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RESEARCH ARTICLE

CELLULAR FIBRONECTIN AS A BIOMARKER FOR ASSESSING ENDOTHELIAL FUNCTION IN DIABETIC PATIENTS

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ABSTRACT

Background: Retinopathy and neuropathy are major chronic complications of diabetes mellitus as result of poor glycaemic control. Cellular fibronectin is derived from an endothelium of the blood vessel, it plays role in sub endothelial matrix assembly. Increased in cellular fibronectin in the plasma is associated with injury to the endothelium of blood vessels. **Aim of the study:** Is to evaluate the plasma levels of cellular fibronectin in diabetic patients. **Materials and methods:** Cellular fibronectin was determined in one hundred (100) diabetic subjects and one hundred (100) non-diabetic subjects using sandwich Enzyme-Linked immunosorbent Assay. While (45%) of the diabetic subjects were males, (55%) were females, (48%) of the non-diabetic subjects were males and (52%) were females. However, out of (45) of the diabetic male subjects, 20(44%) present with erectile dysfunction and 25% of the diabetic subjects have blurred vision. Approval of the study was obtained from the Ethics and Research Committee of the Specialist Hospital, Sokoto. **Results:** Cellular fibronectin was elevated in diabetic subjects ($31.21 \pm 1.62 \text{mg/L}$) compared with non-diabetic subjects ($13.19 \pm 1.20 \text{mg/L}$). Cellular fibronectin was elevated in diabetic subjects with erectile dysfunction ($49.00 \pm 3.76 \text{mg/L}$) compared to diabetic with no erectile dysfunction ($32.20 \pm 2.31 \text{mg/L}$). Cellular fibronectin was also elevated in diabetic subjects with blurred vision ($45.52 \pm 3.32 \text{mg/L}$) compared to diabetic subjects with no blurred vision ($34.00 \pm 1.91 \text{mg/L}$). **Conclusion:** Determination of plasma level of cellular fibronectin in diabetic subjects could be useful as an adjunct in predicting early vascular complication in diabetic patients.

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INTRODUCTION

The endothelium is made up of a single layer of cells, that covered the vascular lumen. (Moreira *et al.*, 2018). It is now identified, however, that endothelial cells are metabolically active with important paracrine, endocrine and autocrine functions which are indispensable for the maintenance of vascular homeostasis under physiological conditions (Sena *et al.*, 2013; Tabit *et al.*, 2010). The functions of vascular endothelium include; regulation of vessel integrity, vascular growth and reconstruction, tissue growth and metabolism, immune responses, cell adhesion, angiogenesis, haemostasis and vascular permeability (Cheok *et al.*, 2018). The endothelium plays an initial role in the regulation of vascular tone, controlling tissue blood flow and inflammatory responses and also, maintaining blood fluidity (Li *et al.*, 2017b; Cheok *et al.*, 2018). It does so, by producing components of the extracellular matrix such as cellular fibronectin (cFn) and

variety of regulatory mediators like; NO (nitric oxide), ET-1 (endothelin-1), Ang II (angiotensin II), t-PA (tissue-type plasminogen activator), PAI-1 (plasminogen activator inhibitor-1), VWF (von Willebrand factor) and tumour necrosis factor alpha (TNF α) (Klemis *et al.*, 2017). Failure of the endothelium to perform its functions from any means is referred to endothelial dysfunction (Yada *et al.*, 2018). The hallmark of diabetes mellitus, chronic hyperglycaemia, has been implicated in the development of endothelial dysfunction (Habib and Ali, 2018). The Principal mechanisms of endothelial dysfunction by hyperglycaemia involve Protein Kinase C (PK C) activation, Activation of the hexosamine and polyol pathways, Formation of advanced glycation end-products (AGEs). These pathways are said to mediate vascular dysfunction through Reactive Oxygen Species (ROS) overproduction, most importantly is the increases in O₂⁻ (Marzona *et al.*, 2017). Activation to the endothelium could result to increased synthesis of cellular fibronectin in response to the endothelial injury (Kassas *et al.*, 2016). Fibronectins (FN) are large glycoproteins found in plasma, in extracellular matrix, and on cell surfaces (Doddapattar *et al.*, 2018). They promote cell-cell and cell-matrix interactions and thus play a

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role in tissue construction and reconstruction (Kanters *et al.*, 2001). The abundant (FN) in plasma is secreted by hepatocytes and lacks extra-domain III (ED) segments, refers to plasma fibronectin (pFn) (Kanters *et al.*, 2001). However, fibronectin found in sub-endothelial and connective tissue matrices are produced locally by endothelial cells and fibroblasts and mostly contain an extra type III structural domain ED-A and is called cellular fibronectin (cFn). It's Usually made up of <2% of the total FN in plasma (Kanters *et al.*, 2001).

Fibronectin: Fibronectins (FNs) are a family of large molecular weight (\approx 440 KDa) dimeric glycoproteins involved in a variety of biological processes (Osawa *et al.*, 2017). The protein is composed of similar but not identical polypeptides, which share a common molecular organization (Cabello *et al.*, 2018). These glycoproteins are found in plasma, in extracellular matrix and cell surfaces (Hamidi and Ivaska, 2017). Fibronectin is one of the most important extracellular matrix proteins that mediate a number of functions in basement membranes (Klemis *et al.*, 2017). It is a major extracellular matrix component closely associated with trabecular cells and capable of interacting with integrins and syndecans (Filla *et al.*, 2017). Fibronectin has been found in the juxtacanalicular region and along the inner wall of Schlemm's canal (Vranka and Acott, 2017). It is also present as a soluble protein in aqueous humor (Yuksel *et al.*, 2017). Granulosa cells in culture secrete high amounts of Fibronectin, and it is one of the major glycoproteins in follicular fluid as well as being a major component of extracellular matrix (Zeebaree *et al.*, 2018). It is an important matrix constituent in cartilage, and its content is markedly elevated in articular cartilage lesions within osteoarthritic joints (Mehta *et al.*, 2018). Fibronectin is a major secretory product of lung fibroblasts which represents 35 - 40% of the parenchymal cell population of normal lung (Jun and Lau, 2018). Fibronectin is also produced by alveolar macrophages, a cell type representing 90% of the inflammatory and immune cells of the lower respiratory tract (Jun and Lau, 2018). Fibronectin is found in plasma, biological fluids, loose connective tissues and some basal lamina (Cappellari *et al.*, 2016). It is synthesized by a variety of cells including liver hepatocytes, lung fibroblasts, endothelial cells, macrophages and Type 2 alveolar epithelial cells (Klemis *et al.*, 2017). Fibronectin has numerous biological functions relating to specific binding domains for cells and for extra cellular macromolecules including collagens (Cappellari *et al.*, 2016).

As a major protein in the extracellular matrix, Fibronectin provides positional assistance to help direct a wide variety of biological processes including cell migration, cell differentiation and apoptosis (Alghadir *et al.*, 2016). These biological activities of Fibronectin are mediated through interactions with various members of integrin family and cell surface proteoglycans (Kanters *et al.*, 2001). Also, these activities are contained within discrete structural domains and that neighbouring domains within Fibronectin can either suppress or enhance a particular activity (Klemis *et al.*, 2017).

Fibronectin modules: The amino acid sequence of fibronectin reveals three types of internally homologous repeats or modules separated by (usually) short connecting sequences (Sankiewicz *et al.*, 2018). There are 12 type I, 2 type II and 15 type III modules, also referred to as fibronectin I, II and III. Each module constitutes an independently folded unit, often referred to as a domain but not to be confused with "functional domains" that frequently contain more than one module

(Sankiewicz *et al.*, 2018). Modules homologous to those in fibronectin are also found in other proteins. The type II which is the most ubiquitous of all modules, is found in about 2 % of animal proteins and occasionally in plants (Eimont *et al.*, 2018). Amino acid sequences of fibronectin modules are highly conserved, differing from their counterparts in other species by only a few percent (Alghadir *et al.*, 2016). The similarity between different modules of the same type within a given protein is much less. All three modules contain several conserved core residues. Type I and II modules each contain four conserved cysteines constituting two disulfide bonds that are crucial for stability and function (Becke *et al.*, 2018). Type III is devoid of disulfide bond but III- 7 and III-15 each contain an unpaired buried cysteine (Zollinger and Smith, 2017). Fibronectin modules fold independently and this can exist in isolation from their neighbors. This was demonstrated in a few cases with proteolytic fragments that were small enough to contain only a single module and which, upon exposure to heat or chemical denaturants, underwent reversible unfolding (Becke *et al.*, 2018).

Forms of fibronectin: Fibronectins exist in two forms; circulating plasma fibronectin (soluble protomeric form) and locally synthesized cellular fibronectin (insoluble multimeric form).

Plasma fibronectin: Plasma fibronectin is synthesized by hepatocytes and circulates in the blood (Ismail *et al.*, 2018). Plasma Fibronectin has been shown to be a good indicator of pathological conditions associated with injury of the reticulo - endothelial system, radiation - induced lung injury, and in patients with Broncho-pulmonary carcinoma (CHOY *et al.*, 2017). Recent studies indicate that a significant increase in plasma Fibronectin occurs following exposure to ozone (O₃) and also that, the measurement of total plasma Fibronectin levels offers a potentially sensitive tool for detecting O₃ effects (Néri *et al.*, 2017).

Cellular fibronectin: Cellular fibronectin is synthesized by several cells including epithelial and connective tissue cells and is deposited into the extracellular matrix (Raitman *et al.*, 2017). Cellular fibronectin is a protein derived from an endothelium, it plays an important role in the assembling of sub endothelial matrix (Osawa *et al.*, 2017). Increased plasma levels of cellular fibronectin therefore, indicate loss of endothelial cell polarization or injury to blood vessels (Osawa *et al.*, 2017). Furthermore, elevated plasma levels of circulating cellular fibronectin have been described in clinical syndrome with vascular damage, although not in diabetes or atherosclerosis (Takii *et al.*, 2017). Both forms of fibronectin are encoded by a single gene, whose transcript can be alternatively spliced at three intron-exon boundaries, yielding fibronectin protein subunits containing, in the rat, the extra type IIIA (EIIIA) repeat, the extra type IIIB (EIIIB) repeat, and/or variable region (V) region. Equivalent regions are designated extra domain - A (EDA), extra domain B (EDB) and connecting segments (III CS) in humans. During splicing of fibronectin mRNA, these exons are included or excluded depending on the tissue and cell type (Huang *et al.*, 2018).

Alternative splicing of fibronectin: There are five sites of alternate splicing of fibronectin mRNA (Murphy *et al.*, 2018). The first two results in the insertion of extra type III domains, EDA and EDB after modules III- II and III - 7 respectively. These modules are virtually absent from adult tissue but are

differentially expressed during embryonic development, in malignant or injured tissue, and during angiogenesis. The inclusion of EDA renders fibronectin a better substrate for cell spreading and migration and has been used as a marker for certain types of cancer (Zhang *et al.*, 2018a). EDA - containing fibronectin is up-regulated as part of response to wound healing (Yan *et al.*, 2017). Both EDA and EDB are increased in human plasma following trauma (Bogoslovsky *et al.*, 2015). This module is reported to serve as a ligand for integrins $\alpha_9\beta_1$ and $\alpha_4\beta_1$. No specific receptors or other ligands have been identified for EDB, although it also is reported to support cell adhesion. EDB is also of interest as a marker for angiogenesis and as a potential target for tumor therapy (Sackey-Aboagye *et al.*, 2016). In plasma fibronectin mRNA, exons encoding the EDA and EDB segments are excluded, whereas the exon encoding the IIICS is either partially or completely included depending on the species. In cellular fibronectin mRNA, EDA and / or EDB exons are wholly retained and the IIICS appears to be at least partially included (Kaspar *et al.*, 2017). The third site of splicing is in the V (variable) (ITICS) region, which can be included either in its entirety or only partially, depending on the tissue. In plasma fibronectin the V region is fully incorporated into one chain but entirely absent from the other. Since the V region has at least two different integrin binding sites, its presence or absence will affect the adhesion of some types of cells. The multiple splice sites within this region produce five IIICS variants in humans namely IIICS-0, IIICS - 120, IIICS - 95, IIICS - 89, and IIICS - 64 (Scanzello *et al.*, 2015). A fourth site of splicing occurs primarily in cartilage where the dominant form of fibronectin lacks not only the entire V region but modules IIICS -15 and 1 - 10 as well. This site is located at the carboxyl termini of fibronectin (Scanzello *et al.*, 2015). A truncated single chain form of fibronectin first identified in zebra fish, has been reported in mammals, including humans (Qiao *et al.*, 2014).

Glycosylation of fibronectin: In addition to the primary mRNA transcript processing, the fibronectin molecule can also undergo post-translational modifications notably O-linked glycosylation (King *et al.*, 2017). Fibronectin is a glycoprotein sites of N - linked glycosylation include the 2nd type II module, the 8th type I module, the 3rd, 5th and 7th type III modules and the V region, which also contains O - linked carbohydrates (King *et al.*, 2017). The extent and type of glycosylation varies depending on the tissue source. Amniotic fluid fibronectin contains about twice as much carbohydrate as the plasma form and this includes the presence of poly-lactosamine in the gelatin - binding domain (Ayodele *et al.*, 2017). Fibronectin isolated from malignant human tissues, fetal tissues or placenta also contains a unique "oncofetal " epitope, involving N-Acetylgalactosaminylation of a threonine in the V region, that" is recognized by the monoclonal antibody FDC- 6 and serves as the basis for a widely used gynecological test to predict pre-term delivery (Kumra and Reinhardt, 2016). The actual function of the various carbohydrate groups on fibronectin, beyond protecting certain domains from proteolysis or stabilizing them toward heat remains unclear. Thus, differential pre-mRNA processing and post-translational modification can lead to 20 different fibronectin isoforms, Usually less than 2% of total fibronectin in plasma is derived of cellular fibronectin (Stewart and O'connor, 2015).

Functions of fibronectin: Fibronectin has numerous biological functions relating to specific binding domains for cells and for extracellular macromolecules including collagens

(Prakash *et al.*, 2015). It plays key roles in various cellular events that include cell migration and proliferation of fibroblasts and endothelial cells (Grigoriou *et al.*, 2017). Physiologically, fibronectin plays an important role in cell adhesion, motility and tissue repair among others (Lenselink, 2015). However, its overproduction may decrease motility and replication of many cells including endothelial cells (Viola *et al.*, 2017). This dimeric glycoprotein has been implicated in a variety of biological processes, including tissue remodeling during wound healing and embryonic development (Grigoriou *et al.*, 2017). Fibronectins influence these processes by affecting cell adhesion and extracellular matrix assembly (Eimont *et al.*, 2018). Fibronectins promotes cell-cell and cell matrix interactions and thus plays a role in tissue construction and reconstruction (Ayodele *et al.*, 2017). This glycoprotein is also important in such diverse activities as cell adhesion, cell migration, cellular differentiation, blood clotting, opsonization, wound healing and neoplastic transformation (Grigoriou *et al.*, 2017). It is also involved in many complex functions, some of which include the enhancement of the binding of a lymphokine to macrophage and the expression of iron receptors on macrophages. It also increases adherence and chemotaxis of phagocytes and helps maintain the oxidative bactericidal capacity of macrophages (Akenami *et al.*, 1997).

Functional domains of fibronectin: The diverse recognition functions of fibronectin are located on distinct fragments or domains, many of which have been expressed on recombinant form or isolated from proteolytic digest with retention of specific binding properties (Piprek *et al.*, 2018). Digestion of fibronectin with any of a variety of proteases generates a collection of fragments that are sometimes referred to as functional domains because they retained the ability to interact with other macromolecules (Piprek *et al.*, 2018). When thermolysin is used, one obtained an N-terminal 29 KDa Fib-1/Hep-1 fragment (1₁₋₅) that binds to fibrin, heparin and some bacteria, and is important for fibronectin matrix assembly. This is followed by a 42 KDa gelatin binding fragment (GBF, I₆II₁₋₂I₇₋₉) that binds to denatured collagen (gelatin) and also to tissue transglutaminase. Then a small ≈ 9 KDa fragment (III-I) that binds weakly to heparin and is thought to be important for self-association and fibril formation. Then the large 110kDa central cell- binding fragment (III₂₋₁₀) contains the famous tripeptide RGD that was the first integrin recognition sequence to be identified. This is followed by either a 30 or 40 kDa Hep-2 fragment (III₁₄ or III₁₂₋₁₅) contains the highest affinity heparin- bindings site. Finally, from the C-terminus one obtains a second fibrin-binding fragment, 19-kDa Fib-2 (1₁₀₋₁₂). The variably spliced V region, also referred to as the type III connecting strand contains two sites for recognition by integrin $4\beta 1$; however, it is not recovered intact from proteolytic digests. Its presence in only one chain of plasma fibronectin and its vulnerability to proteolysis digest accounts for the occurrence of 30 and 40kDa Hep-2 fragments as well as a small 8kDa fragment consisting of module III-15 whose function is unknown (Hecker *et al.*, 2018). The biological activities of fibronectin are mediated via interactions with various members of the integrin family (Park *et al.*, 2018). Integrins are transmembrane receptors composed of an α - and β - subunits covalently associated into a heterodimer (Yang *et al.*, 2018). There are more than 20 different integrin receptors each having their own ligand specificity. In general, however, they bind various extracellular matrix proteins usually through a specific ARG sequence and cell surface proteoglycans.

It has been shown for some time that these biological activities of fibronectin are contained within discrete, structural domains and that neighbouring domains within fibronectin can either suppress or enhance a particular activity (Yang *et al.*, 2018)

Fibronectin and pathological states: Fibronectin has been implicated in many pathological states (Kumar *et al.*, 2014). These include cutaneous wounds, tumour metastasis, rheumatoid arthritis; cardiac allograft rejection and liver fibrosis (Piperigkou *et al.*, 2018). Fibronectin is also markedly elevated in articular cartilage lesions within osteoarthritic joints (Tanoue *et al.*, 2018). Fetal tissues and tumours express a higher percentage of fibronectins with ED-A and ED-B repeats, but in adult life, they are particularly absent (Kumra and Reinhardt, 2016). The extra type III domain fibronectin is prominent during the injury response of adult tissues and mediates important early events in the response (Wang *et al.*, 2016). This form is particularly apparent in acute liver injury, where it has been shown that sinusoidal endothelial cells produce ED-A fibronectin (Kornblihtt *et al.*, 1996). Expression of the ED-A and ED-B isoforms are also increased during wound healing (Maione *et al.*, 2016). Fibronectin synthesis is increased in macrophages and polymorphonuclear leukocytes isolated from inflammatory foci (Akenami *et al.*, 1997). One of the most important extra cellular matrix proteins that are over expressed in retinas, glomeruli and the hearts in diabetes is fibronectin (Hu *et al.*, 2017). As reported by Doddapattar *et al.* (2018), high glucose concentrations increase a number of extra cellular matrix related genes, including fibronectin.

Vascular smooth muscle cells, cultured under hyperglycaemic conditions proliferate at a significantly faster rate than those cultured under normal glucose conditions (Xi *et al.*, 2019). High glucose increases the de novo synthesis of the protein Kinase-C activator, diacylglycerol (Xi *et al.*, 2019). Thus, one hypothesized mechanism by which High glucose induces VSMC proliferation is through the chronic activation of one or more isoforms of protein kinase – C (Xi *et al.*, 2019). Also, Chen and colleagues in 2003 used human macrovascular and microvascular endothelial cell lines to evaluate the key molecular signaling events involved in high glucose -induced fibronectin over-expression. This expression was shown to be dependent on endogenous endothelin receptor - mediated signaling. They also examined the roles played by protein kinase-C and the transcription factors, nuclear factor κ B and activating protein-I with respect to such changes (Chen *et al.*, 2003). Protein Kinase C (PKC), a family of phospholipids dependent serine/ threonine protein kinases, plays a central role in the growth factor signal transduction pathway and regulates a wide variety of cellular functions, including cell proliferation, differentiation and cell growth (Fu *et al.*, 2017). PKC represents a family of II isoenzymes that have been categorized into three groups: group A or cPKCs (α , β I, β II and γ), group B or nPKCs (α , ε , η , θ) and group C or aPKCs (ζ and λ). In addition, PKC resembles nPKCs structurally but aPKCs functionally (Isakov, 2017). While cPKCs require Ca^{2+} - and diacylglycerol and phorbosters for their activities, PKCs are Ca^{2+} insensitive, but they are still activated by DAG or phorbol esters, aPKCs are independent of both Ca^{2+} and DAG/phorbol esters. The PKC isoenzymes differ in biochemical properties, tissue specific distribution and intracellular localization (Frank *et al.*, 2016). Increased activity of PKC in high glucose could be explained by the de novo synthesis of DAG via the polyol pathway - dependent increase in cytosolic $NADH/NAD^+$ ratio (Frank *et al.*, 2016).

Transforming Growth factor B (TGF - B) is a key cytokine for matrix protein production (Roy *et al.*, 2015). TGF- β a multifunctional cytokine that operates in either a paracrine or autocrine manner stimulates fibronectin synthesis and regulates production of the EIIA - fibronectin isoform by acting on fibronectin pre - mRNA (Duran-Salgado and Rubio-Guerra, 2014). TGF - β is derived from a number of cell types including platelets wound macro phages and fibroblasts myofibroblasts, as well as injured proximal tubular epithelial cells (Moses *et al.*, 2016). High glucose has been shown to induce coordinated increases in TGF - β protein, the TGF Type 2 receptor, and the extracellular matrix protein fibronectin). High glucose has also been shown to trigger a series of events including protein kinase - C and mitogen - activated protein (MAP) kinase activation, elevation of reactive oxygen species, and increased expression of c - fos and c-jun proto-oncogenes which form a heterodimer, AP - I, that activates the TGF - β gene promoter (Yang *et al.*, 2014). Also, fibronectin splicing is known to be regulated in part by TGF - β . In the setting of injury in vivo, TGF - β overrides the promoter - dependence of fibronectin splicing in normal cells. This suggest that TGF - β , modifies the spliceosome if not through its known signaling intermediates, then it's through the products of genes regulated by this cytokine (Wang *et al.*, 2017).

DIABETES MELLITUS: Diabetes mellitus is one of the most common metabolic diseases in which either the hormone insulin is lacking or the body's cells are insensitive to insulin effects. The multi - system effects of diabetes such as retinopathy, nephropathy neuropathy and cardiovascular diseases are considered important impinging on the public health (Gingras *et al.*, 2017). A persistently increased blood glucose level characterizes diabetes. The diagnosis of diabetes mellitus is established by a raised blood glucose level: the venous whole blood glucose must be over 10mmol/l in samples taken at random or 2 hours after a 75g oral glucose load (de Oliveira *et al.*, 2018). Diabetes mellitus is generally categorized into two types of disease. Type I diabetes mellitus (also known as insulin - dependent diabetes mellitus (IDDM) is characterized by an absolute deficiency of insulin. In some clinical studies, type I diabetes mellitus is assumed when diagnosis occurs at or before the age of 30 in patients who require continuous insulin use. Type 2 diabetes mellitus (also known as non-insulin -dependent diabetes mellitus (NIDDM) refers to patients with diabetes mellitus characterized by insulin resistance or a state of relative insulin deficiency. Clinical studies will often use diabetes onset after age of 30 years as an operational criterion for type 2 diabetes mellitus (Gospin *et al.*, 2017). These two differ in their clinical characteristics, etiological and pathophysiologic basis for diseases. The main clinical difference is their propensity to develop diabetic ketoacidosis in the basal metabolic state; insulin is required in type I to prevent ketoacidosis, whereas in Type 2, ketoacidosis is unlikely even' when the glycaemic control is poor. Typically, type 1 presents acutely and manifests the typical symptoms of polyphagia, polydipsia and polyuria (Gospin *et al.*, 2017). In contrast, Type 2 is insidious and may be present for years before being diagnosed. Approximately, a good percentage of all diagnosed cases of diabetes mellitus is Type 2 and may be as many undiagnosed cases of Type 2 as diagnosed cases (Matsugasumi *et al.*, 2018).

Prevalence of diabetes mellitus: The number of people with diabetes mellitus is increasing due to population growth,

aging, urbanization, and increasing prevalence of obesity and physical inactivity (Gjesing *et al.*, 2017). The adoption of western lifestyles is known to lead to increasing prevalence of type 2 diabetes mellitus in Africa, although epidemiological studies using standardized methods are rare (Uloko *et al.*, 2018). The prevalence of diabetes for adult worldwide was estimated to be 8.8% in 2015 (Uloko *et al.*, 2018) and 6% in Sokoto (Makusidi *et al.*, 2013). The total number of people with diabetes is projected to rise from 171 million in 2000 to 366 million in 2033 (Alexander *et al.*, 2003). The prevalence of diabetes is higher in men than women, but there are more women with diabetes than men (Rathmann and Giani, 2004). This common endocrine disease affects more than 15 million Americans and contributed to ≈47.9 billion dollars in health care expenditure in 1995 (Bouguerra *et al.*, 2007). Both type 2 diabetes mellitus and type I diabetes mellitus show a wide variation in incidence and prevalence in different populations (Bouguerra *et al.*, 2007). The prevalence of type 2 varies from close to zero in some populations to 40 - 50% in the adults of Nauru (Pacific) and in the Pima Indians (North American) (Misra *et al.*, 2010). The incidence of type I in children under 16 years range from around 301,100,000 children per year in Finland and other Northern European countries, to less than 111,00,000 per year in Japanese children (Dabelea *et al.*, 2014). Many genetic and environmental factors combine to produce this variation. Among the suggestions made for Type 2 are the thrifty genotype and more recently the thrifty phenotype hypotheses of Nee! and of Hales and Barker respectively (Hales and Barker, 2001). Genetic and environmental factors in type I diabetes mellitus in children have combined to create an apparent gradient of decreasing incidence from northern to southern European countries but with at least one local 'hot spot': The Island of Sardinia (Cucca *et al.*, 1995). The factors responsible for this pattern and the increasing incidence overtime require further studies (Cucca *et al.*, 1995)

Type 1: Insulin Dependent Diabetes Mellitus (IDDM): The aetiology and pathogenesis of Type 1 diabetes mellitus are not fully understood, but Mandrup-Poulsen grouped patients with Type 1 diabetes mellitus according to their presenting characteristics as well as immune and genetic markers into early onset, older children and adolescent and late onset (Mandrup-Poulsen, 2003). He reported that Type 1 diabetes mellitus in early childhood is characterized by abrupt onset and by high frequencies of ketoacidosis and of insulin and IA2 autoantibodies (Pociot and Lernmark, 2016). It is associated with the major histocompatibility complex heterozygosity, HLA DR 3/4 (Pociot and Lernmark, 2016). Type 1 diabetes mellitus in older children and adolescents is associated with a high frequency of islet cell cytoplasmic autoantibodies, glutamic acid decarboxylase (GAD) autoantibodies and HLA DR 3. Typical immune markers often accompany autoimmune diabetes mellitus of late onset (Leslie *et al.*, 2016). Whether these groupings represent separate diseases or simply reflect the rapidity with which β -cell destruction takes place is unclear (Asmat *et al.*, 2016). Some learn people with type 1 diabetes mellitus lack the classic manifestations of autoimmune disease, and their diabetes may therefore have a different aetiology and pathogenesis (Leslie *et al.*, 2016). β -cell destruction in immune mediated type 1 diabetes mellitus is caused by an inflammatory reaction in the islets of Langerhans, triggered by environmental factors in genetically susceptible individuals (Mandrup-Poulsen, 2017).

Aetiology: Although the incidence is highest in school children and adolescents, studies have shown that in its classic form, Type 1 diabetes mellitus may develop at any age (Newton *et al.*, 2016). Between 1960 - 1989, steady temporary increases were reported in North America, Northern Europe, Japan and New Zealand (A.D.A., 2015). Data from Finland and United Kingdom show that the growing incidence is mainly explained by an increase in the frequency of onset in children under 5 years (A.D.A., 2016). This finding points to changes in major environmental aetiological factors early in life, such as viral infections and nutritional factors (Enander *et al.*, 2018).

Viral infections: Studies had shown that mothers whose children became diabetic before age 13 had increased titers of low avidity antibody to enteroviruses such as the Coxsackie β group (indicating recent viral infection) compared with mothers whose children were not diabetic by age 15 (Enander *et al.*, 2018). Immune cross reactivity to similar sequences in the Coxsackie β virus and the putative β -cell autoantigen, glutamic acid decarboxylase, has been found, suggesting that molecular mimicry may be the underlying mechanism (Enander *et al.*, 2018).

Nutritional factors : Early exposure to cow's milk proteins is associated with an increased risk of childhood diabetes (Lamb *et al.*, 2015). Molecular mimicry, between bovine serum albumin and ICA 69, another putative β -cell autoantigen, has been suggested (Dubridge *et al.*, 2019). However, use of bovine serum albumin to induce tolerance in animal models of spontaneous type 1 diabetes mellitus failed to show protection against diabetes despite tolerance to the bovine serum albumin (Li *et al.*, 2017a). Also, of interest is formula milks based on cows' milk and on hydrolysed casein contain bovine insulin. Children fed with cows' milk formula had highertiters or IgG antibodies to bovine insulin that cross reacted with human insulin, and the presence of these autoantibodies correlated with the presence of autoantibodies to human insulin (Gil *et al.*, 2017). These findings may explain the high incidence of insulin autoantibodies in young children and raise the possibility that sensitization to bovine insulin could affect immune system auto aggression towards β -cells in people who are genetically predisposed (Gil *et al.*, 2017). Finally, excess intrauterine growth has been reported as an independent risk factor for type 1 diabetes mellitus, but the mechanism remains to be explain (Sauder *et al.*, 2017).

In some genetically susceptible individuals, this immune process can persist In association with chronic progressive beta cell destruction over many months, even years, and lead to IDDM, but in others it may remit spontaneously without diabetes developing (Gil *et al.*, 2017). At diagnosis, about 80% of islets contain no beta cells and the islets may be heavily infiltrated with lymphocytes. There is no evidence that the exocrine pancreatic cells or the other islet cells are involved in this destructive process. The limited secretion of insulin by patients with IDDM results in them being prone to ketoneogenesis (Gormsen *et al.*, 2017). In the absence of insulin treatment such patients will die in diabetic ketoacidosis (Gormsen *et al.*, 2017).

Type 2: Non - Insulin Dependent Diabetes (NIDDM): Type 2 diabetes affects at least 2% of the population and no race is immune from the disease; among the Pima Indians of Arizona and Nauruan from Polynesia, half the adult population is diabetic. The prevalence of type 2 diabetes increases with

increasing age: in the elderly the prevalence can be striking, reaching 45% in men aged 75 - 79 years in east Finland (Rewers and Ludvigsson, 2016). In part this increase may be due to an age-related deterioration in glucose tolerance. Thus, while fasting blood glucose levels remain fairly constant with age the glucose level 2 hours after oral glucose rises steadily. In the absence of prospective studies this age-related effect is ignored for the purposes of defining the disease (Rewers and Ludvigsson, 2016).

Pathogenesis: The islets of Langerhans in-patient with type 2 appear normal except for amyloid deposits and a reduction in the beta cell mass to about 60% of normal (Chiang *et al.*, 2017). In general, the secreted insulin has a normal structure though there is a tendency to secrete a relative excess of pro-insulin. The disease is characterized by a decreased beta cell, secretory capacity, insulin resistance and hepatic glucose overproduction (A.D.A., 2010). The onset of the diabetes is insidious, probably occurring several years before the clinical diagnosis (Alberti and Zimmet, 1998). Since hyperglycaemia itself can induce beta cell dysfunction and insulin insensitivity, it has proved impossible to distinguish primary changes leading to diabetes from those, which are secondary to the disease (Tilg *et al.*, 2017). To address this problem, studies have been performed on non-diabetic children of diabetic patients. These children show fasting hyperglycaemia, impaired glucose tolerance, decreased glucose clearance, fasting hyperinsulinaemia, either decreased or increased insulin response to glucose, and impaired insulin-mediated glucose disposal due to reduced non-oxidative glucose metabolism (DeFronzo *et al.*, 2015). It remains to be determined whether these changes presage diabetes. The one study which followed patients prospectively found that offspring who developed diabetes, as compared with those who do not, were initially more obese and had decreased glucose tolerance and glucose clearance, fasting hyperinsulinaemia and increased second phase insulin responses (Koivusalo *et al.*, 2016). The development of type 2 was both preceded and predicted by defects in insulin-dependent and insulin-independent glucose uptake changes, which could precede the onset of hyperglycaemia and diabetes by more than a decade (Zheng *et al.*, 2018). These observations are consistent with the finding that initial lesion in type 2 is due to peripheral insulin resistance not beta cell dysfunction (Zheng *et al.*, 2018). However, it is widely believed that neither abnormal insulin secretion nor abnormal sensitivity to insulin alone can explain the glucose intolerance of type 2 diabetes (Zheng *et al.*, 2018).

Aetiology: The most powerful evidence that type 2 is predominantly inherited comes from the study of identical twins of diabetic patients concordant for type 2 diabetes (Willemsen *et al.*, 2015). Evidence suggest that some individuals with maturity onset diabetes in the young (MODY) have a defect (both missense and nonsense mutations have been reported) in their glucokinase gene promoter region which might account for up to 40% of these cases. Rare genetic defects in the insulin gene and in the insulin receptor have also been described which may cause diabetes (Willemsen *et al.*, 2015). The estimated heritability of type 2 is about 82%. In assessing the role of the environment in causing NIDDM, it is important to understand that heritability is not an invariant index of a genetic influence. It describes the genetic effect under particular environmental conditions (DeFronzo *et al.*, 2015).

Different estimates of heritability might be obtained if twins were studied in different environments. A number of non-genetic factors have been implicated in the aetiology of this disease, including nutrition, obesity, and aging and reduced exercise (van Dongen *et al.*, 2015).

Nutrition: The incidence of type 2 decreases during food shortage (Ragnarson Tennvall and Apelqvist, 2004). However, high carbohydrate diets improve insulin sensitivity in both normal, and Type 2 subjects through physiological adaptation to an altered fuel supply and not reversal of a pathological process (Wilding, 2014). The change from hunter-gatherer to a modern diet may be responsible, in part, for the virtual epidemic of diabetes in migrant populations and previously isolated communities. At present, however, there is no direct evidence that dietary factors cause type 2 diabetes although they may influence rate of progression to clinical symptoms. Studies do indicate a relationship between low birth weight and impaired glucose tolerance in later life (Crusell *et al.*, 2017). Fetal malnutrition may contribute to the later development of type 2 diabetes mellitus by impairing early fetal growth and reducing the β -cell mass. This might limit β -cell function in adult life and hamper the ability of the β -cell to compensate for insulin resistance caused by obesity, pregnancy, or drugs such as glucocorticoids (Crusell *et al.*, 2017). Support for this view shows that, low birth weight in adult first degree relatives of patient, with type 2 diabetes mellitus is associated with impaired β -cell function, and monozygotic twins who became concordant for type 2 diabetes mellitus after 60 years of follow up were growth retarded at birth compared with discordant twins (Beaumont *et al.*, 2017). Furthermore, low birth weight has been found to predispose to obesity and hypertension, suggesting that intrauterine growth retardation is a risk factor for several features of metabolic syndrome (van Dijk *et al.*, 2015). These data show that increased focus on early fetal growth and nutrition may prevent type 2 diabetes mellitus in the coming generation (van Dijk *et al.*, 2015).

Obesity: Obesity is a major factor and can potentiate type 2 diabetes mellitus in genetically susceptible individuals (Costanzo *et al.*, 2015). Offspring of type 2 diabetes mellitus themselves are more likely to be obese when young. In one study, those subjects with low birth weight who became obese were particularly prone to diabetes mellitus (Duque-Guimarães and Ozanne, 2017). Individual with upper body obesity are also particularly high risk of type 2 diabetes mellitus (Kratochvilova *et al.*, 2019). Nevertheless, there are no differences in the relative contributions of decreased insulin secretion and decreased insulin sensitivity between obese and lean type 2 diabetes mellitus subjects (Sattar and Gill, 2014; Kratochvilova *et al.*, 2019).

Aging and Exercise: Ageing is not associated with decreased insulin secretion nor with decreased insulin sensitivity in physically active subjects (Lalia *et al.*, 2016). However, lack of exercise in the elderly may hasten the appearance of hyperglycaemia. Prospective studies of populations at risk of diabetes suggest that physical activity protects against the development of type 2 diabetes mellitus (Shih *et al.*, 2018).

Genetics of diabetes: Two different methodological approaches have been taken to clarify the genetic bases of insulin dependent and non-insulin dependent diabetes (Burgio *et al.*, 2015).

The random marker approach uses information from the human genome project on the location of so called chromosomal "tags" spread all over the human genome, and the segregation of these tags with diabetes in genetically informative families. This method identifies chromosomal regions that may contain genes predisposing to disease; subsequent analysis is necessary to identify the structural genes of pathogenetic relevance. A complementary method is the candidate' gene approach (Mandrup-Poulsen, 2017). Here genes encoding for proteins believed to have a role in the pathogenesis of a disease (based on formulated pathogenetic model) are searched for variations associated with the disease in case-control studies or linked to 'the disease in family studies, or both (Mandrup-Poulsen, 2017).

Type 1 diabetes mellitus: In insulin dependent diabetes mellitus, the random marker approach has identified 18 chromosomal regions that show evidence of linkage to insulin dependent diabetes mellitus (Gauguier *et al.*, 1996). Formula linkage has been shown only for the major histocompatibility complex class II region, a variable number of tandem repeats 5' to the insulin gene and the CTLA-4 gene, coding for the ligand B7, an important accessory molecule for T cell activation (Nisticò *et al.*, 1996). Studies comparing different racial groups have shown evidence of appreciable genetic heterogeneity between ethnic groups. Much work is needed before the complex genetic susceptibility of insulin dependent diabetes mellitus is understood and the functional role of the products of these genes in pathogenesis is clarified (Peres *et al.*, 2017).

Type 2 Diabetes mellitus: More than 250 candidate genes have been tested for association with or linkage to non-insulin dependent diabetes mellitus, but none has shown consistent results in different study populations (Horikawa *et al.*, 2000). Most success for the candidate gene approach has been provided in the study of maturity onset diabetes in young people, an autosomal dominant form of early onset non-insulin dependent diabetes mellitus (Yamagata *et al.*, 1996). Although maturity onset diabetes in young people may seem to be a homogeneous phenotype, four genetically and pathogenetically distinct forms have been described. The first form is caused by a mutation in the transcription factor hepatic nuclear factor 4 (Yamagata *et al.*, 1996), the second results from different mutations in the glucokinase gene affecting β cell glucose sensing (A.D.A., 2014) the third is caused by mutations in the gene encoding for hepatic nuclear factor 1 and the fourth results from mutation of the insulin promoter factor-1. The precise role of the hepatic nuclear factor 4 and 1 mutations is not known, but it is suspected that mutations in these transcription factors affect the β cell response to glucose (A.D.A., 2017). These genes account for a small percentage of non-insulin dependent diabetes and underline the genetic complexity behind apparently homogeneous diabetic phenotypes (A.D.A., 2017). However, these discoveries may lead to new drugs that modify the β cell response to glucose, and to the development of pharmacological principles for treating common non-insulin dependent diabetes. Linkage studies in non-insulin dependent diabetes are hampered by a general lack of parents of patients with diabetes of adult onset. Very large numbers of affected sibling pairs are needed for linkage studies, and this calls for international collaboration (Mandrup-Poulsen, 2017).

Metabolic consequences of hyperglycaemia: A common health aim is to reduce the morbidity and mortality associated with diabetes (Ottosson-Laakso *et al.*, 2017). Risk factors for macro vascular disease are well established in the non - diabetic population and include hypercholesterolemia, increased plasma fibrinogen, smoking, obesity and hypertension (Zelihic *et al.*, 2015). The evidence is that in the diabetic population the same risk factors operate but, if anything, diabetes has an additive effect with them (Zelihic *et al.*, 2015). In addition, in type 2 diabetes mellitus there is tendency for some of these risk factors to aggregate; these patients are particularly at risk of obesity, dyslipidaemia and hypertension (Corsino *et al.*, 2017). A common feature of these changes is their association with insensitivity to insulin, an observation which has led to the proposal that insulin resistance is the single unifying factor causing an excess risk of macro vascular disease has been called the New World syndrome, the metabolic syndrome or syndrome X (Furukawa *et al.*, 2017).

The cause of microvascular disease is not clearly defined but the belief is that hyperglycaemia is a major factor (Packer, 2018). Microvascular-complications are not simply genetically determined since the non - diabetic twins of diabetic patients do not get them. For complications to develop, hyperglycaemia must be present (Packer, 2018). The cause of hyperglycaemia is irrelevant since microvascular complications are a feature of all types of diabetes mellitus (Magri *et al.*, 2018). As diabetes is defined by hyperglycaemia, it is reasonable to anticipate that; these microvascular complications should result from this hyperglycaemia. It is clear that the risk of diabetic complications is related to the duration of the disease (Targher *et al.*, 2018). A number of studies have demonstrated a relationship between the level of blood glucose and the risk of developing complications (Targher *et al.*, 2018). Thus, in one study, average blood glucose more than 50% above the normal range was associated with a 49% risk of developing severe retinopathy at 14 years: in contrast, this risk was only 5% in those patients with blood glucose levels close to the normal range (Cooper *et al.*, 2017).

Pregnancy: It is known that hyperglycaemia can influence foetal development (Berry *et al.*, 2016). The incidence of major and minor congenital anomalies in children of patients with diabetes mellitus is between 6% and 9% that is up to 3 times greater than in the general population (Basu and Garg, 2018). The most prevalent congenital anomalies in children of diabetic patients include caudal regression syndrome, neural tube defects and cardiac anomalies (Basu and Garg, 2018). The excess in malformations is confined to patients whose diabetes mellitus antedates their pregnancy (Ornoy *et al.*, 2015). In addition, the malformations arise from developmental changes likely to have occurred before the seventh week of gestation (Monteiro *et al.*, 2016). It was proposed that the excess congenital anomalies in children of patients with diabetes mellitus were due to hyperglycaemia in early fetal life (Monteiro *et al.*, 2016). Measuring glycated haemoglobin, an index of blood glucose levels, over the previous two months, tested this hypothesis (Ribeiro *et al.*, 2018). Children of patients with high glycated haemoglobin levels had a striking excess of congenital anomalies, which reached 22% if the glycated haemoglobin was greater than 10% (Ribeiro *et al.*, 2018). The risk of major malformations can be reduced to non - diabetic levels if the diabetic mother is treated to obtain normal glycated haemoglobin levels before

conception (Mañé *et al.*, 2019). The mechanism of this embryopathy is not clear. Pregnancy counselling is now routine in diabetic clinics and patients are advised to obtain near normal fasting plasma glucose levels before conception (Mañé *et al.*, 2019).

Mechanisms of diabetes complications: Exposure to hyperglycaemia can cause acute reversible metabolic changes and, if prolonged, cumulative irreversible changes (Gioacchini *et al.*, 2018). Broad mechanisms have been described for glucose-induced damage (Habib and Ali, 2018). First, glucose and other sugars can bond with any exposed lysine residues or (in the case of haemoglobin) valine, or any protein (Góralczyk *et al.*, 2016). This process of glycation can alter the structure and function of protein (Giri *et al.*, 2018). Further changes can lead to glycation products with extensive cross-linkage called advanced glycation end products - an irreversible change (Giri *et al.*, 2018). These molecules may lead to the production of free oxygen radicals, which could themselves cause tissue damage (Zhang *et al.*, 2018b). The second mechanism is the production of excess sorbitol through a normally redundant pathway involving the enzyme aldose reductase (Negahdar *et al.*, 2015). Sorbitol cannot readily leave a cell and accumulation of the alcohol sugar could lead to osmotically driven over hydration of the tissue and damage (Negahdar *et al.*, 2015).

The third mechanism involves the direct competition between glucose and myoinositol (Ivanov Kavkova *et al.*, 2019). Myoinositol is an important substrate in cellular energy production, and its structure is very similar to that of glucose. Excess glucose can therefore compete for myoinositol uptake by a cell, leading to myoinositol depletion (Zhan *et al.*, 2015). Although, diabetes control and complications trial has identified, hyperglycaemia as a significant risk factor for the development of diabetic complications (Wessells *et al.*, 2018). The full spectrum of the pathophysiological mechanisms of chronic diabetic complications has not been thoroughly elucidated. Some equally tenable hypotheses for the origin of complications are oxidative stress damage (Kortenkamp *et al.*, 2011; Wessells *et al.*, 2018), advanced glycation end products (AGEs) hypotheses (King and Brownlee, 1996), aldose reductase pathway (Yabuuchi *et al.*, 1995) reductive stress (pseudohypoxia) (Oldham *et al.*, 2015), true hypoxia (Sansome *et al.*, 2001), carbonyl stress (Takeda *et al.*, 2015), altered lipoprotein metabolism (Rye and Barter, 2014), increased protein kinase - c activity (Newton *et al.*, 2014), and altered growth factors and- cytokine activities (Sun *et al.*, 2017). The various hypotheses overlap and intersect with one another; AGE formation and altered polyol pathway activity may lead to oxidative stress, oxidative stress may accelerate AGE formation, reductive stress may lead to activation of protein Kinase - C, AGEs may induce growth factors and cytokine production and so on (Baynes and Thorpe, 1999). Among Chronic Complications of DM as a result to endothelial injury include; Nephropathy, Retinopathy, Diabetic Neuropathy, Peripheral vascular disease, and Coronary artery diseases Change in endothelial function is considered an early pivotal step in the development of vascular disease (Kanters *et al.*, 2001). The role of circulating (cFn) in the diagnosis of vascular endothelial injury in rheumatoid vasculitis, pre-eclamptic women, have been documented (Kanters *et al.*, 2001). Vessel wall damage with characteristic endothelial extracellular matrix changes is also found in subjects with diabetes.

Several reports have implicated DM in the pathogenesis of vessel wall damage with extracellular matrix changes (Peters *et al.*, 1989).

Justification: Cellular FN level assessment could be useful as an adjuvant in predicting early vascular complications of DM. The paucity of data on circulating (cFn) in only DM in Nigeria, is the main thrust for this study. The anticipated data could provide useful adjuvant in predicting early vascular complication of DM patients.

Aim: The aim of the present study was to determine the levels of plasma cFn among DM patients attending diabetic clinic in Specialist Hospital, Sokoto.

Objectives: To determine the levels of plasma cellular fibronectin (cFn), nitric oxide (NO), glycated haemoglobin (HbA1c), von Willebrand factor (VWF) and TNF α in DM patients and apparently healthy individuals (controls). To compare levels of plasma cFn, NO, HbA1c, VWF and TNF α among DM patients and controls. To determine effects of duration treatment of DM on cFn, NO, HbA1c, VWF and TNF α level. To compare levels of plasma cFn, NO, HbA1c, VWF and TNF α among DM patients with/ without complications.

MATERIALS AND METHODS

The research was carried out in Sokoto State, basically in the Department of Chemical Pathology and Immunology, College of Health Sciences (CHS) and Department of Medicine Specialist Hospital Sokoto. A total of 200 participants were consecutively selected for the study. Only diabetic and apparently healthy individuals who fulfilled the inclusion criteria and agreed to participate in the study were selected. Diabetic subjects were selected from diabetic clinics in the Department of Medicine Specialist Hospital, Sokoto. Preliminary information's such as age, sex, height, weight of the patients, duration of the disease and medications were obtained using a questionnaire. Patients that were only dieting as means of diabetic controls were also noted. The control subjects were one hundred (100) apparently healthy individuals, of both genders. Those with history of liver diseases and cigarette smoking were excluded from the study. Both type 1 & 2 diabetic patients and apparently healthy individuals aged 18 years to 60 years were recruited into the study. Diabetic patient with hypertension, Diabetic patient with coexisting other endocrine disorders and Diabetic patient that consume alcohol were excluded from the study. Individuals who were non-diabetic and who have never had any family history of diabetes were included in the study as controls. Participants (Diabetics patients and apparently healthy controls) were fully informed, and their consent were obtained before the commencement of the research. Participants were allowed to withdraw from the study at any time and for any reason. Approval was obtained from the Ethics and Research Committee of the Specialist Hospital Sokoto. The study was a descriptive cross-sectional study, which was performed on Diabetic subjects attending Diabetic Clinic at Specialist Hospital Sokoto, for a period of 12 months. The diabetic patients were categorized into 2; group A based on duration of treatment and group B, which served as control. Group A was further subclassified into 5.

Group A1. Treatment naïve

Group A2. on treatment for less than a year (<1yr)

Group A3. on treatment for one to less than two years (1 - <2yr)

Group A4. on treatment two to less than five years (2 - <5yr)

Group A5. On treatment for five years and above (\geq 5yr). From each selected subject, a total of five milliliters (5mls) of fasting venous blood specimen were collected using a sterile syringe and needle. From the 5mls, three mls was placed in EDTA bottles for cFn, HbA1c, VWF and TNF α assay and two ml was be placed into a fluoride oxalate container for Fasting blood glucose assay. The blood sample was centrifuged to separate plasma from the cells immediately. The plasma was refrigerated at -20 $^{\circ}$ c

Statistical analysis: The data obtained were analyzed using Microsoft Office Excel 2007 and SPSS software version 20.0 of 2016. The results of plasma fasting glucose and lipid profile obtained from diabetic subjects were compared with the controls using pair two-tailed student's t-test for matched samples, while analysis of variance (ANOVA) was used to for comparisons of three (3) or more mean values of the parameters in the various groups. In each case where there was significant difference, a post-hoc analysis was carried out using Bonferroni multiple comparisons test. A p-value of less than or equal to 0.05 ($P \leq 0.05$) was considered as statistically significant.

RESULTS

A total of two hundred (200) subjects participated in this study. Of this number, 100 were diabetic patients, 45 males (45%) and 55 females (55%) with their age ranged between 20 and 60years and mean age and standard error of mean of (49.53 \pm 0.92). The remaining 100 were age and sex matched apparently healthy individual comprised of 48 males (48%) and 52 female (52%) who served as controls. The diabetic patients comprised of 4 type I diabetes and 96 type II diabetes. The diabetic patients were categorized into 3; group A based on duration of treatment, group B and group C. Group A was further subclassified into 5 and group B into 2. Group A1. Treatment naïve 10 (10%), Group A2. on treatment for less than a year (<1yr) 12 (12%), Group A3. on treatment for one to less than two years (1 - <2yr) 18 (18%), Group A4. on treatment two to less than five years (2 - <5yr) 20 (20%), Group A5. On treatment for five years and above (\geq 5yr) 40 (40%). Group B1. Diabetic with complication. Group B2. Diabetic patients without complication. Group served as control. Table; 1 shows the results of socio-demographic and diabetics related characteristics of the study subjects including marital status, tribe, and level of education. The anthropometric data of the diabetic subjects were summarized in table 2, the age, body weight, BMI and diastolic blood pressure of the diabetic were found to be similar with the control ($p > 0.05$). However, the systolic blood pressure and height of the diabetic subjects was significantly higher than the control ($p < 0.05$). The mean concentration of cellular fibronectin, glycosylated haemoglobin, nitric oxide, von Willebrand factor and tumour necrosis factor alpha were shown in table 3; There were significantly increased in the mean concentration of cellular fibronectin (13.21 \pm 1.62mg/L), mean concentration of glycosylated haemoglobin (9.47 \pm 0.31%), mean concentration of von Willebrand factor alpha (35.58 \pm 3.94ng/ml) and mean concentration tumor necrosis factor alpha (93.52 \pm 20.23) ($p < 0.05$) in diabetic subjects compared to control.

Furthermore, mean nitric oxide level was significantly higher in control compared to diabetic subjects (106.42 \pm 13.62 μ mol/L) ($p < 0.05$). There were also no significantly difference in SBP and DBP as ($p > 0.05$) in all the classes of duration of diabetes. Table 4. shows the results of plasma concentration of endothelial function biomarker in diabetic subjects according to the duration of treatment of diabetes in years. The mean concentration of plasma cellular fibronectin in group A5 (43.30 \pm 2.02mg/L) were significantly higher ($p < 0.05$) compared to group A1, A2, A3 and A4 of the diabetic subjects and also in the control subjects. The mean plasma concentration of Nitric oxide was significantly lower in group A5 of the diabetic subjects (52.42 \pm 9.21 μ mol/L) compare to group A1, A2, A3, A4 and the control subjects ($p < 0.05$). However, the plasma level of glycosylated haemoglobin (5.71 \pm 0.37%) was only significantly lower ($p < 0.05$) in the control subjects compared to all the groups in the diabetic subjects. The mean concentration of the von willebrand factor was significantly higher in group A5 (35.09 \pm 4.70ng/L) and group A2 (42.80 \pm 8.40ng/L) ($p < 0.05$) compared the controls. The mean plasma concentration of tumor necrosis factor alpha in group A5 was significantly higher compared to all the groups ($p < 0.05$).

Table 5; shows the effect of erectile dysfunction on endothelial function biomarkers in diabetic subjects. The mean concentration of cellular fibronectin in diabetic subjects with erectile dysfunction (49.00 \pm 3.76mg/L) were significantly higher ($p < 0.05$) compared to diabetic subjects with no erectile dysfunction (32.40 \pm 2.31mg/L). There were also, a significantly difference ($p < 0.05$) between the mean concentration of nitric oxide in diabetic subjects with erectile dysfunction (45.21 \pm 3.77 μ mol/L) and diabetic subjects with no erectile dysfunction (49.59 \pm 4.40 μ mol/L). There was no significantly relationship between the mean concentration of glycosylated haemoglobin in diabetic subjects with erectile dysfunction (9.84 \pm 0.53%) compared to diabetic subjects with no erectile dysfunction (10.34 \pm 0.71%). There were no significantly ($p > 0.05$) difference between the mean concentration of von willebrand factor in diabetic subjects with erectile dysfunction (42.93 \pm 8.03ng/L) compared to diabetic subjects with no erectile dysfunction (34.67 \pm 7.95ng/L). There was no significantly ($p > 0.05$) difference between the mean concentration of tumor necrosis factor alpha in diabetic subjects with erectile dysfunction (91.50 \pm 40.46ng/L) and diabetic subjects with no erectile dysfunction (52.93 \pm 13.41ng/L). Table 6; shows the effect of blurring of vision on markers used for assessing endothelial function in diabetic subjects. The mean concentration of cellular fibronectin in diabetic subjects with blurring of vision (45.52 \pm 3.32mg/L) were significantly higher ($p < 0.05$) compared to diabetic subjects with no blurring of vision (34.00 \pm 1.91mg/L). There was no significantly difference ($p > 0.05$) between the mean concentration of nitric oxide in diabetic subjects with blurred vision (53.91 \pm 3.86 μ mol/L) compared to diabetic subjects with no blurred vision (57.94 \pm 4.84 μ mol/L). There was no significantly ($p > 0.05$) relationship between the mean concentration of glycosylated haemoglobin in diabetic subjects with blurred vision (9.76 \pm 0.44%) compared to diabetic subjects with no blurring of vision (9.37 \pm 0.39%). There were no significantly ($p > 0.05$) difference between the mean concentration of von willebrand factor in diabetic subjects with blurred vision (32.17 \pm 6.45ng/L) compared to diabetic subjects with no blurred vision (36.72 \pm 4.81ng/L).

Table 1. Socio-Demographic Characteristics of the Diabetic Subjects

Characteristics	Number of Subjects	Percentage (%)
Marital Status	100	100
Married	79	
Single	1	
Widowed	15	
Divorced	5	
Tribe	100	
Hausa	77	
Fulani	17	
Yoruba	2	
Igbo	2	
Other	2	
Level of education	100	
Primary School Cert.	10	
Secondary School Cert.	9	
Tertiary	12	
Non-formal/ Quranic Sch.	69	
Duration of DM	100	
1-11Months	15	
12-23Months	13	
24-59Months	24	
60 Months and above	48	
Types of DM	100	
Type 1	4	
Type 2	96	
Duration of treating DM	100	
Treatment naïve	10	
Less than a year	12	
1yr – less than 2yrs	18	
2yrs – Less than 5yrs	20	
5years and above	40	

Majority of the diabetic subjects in the population had type II diabetes mellitus (96%),

Table 2. Anthropometric Data of the Diabetic Subjects (Mean ± SEM)

Characteristics	Group A Diabetic Patients (n=100)	Group B Controls (n=100)	P value
Male	45	48	
Female	55	52	
Age (Years)	49.53 ± 0.92	45.9 ± 1.48	0.457
Sbp (mmHg)	115.59±0.84 ^b	111.30±1.54 ^a	0.009
Dbp (mmHg)	74.24±0.77	73.24±1.10	0.460
Body Weight (Kg)	68.11±1.70	65.98±2.62	0.485
Height(m)	1.62±0.01 ^b	1.72±0.01 ^a	0.000
BMI(Kgm ⁻²)	27.19 ± 0.63	25.14 ± 0.82	0.056

Values are expressed as Mean ± SEM; Values of the group with superscript “a” are statistically significantly (p<0.05) and different from group A. Values of the group with superscript “b” are statistically significantly (p<0.05) and different from group B.

Table 3. Endothelial functions Biomarkers in Diabetics Subjects (Mean ± SEM)

Characteristics	Group A Diabetic Patients (n=100)	Group B Controls (n=100)	p value
Cellular Fibronectin (mg/L)	31.21 ± 1.62 ^b	13.91 ± 1.20 ^a	0.000
Glycated Haemoglobin (%)	9.47 ± 0.31 ^b	5.71 ± 0.06 ^a	0.000
Nitric Oxide (µmol/L)	56.93 ± 3.75 ^b	106.42 ± 13.62 ^a	0.000
Von willebrand Factor (ng/ml)	35.58 ± 3.94 ^b	23.56 ± 3.11 ^a	0.018
Tumor Necrosis Factor (ng/L) Alpha	93.52 ± 20.23 ^b	44.49 ± 6.48 ^a	0.045

Values are expressed as Mean ± SEM; Values of the group with superscript “a” are statistically significantly (p<0.05) and different from group A. Values of the group with superscript “b” are statistically significantly (p<0.05) and different from group B.

Table 4. Effect of treatment Duration in Diabetes Mellitus on endothelial function biomarkers on diabetic subjects

Charac Teristics	N	Fibronectin (mg/L)	Nitric Oxide (µmol/L)	Glycated Haemoglobin (%)	Von willebrand Factor(ng/L)	Tumor Necrosis Factor (ng/L)
Group A1	10	30.29±2.17 ^{eq}	64.22±14.31 ^{eq}	9.41±0.48 ^{eq}	40.73±6.90 ^{eq}	71.81±36.29 ^{eq}
Group A2	12	36.44±3.62 ^{ac}	67.39±16.49 ^{eq}	9.53±0.68 ^{eq}	42.80±8.40 ^{eq}	77.83±43.30 ^{eq}
Group A3	18	27.08±3.88 ^{ac}	57.92±17.71 ^{eq}	9.82±0.73 ^{eq}	30.98±9.03 ^{eq}	54.70±46.51 ^{eq}
Group A4	20	29.62±2.86 ^{acq}	58.89±13.03 ^{eq}	9.30±0.53 ^{eq}	34.55±6.64 ^{eq}	97.27±34.23 ^{eq}
Group A5	40	43.30±2.02 ^{abcdq}	52.42±9.21 ^{abcdq}	10.44±0.38 ^{abcdq}	35.09±4.70 ^{abcdq}	173.05±24.20 ^{abcdq}
Group C (Control)	100	13.91±1.98 ^{bcdde}	106.42±9.03 ^{bcdde}	5.71±0.37 ^{bcdde}	13.12±4.60 ^{be}	44.49±23.72 ^{bcdde}

Values are expressed as Mean ± SEM; Values of the group with superscript “a” are statistically significantly (p<0.05) and different from group A1. Values of the group with superscript “b” are statistically significantly (p<0.05) and different from group A2. Values of the group with superscript “c” are statistically significantly (p<0.05) and different from group A3. Values of the group with superscript “d” are statistically significantly (p<0.05) different from group A4. Values of the group with superscript “e” are statistically significantly (p<0.05) different from group A5. Values of the group with superscript “q” are statistically significantly (p<0.05) different from group C.

Table 5. Endothelial function biomarker in diabetes subjects with erectile dysfunction

Characteristics	Erectile dysfunction n=20	No erectile dysfunction n=25	P value
Fibronectin (mg/L)	49.00±3.76 ^b	32.40±2.31 ^a	0.000
Nitric oxide (µmol/L)	45.21±3.77 ^b	49.59±4.40 ^a	0.002
Glycated haemoglobin (%)	9.84±0.53	10.34±0.71	0.591
Von willebrand factor (ng/L)	42.93±8.03	34.67±7.95	0.475
Tumor necrosis factor (ng/L)	91.50±40.46	52.93±13.41	0.329

Values are expressed as mean ± SEM; Values of the group with superscript "a" are statistically significantly (p<0.05) different from group A. Values of the group with superscript "b" are statistically significantly (p<0.05) different from group B.

Table 6. Endothelial function biomarker in diabetes subjects with blurred vision

Characteristics	Blurring of vision n=25	No blurring of vision n=75	P value
Fibronectin (mg/L)	45.52±3.32 ^b	34.00±1.91 ^a	0.003
Nitric oxide (µmol/L)	53.91±3.86	57.94±4.84	0.645
Glycated haemoglobin (%)	9.76±0.44	9.37±0.39	0.602
Von willebrand factor (ng/L)	32.17±6.45	36.72±4.81	0.620
Tumor necrosis factor (ng/L)	92.89±32.96	93.72±24.94	0.986

Values are expressed as mean ± SEM; Values of the group with superscript "a" are statistically significantly (p<0.05) different from group A. Values of the group with superscript "b" are statistically significantly (p<0.05) different from group B.

There was no significantly (p>0.05) difference between the mean concentration of tumor necrosis factor alpha in diabetic subjects with blurred vision (92.89±32.96ng/L) compared to diabetic subjects with no blurred vision (93.72±24.94ng/L).

DISCUSSION

Diabetes mellitus is a common group of metabolic disorders that are characterized by chronic hyperglycaemia resulting from relative insulin deficiency, insulin resistance or both (Tuomilehto, 2019). The clinical complications associated with diabetes are most likely the consequences of hyperglycaemia through altered metabolism, non-enzymatic glycation of protein and advanced glycosylated end -products (AGEs) that accumulated in long-lived proteins such as vascular collagen and reduce the elasticity of vessel walls (Bruce and Mallika, 2019). Endothelial dysfunction in diabetes originates from hyperglycaemia and its immediate biochemical sequelae directly alter endothelial function or influence endothelial cell functioning indirectly by synthesis of cellular fibronectin, growth factors, cytokine and vasoactive agents in other cells (Tabit *et al.*, 2010).

In this present study, out of 100 diabetic subjects recruited for the research 55 (55%) were females this indicates that, most of the patients that visit diabetic clinic in Specialist Hospital, Sokoto were female. Though, the prevalence of diabetes mellitus is common amongst male (Dabelea *et al.*, 2014), however, several studies revealed that, the risk of having diabetes mellitus in female, is inversely related to their socio-economic status (Uloko *et al.*, 2018), this may be why most of the diabetes mellitus patients that visits specialist hospital, Sokoto were female and had no formal education. Another reason, may be because most of the women that visit the hospital seeking medical attention are full time house wife and therefore, can have enough time to go to hospital. This tallied with a study carried out by Uloko *et al.* (2018) which revealed that majority of diabetic patients in Nigeria are female this may not be unconnected that, female are at risk of developing obesity which is an important risk factor of type II diabetes mellitus (A.D.A., 2017). Obesity is a major factor and can potentiate type 2 diabetes mellitus in genetically susceptible individuals (Costanzo *et al.*, 2015). This present study revealed that the plasma level of cellular fibronectin, glycated haemoglobin, von willebrand factor and tumor necrosis factor

alpha which could be used as endothelial function biomarkers were significantly increased in the diabetic subjects compared to control. This may not be unconnected that, diabetic subjects with poor, long standing glucose control have tendencies of developing diabetic complications like retinopathy and nephropathy, due to endothelial injury (Azeze *et al.*, 2018). Risk factors for macro vascular disease are well established in the non - diabetic population and include hypercholesterolemia, increased plasma fibrinogen, smoking, obesity and hypertension (Zelihic *et al.*, 2015). The evidence is that in the diabetic population the same risk factors operate but diabetes has an additive effect with them (Zelihic *et al.*, 2015). In addition, in type 2 diabetes mellitus, there is tendency for some of these risk factors to aggregate; these patients are particularly at risk of obesity, dyslipidaemia and hypertension (Corsino *et al.*, 2017). A common feature of these changes is their association with insensitivity to insulin, an observation which has led to the proposal that insulin resistance is the single unifying factor causing an excess risk of macro vascular disease (Furukawa *et al.*, 2017). The cause of microvascular disease is not clearly defined but the belief is that hyperglycaemia is a major factor (Packer, 2018). Microvascular complications are not simply genetically determined since the non - diabetic twins of diabetic patients do not get them. For complications to develop, hyperglycaemia must be present (Packer, 2018). The cause of hyperglycaemia is irrelevant since microvascular complications are a feature of all types of diabetes mellitus (Magri *et al.*, 2018). As diabetes is defined by hyperglycaemia, it is reasonable to anticipate that; these microvascular complications should result from this hyperglycaemia. It is clear that the risk of diabetic complications is related to the duration of the disease (Targher *et al.*, 2018). In a study carried out in Netherland that assess the plasma levels of cellular fibronectin in diabetes patients, shows that circulating cellular fibronectin were significantly higher in diabetic patients compared to ischaemic stroke patients (Kanters *et al.*, 2001). Similar finding was also obtained in China, though, von willebrand factor was only used for the research, by Chen *et al.* (2013) it revealed that type II diabetic patients older than 60years of age had increased levels of VWF, VWF activation and VWF propeptide compared to younger patients who are non-diabetics. They also found that the total active VWF was associated with the time of being diagnosed with diabetes, indicating that probably the total active VWF is the best

marker in this patient population for endothelial activation and endothelial damage (Chen *et al.*, 2013). However, the mean nitric oxide level was significantly lower in control compared to diabetic subjects. This may be due to impaired conversion of arginine to NO due to oxidative stress in type II diabetes mellitus (Tessari *et al.*, 2010). Though many researches show that nitric oxide was significantly increased in diabetic subjects with complication (Chen *et al.*, 2018). Nitric oxide (NO) is a crucial player in vascular homeostasis, is synthesized within endothelial cells, during conversion of L-arginine to L-citrulline by endothelial nitric oxide synthase (eNOS) (Sambe *et al.*, 2018). It is released from endothelial cells mainly in response to shear stress elicited by the circulating blood or receptor-operated substances such as acetylcholine, bradykinin, or serotonin. NO diffuses to vascular smooth muscle cells (VSMC) and activates soluble guanylate cyclase (sGC), yielding increased levels of cyclic guanosine-3,5-monophosphate (cGMP) and relaxation of VSMC. Additionally, NO also prevents leukocyte adhesion and migration, smooth muscle cell proliferation, platelet adhesion and aggregation, and opposes apoptosis and inflammation having an overall antiatherogenic effect (Todd Milne *et al.*, 2017). This also tally with research carried out by Chen *et al.* (2018), which shows that there is a decrease in bioavailability of NO as a result of oxidative stress caused by hyperglycaemia. The results of plasma concentration of endothelial function biomarkers in diabetic subjects based on the treatment duration of diabetes mellitus in years, this present study shows that, the mean concentration of plasma cellular fibronectin in diabetes subjects with a treatment duration of ≥ 5 years was significantly higher compared to diabetes subjects with < 5 years duration of treatment. A long period of diabetes and poor glucose control is associated with an increased production of glycosylation end products, metabolic derangements, endothelial cell activation, endothelial injury and reactive oxygen species production with increased level of cellular fibronectin (Kanters *et al.*, 2001). Poor compliance of diabetic drugs might be another factor that could result to endothelial injury and consequently increase in plasma level of cellular fibronectin, this is in agreement with research carried out by (Alghadir *et al.*, 2016).

Also, in this study we observed significantly lower plasma NO levels in subjects living with diabetes for more than 5 years compared to subjects having diabetes for less than 5 years and also in control. This corresponds with a study carried out by Assmann *et al.* (2016). However, in another study carried out in India in type II diabetes mellitus subjects revealed higher NO levels with hyperglycaemia (Adela *et al.*, 2015). This study also shows a significant decrease in NO level, in diabetes mellitus subjects with erectile dysfunction compared to diabetic subjects without the complication. NO might have both beneficial and harmful effects depending upon its concentration (Adela *et al.*, 2015). On the other hand, NO causes relaxation of blood vessels reducing blood pressure, prevents platelet aggregation and adhesion, limits oxidation of LDL cholesterol, inhibits proliferation of smooth muscle cells, and decreases the expression of pro-inflammatory genes that are associated with atherogenesis (Yang *et al.*, 2010). However, NO interacts with O_2 leading to NO inactivation and production of peroxynitrite, which post-transcriptionally modifies proteins and negatively affects their function (Cosentino *et al.*, 1997). This may contribute to endothelial dysfunction by inactivating eNOS dimers, a zinc ion is held by four thiols, two from each monomer.

Physiological relevant concentrations of peroxynitrite oxidize the zinc thiolate center in eNOS, releasing zinc and oxidizing the thiols (cite). Upon thiol reduction, eNOS dimers dissociate into monomers. This modification reduces NO bioactivity and enhances O_2 production, which reacts with NO, further generating more peroxynitrite, in a process called as eNOS uncoupling (Aydın *et al.*, 2001). Increase in NO concentrations or its reduced bioactivity due to eNOS uncoupling may not only be relevant to the development of endothelial dysfunction and atherosclerotic complications in diabetes mellitus, but may also affect insulin-mediated postprandial glucose disposal and possibly contribute to the development of insulin resistance (Domingueti *et al.*, 2016). However, the plasma level of glycated haemoglobin in this study was found to be on the increasing side as duration of diabetes increases. This correlates with a study carried out by Verma *et al.* (2006) in India, which revealed that the amount of carbohydrate attached to HbA1c increases with increasing duration of the disease. Also, in another study, it revealed that diabetic subjects with poorly controlled glucose showed a significant correlation between HbA1c and duration of diabetes (van Steen *et al.*, 2018). The mean concentration of the von Willebrand factor was significantly higher in subjects living with diabetes mellitus for more than 5 years compared to subjects with diabetes for less than 5 years and also in controls. This is in agreement with a study carried out in China by Chen *et al.* (2013), which said there is a positive correlation between the period of suffering from diabetes and von Willebrand factor related parameters. Indicating that the duration of the diabetic process influences the endothelium and thereby promotes VWF secretion (Chen *et al.*, 2013). The mean plasma concentration of tumor necrosis factor alpha was significantly elevated in diabetic subjects with the disease for more than 5 years compared to subjects living with the disease for less than 5 years, this is similar to a study done in Japan (Yano *et al.*, 2004) which revealed that circulating levels of TNF- α increase with duration of disease.

This is because hyperglycaemia stimulates TNF- α secretion from monocytes and endothelial cells (Yano *et al.*, 2004). However, TNF- α can induce vascular injury by affecting the balance between coagulation and fibrinolysis (Margetic, 2012). TNF- α stimulates the expression of tissue factor that is the initiator of blood coagulation activation and the secretion of plasminogen activator inhibitor-1 that inhibits fibrinolysis (Margetic, 2012). This present study shows that, the mean concentration of cellular fibronectin, glycated haemoglobin, von Willebrand factor and tumor necrosis factor alpha in diabetic subjects with erectile dysfunction were significantly higher compared to diabetic subjects with no erectile dysfunction. Endothelial dysfunction is the hallmark of erectile dysfunction in diabetes mellitus subjects and might be through PKC activation, activation of the hexosamine and polyol pathways and formation of advanced glycation end product (Brownlee, 2001). This is in agreement with research carried out by (Malavige and Levy, 2009). However, the mean concentration of nitric oxide in diabetic subjects with erectile dysfunction was significantly lower compared to diabetic subjects with no erectile dysfunction. This study revealed that the mean concentration of cellular fibronectin, glycated haemoglobin, von Willebrand factor and tumor necrosis factor alpha in diabetic subjects with blurred vision were significantly higher compared to the diabetic subjects with no blurred vision. This is because microangiopathy is one of the complications that could be sequel to chronic diabetes mellitus

and eye is not exceptional organ. This was proved right in a research carried out in by (Khan *et al.*, 2017). Hyperglycaemia is the hallmark of diabetic retinopathy starting from advanced glycosylation products and activation of cytokines, could result to retinal pigment epithelium, microaneurysm, inter-retinal oedema, haemorrhage, exudate and intraocular neovascularization (Wan *et al.*, 2015).

Conclusions and Recommendations

Conclusions

Endothelial function biomarkers cellular fibronectin, von Willebrand factor and tumour necrosis factor alpha were significantly higher in diabetic subjects compared to control. Endothelial function biomarkers nitric oxide was significantly lower in diabetic subjects compared to control. Diabetic subject with duration of the disease for more than five years had increase level of endothelial function biomarkers compared to diabetic subject living with diseases for less than five years. Diabetic subjects with complications had higher level of cellular fibronectin compared to diabetic subjects without complication.

Recommendations

Endothelial function biomarker especially cellular fibronectin, nitric oxide, von willebrand factor and tumour necrosis factor alpha should include in the routine investigation in the management of diabetic patients.

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