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OXIDATIVE STRESS AND DIABETIC COMPLICATION: A systematic review

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Abstract: Various indicators, such as protein modification biomarkers and lipid peroxidation products, have been used in research to show a relationship between diabetes and oxidative stress. Due to their capacity to damage lipids, proteins, and DNA, free radicals are thought to play a key role in the initiation and progression of late diabetes complications. Oxidative stress causes a number of clinical illnesses, including rheumatoid arthritis, diabetes, and cancer. Coronary artery disease, neuropathy, nephropathy, retinopathy, and stroke are all consequences of DM caused by free radicals and oxidative stress. In-vivo studies back up the theory that hyperglycemia causes oxidative stress, which leads to endothelial dysfunction in diabetes patients' blood vessels. Increased glucose and insulin levels, as well as dyslipidemia, produce macroangiopathies, which lead to oxidative stress and atherosclerosis in diabetic patients.

Objective- Exogenously and endogenously produced reactive oxygen and nitrogen species promote oxidation of cellular molecules by transferring electrons to them. Exogenous antioxidants can be obtained, and the body's antioxidant defence system comprises vitamins A, C, and E, glutathione (GSH), and the enzymes superoxide dismutase (SOD), catalase, and others. Hyperglycemia causes the creation of free radicals, which causes an imbalance between free radical formation and the antioxidant defence system, resulting in oxidative stress. Increasing data suggests that oxidative stress plays a key role in diabetes mellitus development (DM). Increased formation of advanced glycation end-products (AGEs), polyol pathway, increased expression of the receptor for AGEs and its activating ligands, activation of protein kinase C (PKC) isoforms, and overactivity of the hexamine pathway are the main pathways involved in diabetic pathogenic complications

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1. Introduction

More than 180 million individuals in industrialised and developing countries have diabetes, and the number is expected to quadruple by 2030. (1). Diabetes mellitus is characterised by an increase in blood glucose levels and a decrease in insulin production or sensitivity to its goals (2). Diabetic problems damage the eyes, kidneys, nerves, and blood vessels as a result of long-term hyperglycemia and oxidative stress (3). Sulphonylureas and arginine, in addition to glucose, stimulate beta cells to release insulin, and glucose is a major determinant of insulin secretion (4). Type 1 diabetes is an autoimmune illness caused by the death of pancreatic cells. It accounts for around 80% to 90% of diabetes in children and adolescents (5, 6). T-cells and B-cell responses both have a role in autoimmune type 1 diabetes (7). The major symptoms of type 1 diabetes include polydipsia, polyuria, polyphagia, abrupt weight loss, and blurred eyesight (7). Type 2 diabetes mellitus accounts for approximately 90% of all diabetes cases. The majority of these patients are adults, and obesity is the leading cause. Adult diabetes is a term used to describe diabetes that develops beyond the age of 40. The release of plasma free fatty acids (FFA) and tumour necrosis factor alpha (TNF α) by "full" adipocytes is one of the key causes of type 2 diabetes and obesity. [8,9,10]

OxidativestressandDiabetes

Oxidative stress is a type of physiological stress that occurs when the antioxidant defence system and free radical production in the body are out of balance (11, 12). Vitamins A, C, and Glutathione, as well as superoxide dismutase, catalase, and other antioxidant defencesystems(13), and free radical production comprises ROS and RNS in the body. When reactive oxygen and nitrogen species are present in low or normal amounts, they are helpful. However, at high concentrations, they cause oxidative stress, which is the root of many diseases such as Diabetes, Kidney disease, Lung cancer, etc (14). The creation of ATP by cells when they use oxygen, as well as the generation of free radicals by mitochondria, are all probable sources of Reactive Oxygen Species (ROS). Reactive Nitrogen Species (RNS) and other biological redox products are common. Superoxide, hydrogen peroxide, hydroxyl radical, singlet oxygen, and RNS nitrous oxide, peroxynitrate, nitric oxide, and peroxynitrite (15,16) are examples of ROS.ROS and RNS have favourable effects on cellular response and immunological function at low or moderate levels. They cause oxidative stress at high doses, which is a harmful process that can harm all cell structures. (17, 18, and 19). Beta cells have lesser levels of antioxidant enzymes, making them more vulnerable to oxidative and cytotoxic stress (20). Hyperglycemia-induced oxidative stress and inflammation increased apoptosis and disrupted insulin secretion by causing a shift in gene expression regulation (21).

Oxidative Stress-Induced Cellular Damage AProteins

In vitro ROS react with protein amino acid residues, resulting in nonfunctional proteins. Because proteins have distinct primary, secondary, and tertiary protein structures, ROS can react with any biomolecule, including proteins, altering their structure and making them more vulnerable to proteolysis (23).

In vivo studies in diabetic rats revealed a decrease in serum proteins, which could be due to the following factors: a) decreased amino acid uptake (24) b) a significantly lower concentration of a variety of essential amino acids (25) c) an increased conversion rate of glycogenic amino acids to CO2 and H2O, and

d) a reduction in protein synthesis secondary to a decreased amount and availability of mRNA (26).

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B Lipids

Since plasma membrane is made up of

polyunsaturated fatty acids (PUFA), which are an easy target for ROS due to their numerous double bonds, a shift in membrane receptors occurs. (27). Peroxynitrite (ONOO) oxidises LDL but does not attach to LDL receptors, allowing scavenger receptors in macrophages to pick up LDL and generate foam cells, resulting in atherosclerotic plaques (28,29).Sato et al. (30) published the first evidence of lipid peroxidation in DM, noting that the levels of lipid peroxides in plasma of DM patients were substantially higher than that of normal persons

Lipid peroxidation, which is the most researched field of research when it comes to ROS, is a significant biomarker of oxidative stress (31). After reacting it with thiobarbituric

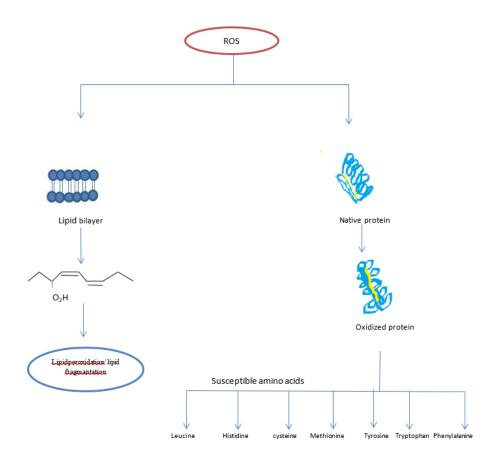


Figure 1. Shows ROS react with susceptible amino acids of proteins and polyunsaturated lipidschange the structure of lipids and proteins.

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acid, malondialdehyde (MDA) is generated as a result olf lipid peroxidation, and it can be used to assess lipid peroxides (32).

ROLES OF ORGANELLES IN OXIDATIVE STRESS

Mitochondria

As electrons transfer through the electron transport system, proton move into intermembrane space, generating a proton gradient that drives ATP synthase, but some electrons leak out of complex 1 or 3 and interact with molecular oxygen to form superoxide anion, mitochondria is rightly called the powerhouse of the cell. Hyperglycemia causes more oxygen to be released, which leads to an increase in the generation of reactive oxygen species (ROS) (33-35).

Furthermore, enzymes such as acyl-CoA dehydrogenase and glycerol phosphate dehydrogenase can produce ROS (36). By increasing mitochondrial superoxide buildup in

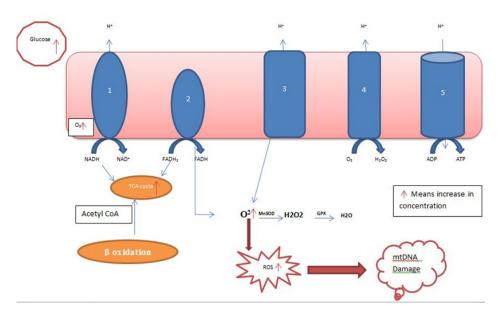
the retina, inactivation of complex 3 activities worsens diabetic retinopathy outcomes (37).

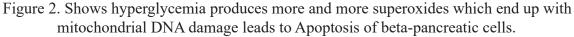
Hyperglycemia caused superoxide to develop in the mitochondrial electron transport chain by raising the inner mitochondrial membrane potential through the Krebs's cycle (37).

Hyperpolarization of the mitochondrial membrane potential and an increase in the ATP/ADP ratio occurred as a result of this scenario, followed by inhibition of complex-III and electron buildup at coenzyme Q. As a result of the partial reduction of O2, free radical production is accelerated, and ATP synthesis is reduced (38,39).

Endoplasmic reticulum stress and insulin resistance

Beta pancreatic cells have a well-developed endoplasmic reticulum (ER), which is the organelle responsible for protein folding and exporting. Because they must produce and secrete significant amounts of insulin, they





have a well-developed ER (40-42)

Protein disulfide isomerases (PDI), endoplasmic reticulum oxidoreductase 1 (ERO1), and glutathione (GSH) combine as a chaperone-like effect in ROS production, which in turn inhibits disulfide bond formation, causing endoplasmic reticulum ER stress by increasing the amount of misfolded proteins in the ER lumen.

Hyperglycemia or an increase in misfolded proteins triggers the unfolded protein response (UPR). In normal physiological settings, the three transmembrane proteins Inositol-required kinase 1 (IRE1), protein kinase-like ER kinase (PERK), and activating transcription factor 6 (ATF6) are all inactive. When the UPR coping mechanisms fail to restore ER stability or homeostasis, the ER stress response ensues. The separation of Bip from ER transmembrane proteins causes ER stress to activate them. (43,44). NO generates ER stress and the production of unfolded proteins by depleting ER Ca2+. The cell responds to unfolded proteins in three ways: upregulation of chaperon proteins, upregulation of chaperon proteins, and upregulation of chaperon proteins.(45,46)

Mechanisms of Hyperglycemia-Induced Damage

Five mechanisms main major are underlying hyperglycemia-induceddiabetic vascular damage which leads to diabetic complication, microvascular and macrovascular damage. ROS responsible for the activation of these five mechanisms which is produced due to mitochondria over activation- polyol pathway, increased intracellular formation of advanced glycation end-products (AGEs), increased expression of the receptor for advanced glycation end products and its activating ligands(RAGE), activation of protein kinase C (PKC) isoforms, and overactivity of the hexamine pathway. . However, the results of clinical studies in which only one of these pathways is blocked have been disappointing (47, 48)

Increase polyol pathway

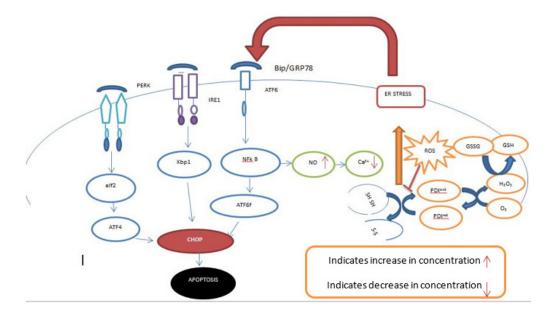


Figure 3. ROS inhibits proper disulfide formation in protein folding cause dissociation of Bip from ER transmembrane proteins like PERK, IRE1, ATF6 cause ER stress. Activation of these proteins causes apoptosis of cells by expression of CHOP gene.

Nerve, retina, lens, glomerulus, and vascular cells rich in Aldose reductase(49).When the body experiences hyperglycemia, the enzymes aldose reductase and sorbitol dehydrogenase are active, converting glucose to sorbitol and sorbitol to fructose, respectively (50). As a result of these reactions, NADPH is oxidised and NAD+ is reduced, which is employed as a cofactor by aldose reductase and sorbitol dehydrogenase, respectively, causing the following:-

1, Because NADPH is utilised as a cofactor to replenish reduced glutathione (GSH), oxidation of NADPH causes a drop in NADPH content in the cell, which lowers the activity of the reducing enzyme Glutathione, resulting in more superoxides being produced (49-50).

Sorbitol does not permeate through the cell membrane and accumulates intracellularly because it is hydrophilic. Furthermore, sorbitol is tough to digest (51,52,53).

2, Sorbitol produced by aldose reductase is oxidised to fructose by sorbitol dehydrogenase, which uses NAD+ as a cofactor to make NADH, which serves as a substrate for NADH oxidase, which produces reactive oxygen species (ROS) (54,55). The synthesis of diacylglycerol, a key activating cofactor for protein kinase-C, is increased by hyperglycemia inside the cell(56,57) PKC changes gene expression and protein function when it is activated by intracellular hyperglycemia (57).

Increased Intracellular AGE Formation

Advanced glycation end-products form nonenzymatically as a result of the reaction of reducing sugars with free amino groups, which occurs not only in proteins but also in lipids and nucleic acids and is known as Maillard's reaction (58). The first reaction produces an unstable product (Schiff base), which undergoes rearrangement to produce a more stable product (amadori product), which then undergoes oxidation, dehydration, and cyclization processes to produce a glycated molecule (59). AGEs cross-link proteins, resulting in proteinase-resistant aggregates. (60).

IgG accounts for 75% of the total immunoglobulin in serum (61) It has the longest half-life of all the immunoglobulins, which makes it a prime target for ROS because the longer the half-life, the more vulnerable to ROS (62)Glycation of IgG causes a change in its structure and function, which is linked to

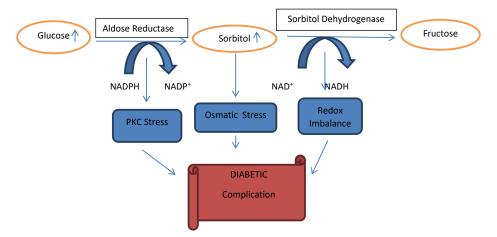


Figure 4: Polyl pathway involved in diabetic complication by osmatic stress,PKC stress and Redox imbalance

inflammation and a target for autoantibodies in individuals with rheumatoid arthritis (63)

Glycation of collagen and elastin can cause skeletal fragility and osteoblast differentiation by modifying characteristics such as the triple helix, solubility, and flexibility, which reduce toughness and stiffness. (64-69)Glycated collagen affects endothelial cell function, which contributes to the formation of atherosclerotic plaques. (69)

Several in vitro investigations have established the critical role of glycated albumin in platelet activation and aggregation (70) and glycated albumin can impact glucose metabolism in both skeletal muscle and adipocyte. (71) Glycated fibrinogen hinders fibrinolysis and increases fibrin gel permeability, resulting in a thrombogenicfibre network. (72,73) The interaction of AGEs with RAGE on macrophages activates p21ras and the mitogenactivated protein (MAP) kinase signalling pathway, which leads to nuclear factor-B (NF-B) activation (74). Which, in turn, modify gene transcription for a variety of reasons, including the production of pro-inflammatory cytokines like interleukin-1 (IL-1), tumour necrosis factor- α (TNF- α), and adhesion molecules like collagen.In order to modulate immune surveillance and inflammation, VCAM-1 works in conjunction with other adhesion molecules. Excessive levels of ROS, oxidised low density lipoprotein (oxLDL), 25-hydroxycholesterol, turbulent shear stress, high hyperglycemia, and microbial stimulation of endothelial cell TLRs all increase VCAM-1 expression. The transcription factors nuclear factor kappa B (NFB), SP-1, Ap-1, and interferon regulatory factor-1 influence the activation of VCAM-1 gene expression. (75-79)

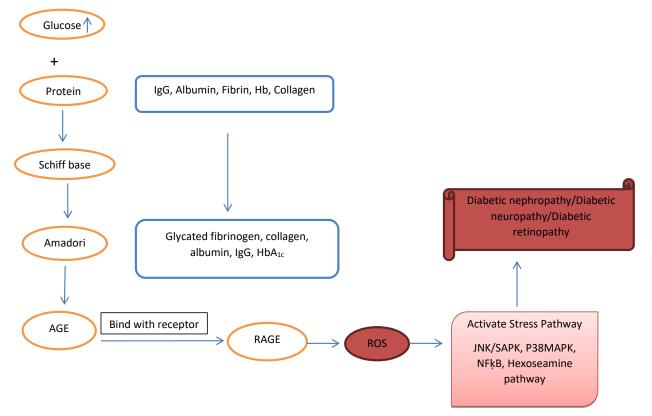


Figure 5: AGE role in diabetic complication by interacting with RAGE , produce ROS activate stress pathway result in diabetic complication.

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Following the interaction of AGEs with RAGE in endothelial cells, NF-B and hemeoxygenase mRNA, both indicators of oxidative stress, are activated .(80) Following the interaction of AGEs with RAGE in endothelial cells, NF-B and hemeoxygenase mRNA, both indicators of oxidative stress, are activated, according to a study (81)

The aetiology of diabetic problems is complicated by the interaction of AGEs with their cellular receptors .(75)

PKC pathway :

Hyperglycemia causes a rise in glycolysis, intermediate dihydroxyacetone phosphate, and glycerol-3-phosphate, all of which boost de novo DAG synthesis. (82) There are two forms of PKC conventional PKC (cPKC) isoforms (PKC- α , - β 1, - β 2, and - γ) are activated by phosphatidylserine (PS), calcium, and DAG or phorbol esters, whereas novel PKCs (nPKC; PKC- δ , - ϵ , - θ and - η) by other than calcium. The atypical PKCs (aPKC; PKC- ζ and - ι/λ) are not activated by calcium, DAG, or PMA(83). Diabetes sequelae indicated a rise in total DAG content in both vascular and nonvascular tissues from diabetic animal models and patients, including renal glomeruli,(84-86), heart, aorta,(87), and retin(88), liver (89)), skeletal muscles,(90). Tight junction between endothelial cells provide a vascular barrier, although PKC enhances the permeability of EC albumin and other macromolecules in diabetic circumstances. (91,92) PKC influences the expression and activity of vascular endothelial growth factor (VEGF) and increases the synthesis of thromboxane endothelin-1 (ET-1) through modifying the bioavailability of NO (93,94,95)Endothelial dysfunction occurs as a result. Different forms of PKC activated by Hyperglycemia among which PKC- α , - β - δ , and $-\varepsilon$ involved in retina dysfunction. PKC- α , - β - δ , and $-\varepsilon$ altering enzymes NO, ET-1, VEGF, PDGF, etc involved EC and pericytes. (96-99) Many in vitro investigations have shown that different versions of PKC perform diverse roles in the proximal tubules of the kidney and the glomerulus. By changing NADPH and boosting renal serum and urinary VEGF, PKC- α causes superoxide to develop. (100,101)The deletion

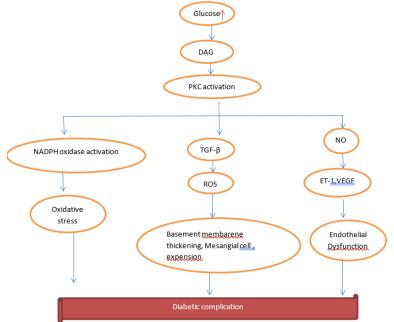


Figure 6: Denovo synthesis of DHA in hyperglycemia activate PKC pathway, result in activation of different factors lead to produce ROS, complicate diabetes.

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of the TGF gene protects diabetic kidney disease in animals. TGF- causes the development of reactive oxygen species (ROS), which impairs glomerular filtration.(102)

Hexosamines Pathway

Excess glucose is provided by hyperglycemia, and the rate-limiting enzyme glutamine: fructose-6-phosphate (F-6-P) amidotransferase (GFAT) allows additional fructose 6 phosphate to enter HSP, forming the primary end product UDP-N-acetylglucosamine (UDP-GlcNAc). Which modulates glucose entrance into the HSP pathway by acting as an allosteric feedback inhibitor of GFAT (103) UDP-Nacetylglucosaminetransferase (O-Glc NAC transferase) is an enzyme that uses UDP-Nacetylglucosamine (UDP-GlcNAc). Many proteins/transcription factors are acylated by transferring N-acetyl glucosamine to Olinkage of serine or threonine residues in certain proteins55 IRS-1 and 2, Glut 4, glycogen synthase (104,105)and RNA polymerase II (106)

Sp1 protein with an O-GlcNAc modification that binds to Sp1 binding sites of plasminogen activator inhibitor 1, (PAI-1) triggered by hyperglycemia. In vascular smooth muscle cells (107), aortic endothelial cells (108) and mesangial cells (109), Sp1 regulates hyperglycemia-induced activation of the PAI-1 promoter, which is regarded to be an important element in the development of vascular disease in diabetes.

The promoter sequence of TGF- β homologs glucose response elements of genes of glucose metabolic proteins involved in glycolysis, such as pyruvate kinase. As a result of their association with the stimulatory factors USF-1 and 2, GREs produce TGF- β 1 overexpression in hyperglycemic conditions. TGF- β synthesis has been shown to be a distinct impact of high glucose levels (110,111).

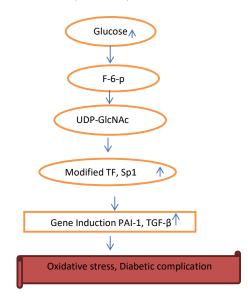
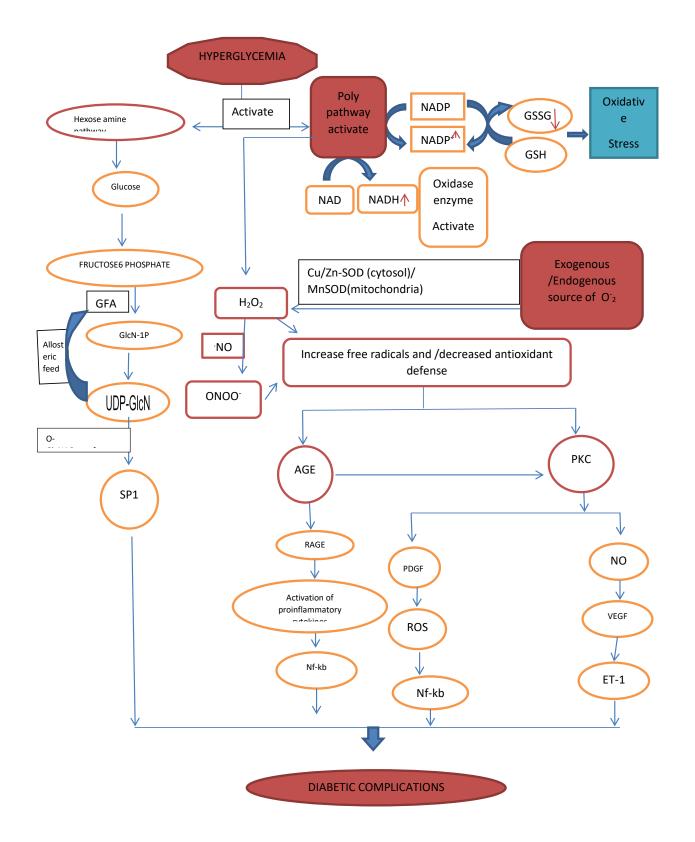


Figure 7:Hyperglycemia supplies extra glucose, and the rate-limiting enzymes enables more fructose 6 phosphate into HSP, resulting in the major end product UDP-N-acetylglucosamine (UDP-GlcNAc).

CONCLUSION-

Several research investigations have indicated that oxidative stress is a crucial factor in the occurrence and consequences of diabetes, as mentioned in this review.

As discussed above, there is substantial evidence from in vitro and in vivo studies that hyperglycemia and perhaps raised FFA levels cause the formation of ROS and RNS, resulting in enhanced oxidative stress in a number of organs. The mechanism becomes overwhelmed in the absence of a suitable compensating response from the cell's natural antioxidant network, resulting in redox imbalance and aggravating the problem. The reactive species not only cause direct cell damage by oxidising DNA, protein, and lipids, but they also cause indirect cell damage by activating stresssensitive intracellular signalling pathways such NF-B, p38 MAPK, JNK/SAPK, hexosamine, UDP-N-acetylglucosaminetransferase use (UDP-GlcNAc) to acylate .proteins55 IRS-1 and 2, Glut 4, glycogen synthase,Sp1 proteindevelopment of vascular disease in diabetes.



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PKC, AGE/RAGE, sorbitol, and others. Activation of these pathways leads to increased production of a variety of gene products that induce cellular damage and play a key role in the development of diabetes' late consequences. Furthermore, recent evidence from in vitro and in vivo studies suggests that activation of the same or similar stress pathways leads to insulin resistance and decreased insulin production. As a result, we propose that hyperglycemia and FFAinduced increases in ROS and oxidative stress, activation of stress-sensitive pathways, and the development of not just late complications of diabetes, but also insulin resistance and -cell dysfunction, are all linked.

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Declarations

I KANEEZ FATIMA, declare that the Submitted Research Paper is my original work and no part of it has been published anywhere else in the past. I take full responsibility, that if in future, the paper is found invalid according to basic rules, the last decision will be of the Authorities concerned. Any form of plagiarism will lead to the disqualification of the paper.

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Competing interests

The authors have no conflicts of interest to Availability of data and material

declare. All co-authors have seen and agree with the contents of the manuscript and there is no financial interest to report. We certify that the submission is original work and is not under review at any other publication.

1. KANEEZ FATIMA SIGN. --- ------2020-21

Katino

2. ZAKIR HUSSAIN SIGN. -----2020-21

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Consent for publication

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Author contribution

The authors confirm contribution to the paper as follows: study conception and Design, data collection, analysis and interpretation of results: Kaneez Fatima; draft manuscriptpreparation:, Zakir Hussain and Dr Rabia Hamid. All authors reviewed the results and approved the final version of the manuscript.

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The data and material is available online and can view by using the reference address.

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