th>Diabetes mellitus (n=160)</th>
<th>Controls (n=180)</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>TLC (thousands)</td>
<td>8395±1843</td>
<td>7289±1404</td>
<td>0.001</td>
</tr>
<tr>
<td>IL-6 (pg/ml)</td>
<td>19.2±7.2</td>
<td>3.0±1.4</td>
<td>0.001</td>
</tr>
<tr>
<td>TNF-α (pg/ml)</td>
<td>34.5±8.8</td>
<td>12.7±3.4</td>
<td>0.001</td>
</tr>
<tr>
<td>CRP (mg/dl)</td>
<td>4.2±0.9</td>
<td>1.4±0.6</td>
<td>0.001</td>
</tr>
</tbody>
</table>

Data are presented as means ± SD. The Significance of the difference is based on the one-way T-test. p value < 0.05 are statistically significant

Table 6. Comparison of Inflammatory mediators in T2DM Male and Female patients versus control subjects

<table>
<thead>
<tr>
<th>Inflammatory Mediators</th>
<th>Male T2DM (n=81)</th>
<th>Female T2DM (n=79)</th>
<th>Male controls (n=94)</th>
<th>Female controls (n=86)</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>TNF-α (pg/dl)</td>
<td>8.8±0.9</td>
<td>8.9±1.0</td>
<td>3.5±0.3</td>
<td>3.3±0.3</td>
<td>0.000</td>
</tr>
<tr>
<td>IL-6 (pg/dl)</td>
<td>6.8±0.7</td>
<td>7.6±0.8</td>
<td>1.4±0.1</td>
<td>1.4±0.1</td>
<td>0.000</td>
</tr>
<tr>
<td>CRP (mg/dl)</td>
<td>0.8±0.1</td>
<td>1.0±0.1</td>
<td>0.6±0.06</td>
<td>0.6±0.06</td>
<td>0.000</td>
</tr>
<tr>
<td>WBC (thousands)</td>
<td>1944±216</td>
<td>1741±195</td>
<td>1448±149</td>
<td>1350±145</td>
<td>0.072</td>
</tr>
</tbody>
</table>

Data are presented as means ± SD. The significance of the difference is based on the one-way T-test. p value < 0.05 is statistically significant.
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cases levels of inflammatory mediators was highly significant (p<0.05) as compared to male cases while WBC was not statistically significant, which provides us the information that females may be at higher risk to T2DM.

Correlation of inflammatory mediators in the study group

Figure 1 (a) to 1 (d) show the correlation of inflammatory mediators in T2DM cases and controls with glycemic profile and insulin sensitivity and was studied by Pearson’s correlation analysis. TNF-α shows a positive correlation with the glycemic profile (Glucose fasting, Hba1c) and insulin sensitivity (Insulin assay, HOMA-IR) in T2DM cases and were statistically significant (p<0.05).

Table 7 describes the association of inflammatory mediators with glycemic profile and Table 8, depicts the association of insulin sensitivity as per gender-wise in cases and controls. We observed in males and female T2DM cases that there was a positive correlation (p=0.001) of TNF-α with glycemic profile and insulin sensitivity and other inflammatory mediators show negative and weak correlation.
Correlation of inflammatory mediators with BMI in the study group

Table 9 shows the correlation of inflammatory mediators with anthropometric parameter (BMI) within T2DM cases and healthy controls and it was found that there was no any association between these parameters (p<0.05).

Discussion

Worldwide, people are suffering from T2DM and it is projected to increase from the present 415 million people to 642 million by 2040. In all developing countries, it was seen that the number of T2DM patients is increasing and 75% of people with T2DM are living in these developing countries (31). In this study, we observed that sociodemographic factors like education, lifestyle and smoking have significant associations with T2DM except for residence (urban and rural areas of the same geographical area) which had no substantial influence on the levels of inflammatory mediators of study like, TNF-α, CRP, IL-6 and WBC. After a thorough literature review on the dynamics of T2DM and inflammatory mediators on the Kashmiri population, this is the first kind of study conducted in the mountainous valley of Kashmir on inflammatory mediators with T2DM except for residence in the study group.

Our study proved that only pro-inflammatory cytokine TNF-α IL-6, could leads to the pathogenesis of T2DM while other inflammatory cytokines show negative and weak correlations with T2DM. This research study showed vibrant changes in concentrations of pro-inflammatory cytokines, in T2DM. Our findings are in concurrence with the results of Samuel et al. which showed serum expression of candidate mediators (TNF-α, IL-6 and CRP) are elevated in T2DM cases which are independent of physical activity and other risk factors (35). It is suggest that TNF-α is an important predictor for the development of T2DM for males and females, in both rural and urban populations of the Kashmir population. These results which support the hypothesis that systemic inflammation is a common precursor for T2DM (24, 25, 35).

Interestingly, the results of our study showed a high degree of correlation between these promising cytokines (CRP, TNF and IL-6, WBC) and BMI and showed increased levels of these candidate cytokines (WBC, CRP, TNF-α, IL-6) in T2DM in comparison to healthy controls. The results are statically significant. Hence, our study support in this ethnic population of Kashmir valley that TNF-α inflammatory mediator’s association and correlation are higher in T2DM cases as compared to controls, which somehow in mechanism create insulin resistance and predispose patients to T2DM if not treated at an early stage. IL-6 Inflammatory marker shows a weak correlation with T2DM. As we know, IL-6 has a pleiotropic nature that can act as an anti-inflammatory agent. In this case-control study, we found in our T2DM cases a significantly higher concentration of TNF-α as compared to those of controls which may be the possible cause of low-grade inflammation and predisposed subjects to the T2DM or towards its complications. These assertions aggress with the findings of AL-Shukaili et al. (37). Furthermore, our experimental finding provides evidence that the pattern and variation of these cytokines (TNF-α, IL-6, CRP and WBC) are important in the pathogenesis of T2DM. The significant correlation of TNF-α inflammatory mediator in T2DM cases with glycemic profile and insulin sensitivity leads to the pathogenesis of diseases in this ethnic population. These findings are in agreement with the fact that inflammatory reactions depend on the group of cytokines rather than a single one. The role of inflammation in the pathogenesis of T2DM has been elucidated in several studies in different populations (36-38). Furthermore, investigations of these cytokine inflammatory mediators in this population is needed to establish their importance both as early biomarkers and diagnostic predictors for T2DM.
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Conclusion

Our data confirms that TNF-α, IL-6, inflammatory mediator i.e., cytokines, plays a positive role in the pathogenesis of T2DM in this ethnic population of Kashmir and can act as early prediction biomarkers which can prevent T2DM in this population. Further studies on the wider range of these inflammatory mediators in association with other biochemical, immunoassay and hematological parameters are needed to establish role of inflammatory markers as early prediction biomarkers which can prevent T2DM in this population.

Conflict of Interests

The authors declare that they have no competing interests.

References


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