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Some physiochemical changes associated with type 2 diabetes mellitus in Benin City, Nigeria

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ABSTRACT

The present study was designed to determine the level of malondialdehyde (MDA) concentration, serum c reactive protein (CRP) and CD4⁺ T cells in type 2 diabetes mellitus in a view to assessing the complications associated with the non specific activation of CD4⁺ T cells. A total of 80 subjects aged between 18 and 50 years gave their consent to participate in the study. They consist of 40 known type 2 diabetes mellitus and 40 aged matched apparently healthy control subjects on routine checkup. Malondialdehyde concentration was determined by spectrophotometric method. Serum c reactive protein was determined by enzyme-linked immunosorbent assay method. CD4⁺ T cells were estimated using Partec cyflow counter. There was a significant increase in MDA concentration, CRP and CD4⁺ T cells in type 2 diabetes mellitus subjects when compared with the aged matched control group ($P < 0.05$). Type 2 diabetes mellitus causes an increased production of free radicals which is possibly linked to the complications associated with non specific activation of CD4⁺ T cells.

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Keywords: Free radicals, CD4+ T cells, diabetes mellitus, cytokines, c reactive protein, malondialdehyde.

INTRODUCTION

The prevalence of diabetes mellitus is increasing rapidly worldwide and the World Health Organization (2003) has predicted that by 2030 the number of adults with diabetes would have almost doubled worldwide, from 177 million in 2000 to 370 million (Rowley and Bezold, 2012). Diabetes mellitus is a complex syndrome characterized by absolute or relative insulin deficiency leading to hyperglycemia and an altered metabolism of glucose, fat, and protein (Remuzzi et al.,

2002; Caramori and Mauer, 2003; Wolf and Ritz, 2003).

Since more than 90% of the global cases of diabetes mellitus are type 2, it is evident that the epidemic is mainly due to the escalation of the causes of type 2 diabetes mellitus. According to the current definition, two fasting glucose measurement above 126 mg/dl (7.0 mmol/L) are considered diagnostic for diabetes mellitus or Glycated hemoglobin (HbA1C) above 6.5% (Sayalah et al., 2001; WHO, 2003; Motala, 2008; Chinenye et al.,

2008). Experimental evidence demonstrated that c reactive proteins is a sensitive physiological biomarker of sub clinical systemic inflammation which is associated with hyperglycemia, insulin resistance and overt type 2 diabetes mellitus (Su et al., 2010).

Lipid peroxidation is a well established mechanism of cellular injury and is used as an indicator of oxidative stress in cells and tissues. The measurement of malondialdehyde is widely used as an indicator of lipid peroxidation and increased level of the peroxidation products have been associated with a variety of acute and chronic pathophysiological processes including type 2 diabetes mellitus (Killic et al., 2003). Recent research suggests that patients with elevated basal level of CRP are at an increased risk of diabetes mellitus, hypertension and cardiovascular disease (Clearfield, 2005). However; could the elevation of MDA and CRP play a major role in diabetogenesis? This question, if explored, may open new doors in the understanding of the complication associated with type 2 diabetes mellitus. Thus, the present study was designed to determine the level of MDA, CRP and CD4⁺ T cells in type 2 diabetes mellitus in a view to assessing the complications associated with the non specific activation of CD4⁺ T cells.

MATERIALS AND METHODS

Subjects

A total of 80 subjects aged between 18 and 60 years gave their consent to participate in the study. They consist of 40 known type 2 diabetes mellitus and 40 aged matched apparently healthy control subjects on routine checkup. They were confirmed to be diabetics and non diabetics respectively using the glucose oxidase method as previously described by Ehiaghe et al. (2013).

Inclusion criteria

Both males and females were included in the study.

Exclusion criteria

Pregnant women and those diagnosed with other systemic diseases were excluded from this study.

Physiological measurements

The subjects resting systolic blood pressure and diastolic blood were measured from the right arm as described by Musa et al. (2002). Using an automated digital electronic BP monitor (Omron digital BP monitor, Model 11 EM 403c; Tokyo Japan).

Collection of blood samples

Nine milliliters of fasting venous blood was collected from the medial cubital vein using vacutainer and needle from each of the subjects and shared equally into ethylene diamine tetra acetic acid, sodium fluoride potassium oxalate and plain containers.

Determination of blood glucose level

Glucose oxidase method was used in the determination of blood glucose as previously described by Ehiaghe et al. (2013). In brief, 10 µl of the standard or test samples was added to 1000 µl of glucose reagent 1, mixed and incubated for 25 minutes at 20 °C. The absorbance of the standard and test samples was measured against the reagent blank within 60 minutes at a wavelength of 540 nm using a spectrophotometer.

Determination of serum c reactive protein

Enzyme-linked immunosorbent assay (ELISA) was used in the determination of the level of c reactive protein in the serum. The assay employs an antibody specific for c reactive protein coated on a 96 well plate. Briefly, 100 µl of the assay diluents was added to each well. 50 µl of standard or

sample(s) was added per well and mixture was incubated for 2 hours. The solution was discarded and microplates washed four times with 300 μ l of 1X wash solution. 200 μ l c reactive protein conjugate was added to the standard or sample(s) and covered with a sealing tape. It was incubated at room temperature for 2 hours. The solution was discarded and microplates washed four times with 300 μ l of 1X wash solution. 200 μ l of tetraethyl benzidine substrate was added to each well and incubated for 30 minutes at room temperature in the dark with gentle shaking. 50 μ l of stop solution was added to each microplate. The intensity of the color developed was measured at 450 nm wavelength using stat fax® 4700 micro strip reader as previously described by Ehiaghe et al. (2013).

CD4⁺ T cell count estimation

CD4⁺ T cell count was estimated using Partec cyflow counter as described by Partec cyflow counter (2006).

Malondialdehyde estimation

Reagent 1: (Trichloroacetic acid 17.5%);

Reagent -2 :(Trichloroacetic acid 70%);

Reagent – 3: (Thiobarbituric acid 0.6%) TBA

Malondialdehyde in the serum was determined as conjugated with TBA. The reaction was performed in 18 \times 150 mm Pyrex test tube labeled as: test and blank, into which 1ml each of reagents 1, 2, and 3 was added into them respectively. 1 ml of the serum was added to the test tube labeled test while 1 ml of distilled water was added to the test tube labeled blank. The tubes were mixed well and incubated in boiling bath for 15 minutes, allowed to cool. The tubes let to stand at room temperature for 20 minutes. Then the tubes centrifuged at 2000 rpm for 15 minutes. The supernatant layer was read at 534 nm with a spectrophotometer. The concentration of MDA

(μ mol/L) was calculated by using the following formula: Concentration of the test= Abs (test) – Abs (blank) / 1.56 x 1000000 as previously described by Mossa et al. (2009).

Statistical analysis

All numerical variables were expressed in mean (\pm SD) and analyzed using one way analysis of variance (ANOVA).Using SPSS version 20.0. Significant level was considered at $P < 0.05$.

Ethics

Ethical approval was obtained from the ethical committee of the Lahor Research Laboratories and Medical centre in Benin City, Edo State, Nigeria with reference number LRL/008/014.

RESULTS

Our findings revealed that there was no significant difference in SBP, DBP, Glucose, CRP, CD4⁺ T cells and MDA concentration of the males type 2 diabetes mellitus when compared with females type 2 diabetes mellitus ($P > 0.05$). There was also no significant difference with males control group when compared with females control group ($P > 0.05$), (Table 1).

There was a significant increase in SBP, DBP, Glucose, CRP, CD4⁺T cells and MDA concentration of type 2 diabetes mellitus when compared with the aged matched control group ($P < 0.05$), (Table 2). Our finding also revealed a significant increase in SBP, DBP, Glucose, CRP, CD4⁺ T cells and MDA concentration of the male type 2 diabetes mellitus when compared with the males control group ($P < 0.05$), (Table 3). A significant increase in SBP, DBP, Glucose, CRP, CD4⁺ T cells and MDA concentration of the females type 2 diabetes mellitus compared with the females control group ($P < 0.05$), (Table 3).

Table 1: Comparison of mean (\pm SD) SBP (mm.Hg), DBP (mm.Hg), Glucose (mg/dl), CRP (ng/l), CD4⁺ T cell count (cells/ μ l), and MDA (μ mol/L) between type 2 diabetes mellitus (DM) groups and the control groups respectively.

Parameter	SBP	DBP	Glucose	CRP	CD4 ⁺ T cell count	MDA
Male type 2 DM(n=22)	140 \pm 10.4	83 \pm 7.1	162 \pm 43	144 \pm 46	660 \pm 69	3.1 \pm 0.29
Female type 2 DM (n=18)	140 \pm 8.8	83 \pm 7.5	151 \pm 41	139 \pm 52	661 \pm 80	2.9 \pm 0.34
P- value	>0.05	>0.05	>0.05	>0.05	>0.05	>0.05
Male control group (n=22)	116 \pm 3.5	74 \pm 4.2	89 \pm 11	1.5 \pm 1.7	504 \pm 74	1.56 \pm 0.4
Female control group (n=18)	117 \pm 3.3	73 \pm 4.6	88 \pm 14.7	1.8 \pm 2.3	508 \pm 60	1.49 \pm 0.35
P-value	>0.05	>0.05	>0.05	>0.05	>0.05	>0.05

P < 0.05 was considered significant.

Table 2: Comparison of mean (\pm SD) SBP (mm.Hg), DBP (mm.Hg), Glucose (mg/dl), CRP (ng/l), CD4⁺ T cell count (cells/ μ l), and MDA (μ mol/L) of the type 2 DM with the control groups irrespective of gender.

Parameters	SBP	DBP	Glucose	CRP	CD4 ⁺ T cell count	MDA
Type 2 DM (n=40)	140 \pm 1.51	83 \pm 1.13	160 \pm 6.7	141 \pm 7.6	661 \pm 75	3.0 \pm 0.05
Control group (n=40)	116 \pm 0.54	73.4 \pm 0.69	88.5 \pm 2.02	1.7 \pm 0.3	506 \pm 67	1.53 \pm 0.06
P- value	<0.05	<0.05	<0.05	<0.05	<0.05	<0.05

P < 0.05 was considered significant

Table 3: Comparison of mean (\pm SD) SBP (mm.Hg), DBP (mm.Hg), Glucose (mg/dl), CRP (ng/l), CD4⁺ T cell count (cells/ μ l), and MDA (μ mol/L) of type 2 DM and control groups with gender biased.

Parameter	SBP	DBP	Glucose	CRP	CD4 ⁺ T cell count	MDA
Male type 2 DM(n=22)	140 \pm 10.4	83 \pm 7.1	162 \pm 43	144 \pm 46	660 \pm 69	3.1 \pm 0.29
Male control group (n=22)	116 \pm 3.5	74 \pm 4.2	89 \pm 11	1.5 \pm 1.7	504 \pm 74	1.56 \pm 0.4
P- value	<0.05	<0.05	<0.05	<0.05	<0.05	<0.05
Female type 2 DM (n=18)	140 \pm 8.8	83 \pm 7.5	151 \pm 41	139 \pm 52	661 \pm 80	2.9 \pm 0.34
Female control group (n=18)	117 \pm 3.3	73 \pm 4.6	88 \pm 14.7	1.8 \pm 2.3	508 \pm 60	1.49 \pm 0.35
P-value	<0.05	<0.05	<0.05	<0.05	<0.05	<0.05

P < 0.05 was considered significant.

DISCUSSION

The prevalence of diabetes mellitus is increasing rapidly worldwide and World Health Organization (2003) has predicted that by 2030 the number of adults with diabetes would have almost doubled worldwide. There is scarcity of published report on the physiochemical changes associated with type 2 diabetes mellitus in Benin City, hence these study. A significant increase in malondialdehyde concentration seen in both males and females type 2 diabetes mellitus subjects as compared with the control groups (Tables 2 and 3). This could be attributed to the increased production of free radical occasioned by the auto-oxidation of glucose and lipid peroxidation. This is in accordance with these findings. The increased level of lipid peroxides can cause oxidative injury to blood cells, cross-linking of membrane lipids and protein, increasing of cell ageing, imbalance of prostacyclin/prostaglandin and vasoconstriction (Jain et al., 1996). Our findings also revealed that there was no significant difference in SBP, DBP, Glucose, CRP, CD4⁺ T cells and MDA concentration of the males type 2 diabetes mellitus when compared with females type 2 diabetes mellitus (Table 1). This may be attributed to the common outcome of diabetogenesis regardless of gender affiliation. The aberrant production of inflammatory cytokines and chemokines as well as differential activation of T helper cells and monocytes is the underlying immunopathological mechanism of type 2 diabetes mellitus patients (Lim et al., 2009).

The significant increase in CD4⁺ T cells, CRP, SBP and DBP of the diabetics groups (Tables 2 and 3) when compared with the non diabetics groups (Table 1 and 2). This could be linked to the non specific activation of the T helper cells by free radical generated from the auto-oxidation of glucose, lipid, protein peroxidation and a possible

destruction of endothelial cells of the renal and cardiovascular system by the non specific mediator released by the activated T cells. This is in accordance with the findings. Activated T cells can cause injury directly through cytotoxic effects and indirectly by recruiting and activating macrophages. Proinflammatory cytokines secreted by T (CD4⁺, CD8⁺) cells could activate neighboring macrophages directly or by stimulating mesangial cell production of colony stimulating factor-1 and MCP-1 indirectly. Once macrophages have activated, they can release nitric oxide, reactive oxygen species, IL-1, TNF- α , complement factors, and metalloproteinase, all of which promote renal injury (Wang and Harris, 2011). An increased body of evidence indicates that immigrated blood leukocytes might considerably alter the phenotype of endothelial cells and increase inflammation of the vascular bed (Endemann and Schiffrin, 2004). Endothelial dysfunction is associated with most forms of cardiovascular disease, such as coronary artery disease, chronic renal failure and Diabetes mellitus (Stehouwer, 2004). The interaction of renal tissue macrophages and T cells produces various reactive oxygen species, proinflammatory cytokines, metalloproteinase, and growth factor, which modulate the local response and increase inflammation within the cells (Yamagishi et al., 2007). Experimental evidence demonstrated that c reactive proteins is a sensitive physiological biomarker of sub clinical systemic inflammation which is associated with hyperglycemia, insulin resistance and overt type 2 diabetes mellitus (Su et al., 2010).

Conclusion

It was observed that type 2 diabetes mellitus causes an increased production of free radical which is possibly linked to the non specific activation of CD4⁺ T cells. The

molecular mechanism underlying the complications associated with the non specific activation of CD4⁺ T Cells needs further investigation.

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