

Review

Effects of Diabetes Mellitus Related Dysglycemia on the Functions of Blood-Brain Barrier and the Risk of Dementia

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Abstract: Diabetes mellitus is one of the most common metabolic diseases worldwide, and its long-term complications include neuropathy, referring both to the peripheral and to the central nervous system. Detrimental effects of dysglycemia, especially hyperglycemia, on the structure and function of blood-brain barrier (BBB) seem to be a significant backgrounds of diabetic neuropathy pertaining to the central nervous system (CNS). Effects of hyperglycemia, including excessive glucose influx to insulin independent cells, may induce oxidative stress and secondary innate immunity dependent inflammatory response which can damage cells within the CNS, thus promoting neurodegeneration and dementia. Advanced glycation end products (AGE) may exert similar, pro-inflammatory effects through activating receptors for advanced glycation end products (RAGE), as well as some pattern-recognition receptors (PRR). Moreover, long-term hyperglycemia can promote brain insulin resistance, which may in turn promote A β aggregate accumulation and tau hyperphosphorylation. This review is focused on a detailed analysis of the effects mentioned above towards the CNS, with special regard to mechanisms taking part in the pathogenesis of central long-term complications of diabetes mellitus initiated by the loss of BBB integrity.

Keywords: diabetes mellitus; blood-brain barrier disruption; dysglycemia; hyperglycemia; insulin resistance; neurodegenerative disease; diabetic complications; dementia; diabetic encephalopathy

1. Diabetes mellitus - a short resume

Diabetes, also known as diabetes mellitus (DM), is a group of the most common endocrine/metabolic diseases worldwide, characterized by sustained high blood glucose levels [1]. It results from deficiency of insulin actions, either because of insulin deficiency or because of insulin resistance. [2] It impairs mainly glucose metabolism, but it results in a disease affecting almost each tissue in the organism. Complications of diabetes mellitus may include myocardial infarction, blindness, chronic renal failure, neuropathy, and loss of cognitive functions [3]. Impaired transport of glucose to insulin-dependent cells results in glucose excess in extracellular fluids and in insulin-independent cells, which is accompanied by severe deficiency of glucose within insulin-dependent cells. Thus, some manifestations of diabetes mellitus are related to glucose depletion inside insulin-dependent cells, others are related to glucose excess toxicity in relation to other cells and tissues, yet others - to oxidative stress or using alternative substances as energy sources. Many long-term complications of diabetes mellitus depend on disorders of blood vessels, which can be divided into those referring to large vessels (macroangiopathies) and those referring to small vessels (microangiopathies). Diabetic macroangiopathies increase the risk of atherosclerosis and its related cardiovascular diseases, while diabetic microangiopathies can lead to chronic renal failure, neuropathy, and blindness, among others. While diabetic retinopathy is regarded as the most common cause of blindness in western countries, effects of diabetes mellitus towards blood-brain barrier (BBB) are much less studied. It is quite astonishing, since retina is a specialized part of the

central nervous system (CNS), and thus blood-retina barrier is regarded as a specialized region of blood-brain barrier [4]. Optic nerve is a part of CNS, and thus its capillaries carry out some barrier functions. Complications of DM in relation to BBB share many common features with those referring to diabetic retinopathy.

There are two main kinds of diabetes mellitus - type 1 DM and type 2 DM. Type 1 DM is usually an acute onset disease of young people, underlined by autoimmune destruction of beta cells of pancreatic islets (islets of Langerhans) and subsequent deficiency of insulin. Type 2 DM is usually a chronic disease with insidious onset, underlined by tissue insulin resistance and resulting deficiency of insulin actions despite quite decent concentration of insulin in the plasma. Type 2 DM is strongly associated with obesity and accounts for 90% of DM cases worldwide. The incidence of type 2 DM has been increasing for about 40 years and nowadays affects about 9% of people worldwide [5].

2. Blood-brain barrier

Blood-brain barrier has been described in detail, both morphologically and physiologically [6]. At the cellular level BBB is formed by cerebral capillary endothelial cells and the closely apposed astrocyte end-feet processes [7]. Endothelium within BBB shows a specific pattern of expression of transmembrane transport systems, regulating the transport of various substances inside and outside the cerebrospinal fluid [8,9]. BBB endothelial cells are connected to one another with tight junctions, are devoid of fenestrations, and hardly perform pinocytic transport, which promotes intracerebral environment homeostasis and is a unique trait of endothelium within cerebral microcirculation. Tight junctions connecting the adjacent endothelial cells comprise a diffusion barrier that prevents most plasma components, such as electrolytes, ionized xenobiotics, and other water-soluble substances from crossing BBB through paracellular route. On the other hand, the co-existing specialized efflux transport mechanisms, dependent on such proteins as P-glycoprotein (P-gp), breast cancer resistance protein (BRCP), and multidrug resistance protein (MRP-4), regulate trafficking of amphipathic and hydrophobic substances across BBB to protect the brain from the exposal for potentially toxic substances [10]. section may be divided by subheadings. It should provide a concise and precise description of the experimental results, their interpretation, as well as the experimental conclusions that can be drawn.

2.1. Glucose transport across BBB

Glucose transport across BBB may depend on two independent classes of glucose transporters: facilitated diffusion transporter proteins (GLUTs) and secondary active transporter proteins, also called sodium-glucose cotransporters (SGLTs) [11,12]. GLUTs include 14 proteins, the first identified of which, and the one mainly accounting for glucose transport across BBB is GLUT-1 [13,14,15]. GLUT proteins are saturable glucose carriers, transporting glucose down its concentration gradient [16,17]. While GLUT-1 molecules present in astrocytes and endothelial cells transport glucose across BBB, GLUT-3 is thought to be the most important glucose carrier transporting glucose from the cerebrospinal fluid to the neurons, although not the only one [14]. In mature and completely differentiated endothelial cells, GLUT-1 transporters are not homogenously distributed, and more of them can be found at luminal part of their cell membrane (i.e. adjacent to the lumen of brain capillaries) than at the opposite (abluminal) part [18,19].

Increased glucose transport across BBB is correlated with increased luminal expression of GLUT-1 molecules. On the other hand, increased abluminal expression of GLUT-1 molecules is observed when there is a downregulation of GLUT-1; in such situation, luminal expression of GLUT-1 molecules is reduced, while its abluminal expression rises [20], suggesting that altered distribution of GLUT-1 molecules can modulate glucose transport from the plasma to the cerebrospinal fluid.

GLUT-4, an insulin dependent glucose transporter, together with insulin receptors located within endothelial cells of brain capillaries, can also play a significant role in glucose transporting to the CNS. In normal conditions, insulin is hardly synthesized in the brain, but insulin transport across BBB has been found to increase in the course of DM [21-24]. Insulin can get to the brain through the cerebrospinal fluid, reaching it within locations devoid of BBB, e.g. in circumventricular organ, or

crossing BBB via insulin receptors acting also as insulin transporters [25,26]. A schematic diagram of BBB with participation of individual cells in glucose transport is presented in Figure 1.

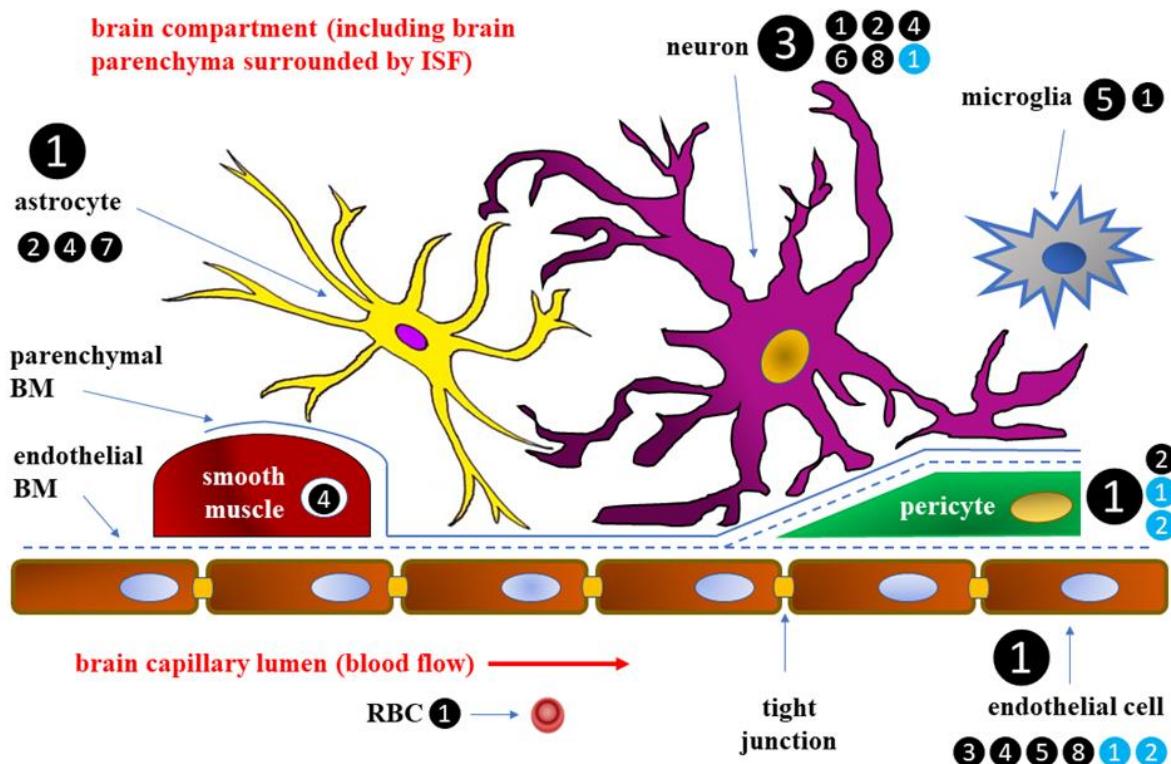


Figure 1. Schematic diagram of blood-brain barrier (BBB) [8,9,11,12,17,27,28]. The numbers in black circles (1–8) correspond to the glucose transporters (GLUTs) 1–8, respectively; the numbers in blue circles (1–2) correspond to the sodium-glucose cotransporters (SGLTs) 1–2, respectively; the main glucose transporter for a given cell type under physiological conditions is marked with an enlarged symbol. *BM* – basement membrane; *ISF* – interstitial fluid; *RBC* – red blood cell.

Effects of insulin action towards the brain are interesting in the context of discovery suggesting that DM can promote Alzheimer's disease (AD) due to mechanisms including attenuation of intracellular insulin receptor dependent signaling within BBB. Although complete elucidation of those mechanisms may require some additional studies, insulin may play a role in regulating tau protein phosphorylation in the course of tauopathies, through glycogen synthase kinase-3 (GSK-3) activation and stimulating the expression of soluble receptors for advanced glycation end products (s-RAGE) [29,30]. High expression of these receptors is correlated with an increased incidence of ischemic heart disease as a complication of DM, as well as with higher mortality due to all complications of DM, both type 1 and type 2 [31,32]. Higher levels of these receptors can also reflect the expression of tissue receptors for advanced glycation end products (RAGE) in the course of DM [33]. Advanced glycation end products (AGE)-RAGE system activation seems to play a crucial role in the pathogenesis of DM-related angiopathy and thrombosis [34,35]. Disruption of BBB integrity and function may have a profound effect on the CNS, thus being a prodromal manifestation of serious neurological disorders. DM-related disruption of BBB integrity and function can have an essential effect on the CNS.

2.1.1. Hyperglycemia and its effect on glucose transport across BBB

On the basis of general physiology, it is thought that when a substance occurs in excess, expression of receptors for the substance falls, undergoing an effect known as downregulation. These effects are aimed at homeostasis maintenance in terms of substrates delivery to the cells, and

adjusting it to the cell demand. Therefore, it is suggested that biological response to hyperglycemia may include downregulation of proteins accounting for glucose transport from plasma to peripheral tissues. Downregulation of glucose transporters in response to hyperglycemia has been confirmed in some research studies [36–38] but not in others [39–43].

Studies on the expression of glucose transporters in rats with streptozotocin-induced DM show that chronic hyperglycemia reduces both GLUT-1 and GLUT-3 expression at the level of transcription and translation; downregulation of these transporters *in vivo* is observed regardless of the method of DM induction [37]. Local brain glucose utilization rises in the course of chronic, but not acute, hyperglycemia in Sprague-Dawley rats. It is accompanied by moderate, yet clinically significant, fall of GLUT-1 expression in cerebral capillaries, but not of GLUT-3 expression [44].

2.1.2. Hypoglycemia and its effect on glucose transport across BBB

Hypoglycemia can be a major threat in the course of DM, since it can result in a serious disruption of CNS functions. Frequent attempts of reducing the plasma glucose concentration in patients with DM may lead to hypoglycemia and hypoglycemic damage of the brain cells, which can induce hypoglycemia-associated autonomic failure (HAAF). General concept of HAAF background assumes that adrenalin release from adrenal medulla, which normally would inhibit insulin secretion and boost glucagon secretion, is blunted in the course of DM. This can be accompanied by being unaware of hypoglycemia, which can lead to a vicious circle of recurrent hypoglycemia and further impairment of hypoglycemia-preventing mechanisms. Although the clinical significance of HAAF is widely known, its detailed mechanisms and mediators remain grossly unknown [45,46]. Most DM patients have difficulties with maintenance of normal plasma glucose concentration because of HAAF and hypoglycemia unawareness. Diabetic rats exposed to acute hypoglycemia does not lead to increased glucose concentration in extracellular fluid of the inferior colliculus [47] thus showing that short-lasting recurrent hypoglycemia affect neither glucose transport to the brain nor brain glucose metabolism rate. However, increased expression of GLUT-1 within BBB at the level of transcription and translation in response to chronic hypoglycemia indicates the existence of a compensatory mechanism, increasing glucose transport across BBB in case of chronic hypoglycemia [48].

Increased glucose transport across BBB in response to chronic hypoglycemia may result both from increased expression of GLUT-1 and from GLUT-1 molecules redistribution within BBB [48]. Brain glucose uptake *in vivo* in response to chronic hypoglycemia increases regardless of hypoglycemia-inducing method [49]. Similar observations (glucose concentration in the brain increased by 48%) can be made in rats exposed to hypoglycemia for 12–14 days [49]. Furthermore, hypoglycemia results in the increase of brain glucose uptake by 25–45%, as well as increased GLUT-1 expression by 23% and GLUT-1 molecules redistribution from abluminal to luminal part of endothelial cells within BBB [39].

Increased GLUT-1 expression, combined with redistribution of its molecules to luminal pole of endothelial cells can additionally aggravate hypoglycemia through stimulating glucose transport across BBB. In addition, acute or mild hypoglycemia stimulates the expression of GLUT-1, GLUT-4, angiotensinogen and mitogen activated protein kinase (MAPK) phosphatase-1 [50]. Increased expression of angiotensinogen may promote vasodilation with locally increased blood flow. This may in turn locally elevate glucose concentration, resulting in overestimation of blood glucose level by the hypothalamus and subsequently inhibit hypothalamic mechanisms aimed at counteracting hypoglycemia.

2.2. Alterations in amino acid transport across BBB in the course of DM

Choline is the precursor of acetylcholine – a neurotransmitter taking part in muscle control and memory, among others. Choline is transported to the brain by a saturable transporter acting within BBB. Prolonged hyperglycemia can inhibit choline transport across BBB [51]. As to other amino acids, transport of branched chain neutral amino acids is increased in DM, while the opposite is true for basic amino acids and some essential amino acids, such as methionine, lysine, phenylalanine, and

tryptophane. Interestingly, these alterations in amino acid transport across BBB result from their altered concentration in the plasma, not from the altered function of amino acid transporters across BBB [52].

2.3. Effects of DM on BBB integrity and permeability

Both *in vitro* and *in vivo* studies show that DM disrupts BBB integrity, which results in its increased permeability [53–56]. *In vitro* studies using co-culture of human brain microvascular endothelial cells and juxtaposed astrocytes indicate loss of BBB integrity (evaluated on the basis of transendothelial electric resistance) when hyperglycemia-mimicking conditions (glucose concentration in the medium elevated to 25 mmol/l) are maintained for at least 5 days. The integrity can be brought back to normal level after normoglycemia-mimicking conditions are restored, or when antioxidants are added [57]. In other studies [58] hyperglycemia has been found to significantly increase the expression of pro-inflammatory cytokines (TNF- α , IL-1, IL-4, IL-6) which is followed by upregulation of NF- κ B and signal transducer and activator of transcription 3 (STAT-3) proteins. There are also research studies performed on brain slices obtained from diabetic rats, showing impaired communication between astrocytes and endothelial cells through gap junctions and increased production of reactive oxygen and nitrogen species, although in an unclear mechanism [59,60]. In addition, increased production of VEGF has been observed in response to the presence of advanced glycation end products [61].

High glucose concentration (30 mmol/l) in endothelial cell cultures stimulates the expression of hypoxia-inducible factor 1 alpha (HIF-1 α) transcription factor and vascular endothelial growth factor (VEGF) as its downstream effector. VEGF enhances GLUT-1 molecule translocation to cell surface within BBB, as well as downregulates proteins that are responsible for proper functioning of tight junctions between endothelial cells (e.g. zonula occludens-1 (ZO-1) and occludin) [62]. In this way, hyperglycemia may increase BBB permeability. Occludin expression in brain microcapillaries in diabetic mice is significantly lower than in their healthy counterparts [63]. Inhibition of VEGF expression normalizes the expression of occludin and ZO-1, thus attenuating interendothelial leakage. Similar effect can be achieved through inhibition of HIF-1 α activity, which normalizes BBB integrity and permeability [64].

Research study results in humans are a bit different. Human plasma VEGF level falls in hyperglycemia and rises in hypoglycemia, which may constitute a neuroprotective mechanism, maintaining a constant rate of glucose influx to the brain [65]. Such findings implicate a key role of VEGF as a regulator of vascular permeability in the course of BBB dysfunction due to hypo- or hyperglycemia. Regardless of VEGF, recent studies suggest a significant role of matrix metalloproteinases (MMPs) in the course of dysglycemia-induced loss of BBB integrity. In diabetic rats, BBB permeability for 14C radiolabeled sucrose rises in response to increased MMP-2 expression, which is accompanied by reduced expression of occludin and ZO-1 [53, 66–68]. Advanced glycation end products, similarly to VEGF, stimulate MMP-2 release [61].

Acute transient hyperglycemia may also induce inflammatory response and endothelial damage, which has been studied on rat model of ischemia-reperfusion damage. Increased expression of HMGB1 and intercellular adhesion molecule 1 (ICAM-1) has been found as a result of ischemia-reperfusion damage, both with co-existing mild hyperglycemia (plasma glucose concentration = 150 mg/dl) and with transient severe hyperglycemia (plasma glucose concentration = 400 mg/dl). These effects have been correlated with disrupted integrity of BBB [69]. In addition, increased expression of ICAM-1 is accompanied by increased expression of IL-1 β in diabetic rats after 3 days of reperfusion [68].

Hyperglycemia-induced expression of HIF-1 α and VEGF may act synergistically, enhancing detrimental response of BBB to occlusion and stopped blood flow [70], thus additionally contributing to loss of BBB integrity observed during reperfusion, although this hypothesis requires confirmation. Both brain capillary permeability and the level of pro-inflammatory cytokines rise in DM type 1 patients with ketoacidosis. Diabetic ketoacidosis, especially pronounced in the course of type 1 DM, increases BBB permeability and promotes such complications as brain edema [71]. What can be

observed in a similar study is the absence of proteins responsible for proper functioning of tight junctions, such as occludin, claudin-5, ZO-1, and junctional adhesion molecule-1 (JAM-1), as well as albumin extravasation and increased expression of inflammatory response markers, such as NF- κ B, C-C motif chemokine ligand 2 (CCL-2), and nitrotyrosine. This indicates that neuroinflammation combined with loss of BBB integrity plays an essential role in the pathogenesis of brain edema in the course of diabetic ketoacidosis [67]. All these findings suggest that DM significantly impairs BBB integrity and maintenance of CNS homeostasis, which may promote the risk of major neurological disorders.

Hyperglycemia can also impair vitamin C delivery, both to the retina and to the brain. Vitamin C is transported across BBB with GLUT-1 transporter, in the form of dehydroascorbic acid (DHA), subsequently transformed to ascorbic acid. Vitamin C is required for biosynthesis of collagen, catecholamines and peptide neurotransmitters. In rats with streptozotocin-induced DM, transport of DHA to the brain is reduced by 84,1 % [10]. Expression of P-glycoprotein (a glycoprotein ATP-binding cassette transporter) may also be altered in diabetic mice [72–74] and brought back to normal with insulin treatment [73].

Additional histologic alterations and vascular abnormalities observed within cerebral microcirculation in the course of DM may include thickening of capillary basal membrane, collagen deposition, accumulation of lipid peroxidation byproducts, and endothelial degeneration [75]. These alterations may promote aberrant neovascularization and brain remodeling, which can contribute to vascular damage, increased risk of hemorrhage, and neurodegeneration as complications of DM [76].

Glucose metabolism for energy obtaining begins from glycolysis, during which glucose is converted to pyruvate that can be further irreversibly transformed to acetyl-coenzyme A. Acetyl-coenzyme A may further react with oxalacetate to begin tricarboxylic acid cycle (TCA) – an eight step enzymatic process coupled with nicotinamide-adenine dinucleotide (NAD) conversion to the reduced form (NADH) and flavin adenine dinucleotide (FAD) conversion to the reduced FADH. NADH and FADH transfer free electrons to respiratory chain which is coupled with ATP synthesis. Oxidative glycolysis of one glucose molecule, with H₂O and CO₂ as final products, can yield 6 NADH molecules, 2 FADH molecules and 2 ATP molecules, which is an equivalent of 36 ATP molecules. Aerobic glycolysis also provides production of reducing equivalents, such as NADH and NADPH, which counteract oxidative stress caused by endogenous and exogenous reactive oxygen species (ROS) [77–79]. In normal conditions, bioenergetic demand of mechanisms transporting various substances across BBB is provided directly or indirectly by ATP. Neurons and astrocytes cooperate to fulfill their energy requirements. Glycolysis occurs mainly in astrocytes, while TCA – in neurons. Each type of cell contains both sets of enzymes, and metabolites produced in the course of glycolysis may enter other metabolic pathways, such as pentose phosphate pathway (PPP), hexosamine biosynthetic pathway (HBP), protein kinase C pathway (PKC) and AGE biosynthesis pathway. Glucose may also enter polyol biosynthetic pathway, being finally isomerized to fructose [77–79].

3. Correlations between hyperglycemia and oxidative stress within BBB

Hyperglycemia can result in the excessive influx of glucose to the cells, which especially refers to insulin independent cells and may result in the increased ROS production, through accelerating TCA reactions with inhibition of respiratory chain and ATP synthesis at the same time. This „hyperglycemic stress” is due to the fact that TCA reactions are coupled with transforming NAD into NADH, and in normal conditions, this NADH is used to transfer free electrons to the respiratory chain. In hyperglycemia, however, intracellular ATP level is already high, which inhibits the rate of respiratory chain reactions and ATP synthesis. Thus, there is too many NADH molecules in the mitochondria, and those NADH molecules may finally transfer the electrons into atomic oxygen, producing ROS. Increased ROS production in endothelial cells which are susceptible to this kind of stress results in microangiopathies and macroangiopathies as complications of DM [77,78,80,81]. In addition, this correlation between hyperglycemia and increased ROS production may be essential for elucidating the pathogenesis of neurodegeneration as a possible complication of DM [82,83].

On the other hand, both acute and chronic hyperglycemia may stimulate the activity of PPP in cultured astrocytes, which is a source of reduced glutathione (GSH) – a coenzyme for glutathione peroxidase, inactivating ROS. Increased activity of PPP promotes reduced glutathione regeneration, thanks to which the cells are capable of counteracting the oxidative stress [84]. In accordance with this fact, nuclear factor erythroid 2-related factor 2 (Nrf2) transcription factor has been found to translocate to the cell nucleus, together with binding immunoglobulin protein (BiP). Nrf2, taking part in cell defense against oxidative stress, is usually found in the cytoplasm, but is translocated to the cell nucleus in response to oxidative stress, thus initiating an antioxidative response dependent on stimulation of expression of some endogenous antioxidant enzymes, such as NADPH-dependent quinone oxidoreductase, heme oxygenase, and glutathione S-transferase [84].

Increased ROS production in the course of hyperglycemia inhibits GADPH activity, and thus promotes glucose entering alternative metabolic pathways (glyceraldehyde 3-phosphate processing to protein kinase C (PKC) activating metabolites and AGE-producing metabolic pathway) [75,84] (Figure 2.).

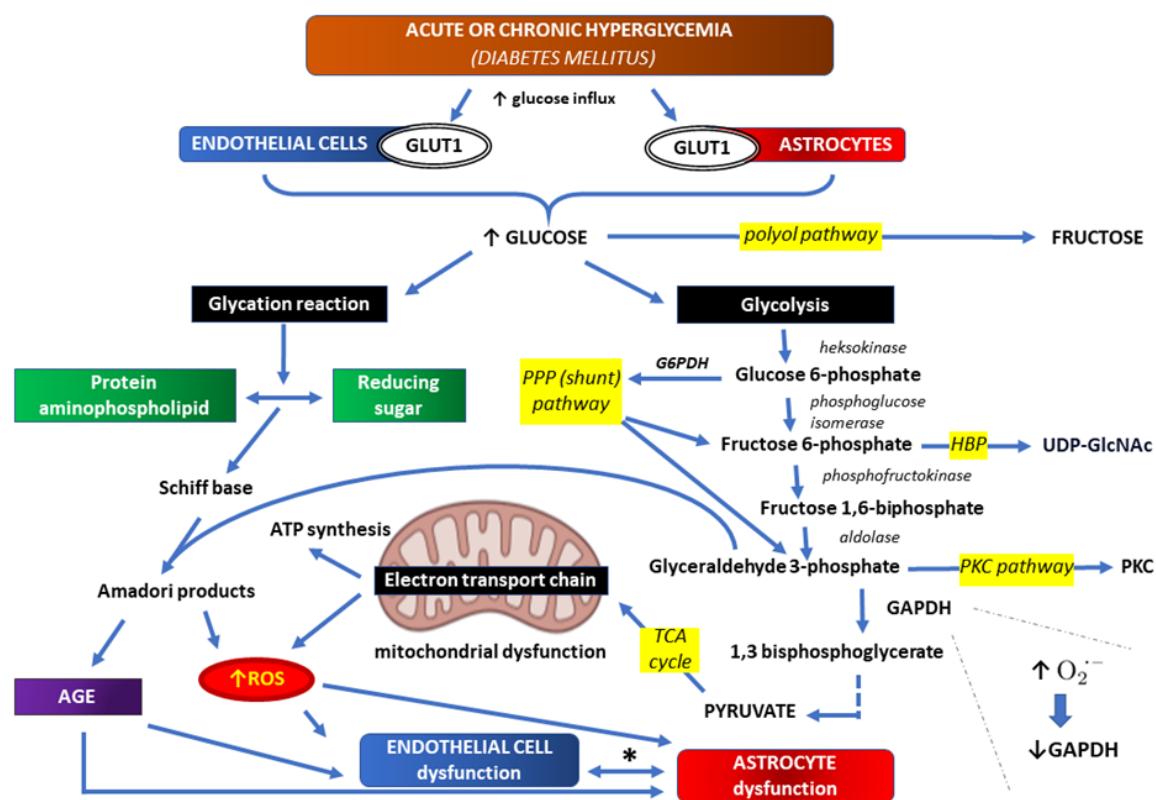


Figure 2. Hyperglycemia and oxidative stress in endothelial cells and astrocytes resulting from increased glucose influx [84,85–89].

Both endothelial cells and astrocytes use insulin-independent facilitative glucose transporter 1 (GLUT1). These cells overloaded with glucose in hyperglycemic conditions show mitochondrial dysfunction in which more than usual electrons are directly transferred to O₂ to generate reactive oxygen species (ROS) in the electron transport chain. Thus, hyperglycemia-driven mitochondrial tricarboxylic acid (TCA) cycle and its intermediates orchestrating mitochondrial oxidative phosphorylation, are a significant center for ROS production. Increased ROS production causes in turn inhibition of glyceraldehyde 3 phosphate dehydrogenase (GADPH) activity, and thus promotes glucose entering alternative metabolic pathways: glyceraldehyde 3-phosphate processing to protein kinase C (PKC) activating metabolites and advanced glycation end products (AGE)-producing metabolic pathway. Other glucose metabolic pathways are also overloaded, including polyol pathway, pentose phosphate pathway (PPP), and hexosamine biosynthetic pathway (HBP). The last

of the mentioned metabolic pathways is used for synthesis uridine diphosphate N-acetylglucosamine (UDP-GlcNAc), a nucleotide sugar and a coenzyme in metabolism.

* interdependent diabetic complications at the level of brain-blood barrier (BBB), including micro- and macroangiopathies as well as neurodegenerative disorders

Angiogenic edema occurring in the course of strokes accompanied by hyperglycemia has been found to result mainly from excessive activation of β isoform of PKC (PKC β). Further activation of PKC promotes increased BBB permeability through ZO-1 phosphorylation, impaired function of tight junctions, and raised expression of VEGF [90]. Increased intracellular AGE concentrations may damage the cells through AGE-dependent modification of various proteins, affecting their interactions with surface components of cell membranes (e.g. integrins) and receptors for advanced glycation end products (RAGE). It can refer to macrophages, endothelial cells, and smooth muscle cells. RAGE activation enhances ROS production, which in turn activates NF- κ B dependent metabolic pathways, promoting the expression of pro-inflammatory mediators [77,78,91,92] and enhancing innate immunity dependent inflammatory response. As a matter of fact, it has been found that in people suffering from AD, AGE accumulation promotes neuronal death and degeneration, which confirms a hypothesis that DM can increase the risk of AD and dysglycemia can be detrimental to BBB. In addition, oxidative stress activates matrix metalloproteinases, such as MMP-1, MMP-2 and MMP-9, which is accompanied by reduced activity of their tissue inhibitors (TIMP-1 and TIMP-2) and occurs in tyrosine kinase dependent manner [82].

Although all insulin independent cells are exposed to increased glucose concentration in the course of DM, only some kinds of cells become damaged in hyperglycemia dependent manner (e.g. retinal cells, endothelial cells), probably because they cannot reduce the expression of glucose transporters. As it has been mentioned above, hyperglycemic cell damage is dependent on five mechanisms: ① - increased influx of metabolites to polyol pathway; ② - increased intracellular production of AGE; ③ - increased expression of RAGE; ④ - activation of protein kinase C; and ⑤ - increased influx of metabolites to hexosamine biosynthetic pathway. All these mechanisms share the same downstream effector, which is increased mitochondrial ROS production [77,93].

4. Role of HMGB1 as RAGE ligand in detrimental effects of DM towards the CNS

High mobility group box 1 (HMGB1) protein is a non-histone chromosomal protein [94] regulating gene transcription through binding to DNA or chromatin, mediated by specific receptors, including RAGE and TLRs [95–98]. Because of its binding to RAGE and TLR4, HMGB1 may act as a pro-inflammatory mediator taking part in the pathogenesis of neurodegenerative diseases, such as AD [99,100]. RAGE, initially discovered as binding advanced glycation end products, can also bind other ligands, including beta-amyloid (A β) peptide, S100 proteins, and HMGB1 [101–103]. Some research studies suggest that hyperglycemia and insulin resistance underlying type 2 DM increase HMGB1 and RAGE expression both in diabetic mice and humans [104–106]. Furthermore, it has been found that in the course of DM complications, HMGB1 activates NF- κ B signaling pathways through interactions with RAGE and TLR4 [95] (Figure 3.).

It was demonstrated that hyperglycemia and insulin resistance may increase expression of both, high mobility group box 1 protein (HMGB1) and receptors for advanced glycation end products (RAGE). HMGB1 protein is a non-histone chromosomal protein regulating gene transcription through binding to DNA or chromatin, mediated by specific receptors, including RAGE and toll-like receptors (TLRs). Therefore, activation of HMGB1-RAGE-TLR4 axis in type 2 DM may induce inflammatory response via NF- κ B signaling pathway. Pro-inflammatory cytokines, activated matrix metalloproteinases, and reactive oxygen species (ROS), accumulating at the border of the brain and the capillary (vascular) compartments, are responsible for degradation of tight junction proteins such as zonula occludens-1 (ZO-1) and claudin-5. Consequently, cerebral microvessel leakage leads to increased blood-brain barrier (BBB) permeability with glial activation and the infiltration of immune cells into the brain parenchyma. Cognitive and memory impairments caused by chronic inflammation and oxidative damage are responsible for the clinical picture of diabetic encephalopathy. AGE – advanced glycation end products; A β – beta-amyloid; MyD88 – myeloid

differentiation primary response 88 (adapter protein); JNK – c-Jun N-terminal kinase; p50/p65 – NF- κ B heterodimer.

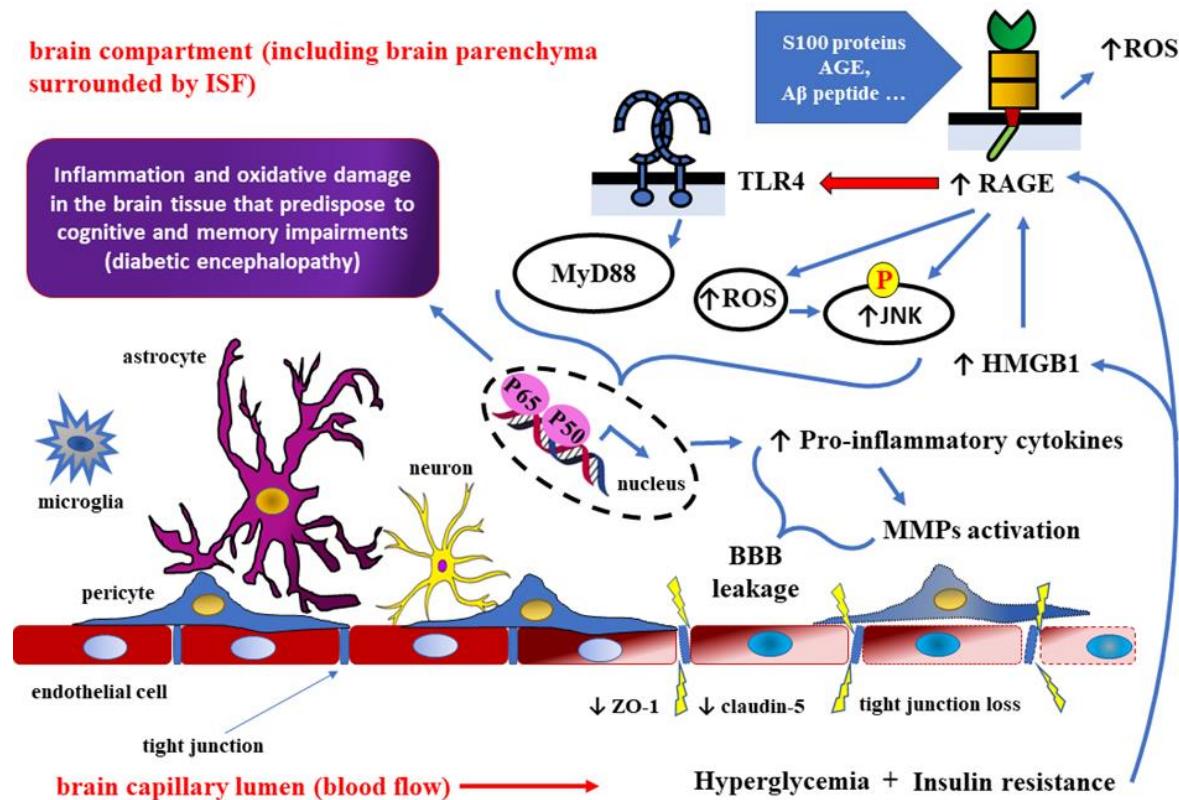


Figure 3. HMGB1-RAGE-TLR4 signaling in type 2 diabetes mellitus (DM) may compromise the integrity of BBB and increase the risk of neurodegenerative diseases [94–107].

Recent studies suggest that both in the brains of AD patients and in the CSF samples collected from them, HMGB1 concentration is elevated, just like in a mouse model of AD [108,109]. Several in vitro studies have revealed that stimulation of HMGB1-RAGE-TLR4 signaling pathway promotes hippocampal neuron damage and memory loss in the course of AD [110–112]. In addition, RAGE interactions with HMGB1 are correlated with axonal overgrowth and neuroinflammation [103,113,114]. Another study has found that HMGB1 and TLR4 interaction promotes hippocampal neuron death in patients with DM [115]. HMGB1 activates astrocytes and promotes the release of proinflammatory cytokines, as well as stimulates inducible nitric oxide synthase (iNOS) expression in cortical astrocytes, thus stimulating TLR4 signaling [116]. Interaction between HMGB1, RAGE, and TLR4 promotes A β aggregate accumulation, stimulates neuroinflammation, dampens insulin dependent signaling and impairs spatial memory [117,118]. Interactions between HMGB1, RAGE, and TLR4 are related to both DM and AD associated complications, such as A β accumulation, neuroinflammation, insulin dependent signaling, memory deficits, and microglial cells activation.

In the course of AD, there is an increased activity of metalloproteinases which can disrupt BBB integrity, thus contributing to neuronal and cognitive dysfunctions [119,120]. DM and dementia share some common features, such as severe and chronic neuroinflammation, brain insulin resistance, overaccumulation of A β and disrupted BBB integrity [121].

Elevated levels of HMGB1 are correlated both with type 2 DM, and with hyperglycemia [122–124]. In addition, HMGB1 impairs axonal growth through its interaction with RAGE [103,125], which may impair cognitive functions [126]. Besides, HMGB1 promotes A β accumulation and disrupts BBB integrity [127,128]. On a mouse model of AD, HMGB1 promotes axonal degeneration through myristoylated alanine rich protein kinase C substrate (MARCKS) protein phosphorylation, dependent on TLR4 signaling [127]. TLR4, a transmembrane protein belonging to pattern-

recognition receptors (PRR) family, often takes part in innate immunity dependent inflammation which has been correlated with AD associated pathology. NF- κ B signaling, which is downstream to TLR4 activation, promotes biosynthesis and release of pro-inflammatory cytokines [129]. TLR4 expression is markedly increased in the brains of AD patients, which promotes amyloid peptide binding and phagocytosis by microglial cells [130,131]. In the course of A β -induced neuroinflammation, HMGB1 can be localized in hippocampal neurons, where it is co-responsible for AD progression through activating RAGE- and TLR4-dependent signaling pathways [100]. In AD patients, HMGB1 accumulates both in the extracellular space and intracellular space of some brain regions [132].

Studies on in vitro models of AD show that HMGB1 is activated upon A β injection, which is accompanied by pro-inflammatory cytokine release and NLRP3 inflammasome assembly in microglial cells [133]. Furthermore, extracellular HMGB1 can impair microglia-dependent A β clearance, thus promoting AD through interactions with RAGE and TLR4 [134,135]. The same interactions may take part in the impairment of memory formation in mice [136].

HMGB1, as RAGE ligand, may promote insulin resistance of the brain through activating TLR4-JNK signaling pathway [118,137,138], as well as through stimulating TNF- α dependent signaling pathway [139,140] (Figure 4.).

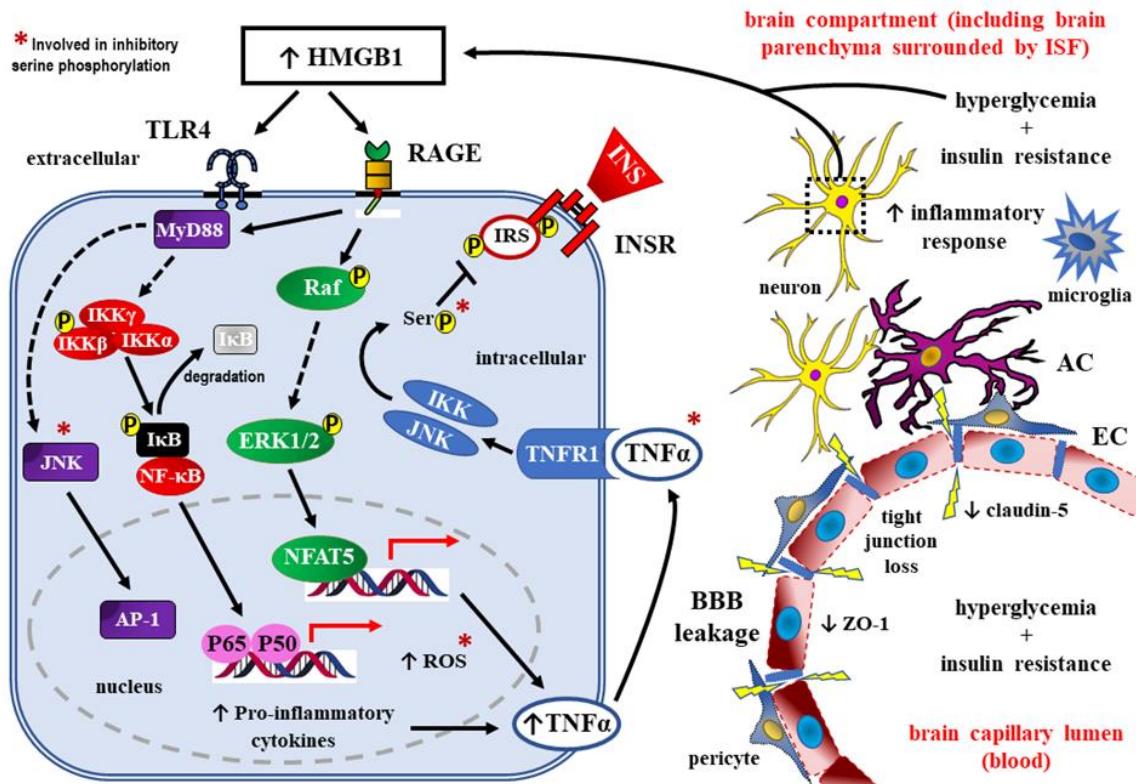


Figure 4. Increased HMGB1 expression in the brain microvessels of type 2 DM patients and ultimately in the brain compartment, may promote insulin resistance of the brain via activation of both RAGE and TLR4 signaling pathways [106,118,137–142].

Downstream signaling then activates inducible transcription factors: NFAT5, AP-1, and NF- κ B. Pro-inflammatory environment develops, because the target genes for all these transcription factors include cytokines such as IL-6, IL-1 β , IL-18, and TNF- α . TNF- α , acting on its own receptor TNFR1, may interfere with insulin signaling through phosphorylation of some serine/threonine residues in IRS, especially IRS-1. Such an inhibitory phosphorylation of IRS-1 was also demonstrated for the components of the signaling pathways (marked in the figure with “*”). Interestingly, the activation of NF- κ B pathway could in turn induce the expression of HMGB1 and its receptors, forming a

positive feedback loop to sustain inflammatory conditions. AC – astrocyte; AP-1 – activator protein 1; BBB – blood-brain barrier; EC – endothelial cell; ERK1/2 – extracellular signal-regulated kinases 1 and 2, also known as classical mitogen-activated protein (MAP) kinases; HMGB1 – high mobility group box 1; I κ B – inhibitory κ B protein; IKK α , IKK β , IKK γ – the members of the inhibitor of κ B (I κ B) kinase (IKK) family; INS – insulin; INSR – insulin receptor; ISF – interstitial fluid; p65/p50 – nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B) heterodimer; JNK – c-Jun N-terminal kinase; MyD88 – myeloid differentiation primary response 88 (adapter protein); NFAT5 – nuclear factor of activated T cells 5; P – phosphorylation; Raf – rapidly accelerated fibrosarcoma; RAGE – receptors for advanced glycation end-products; ROS – reactive oxygen species; Ser – serine; TLR4 – toll-like receptor 4; TNF- α – tumor necrosis factor alpha; ZO-1 – zonula occludens-1 (also known as tight junction protein-1)

5. Summary and conclusions

Insulin dependent signaling in the CNS seems to be crucial for maintenance of cognitive functions, through regulation of neurotransmitter release, synaptic transmission, and glucose uptake by neurons [143–145]. Disrupted brain glucose metabolism in combination with insulin resistance of the hippocampus may contribute to synaptic dysfunction, cognitive function impairment, and development of AD [146,147]. In addition, brain insulin resistance reduces cerebral blood flow [148] and cerebral cortex perfusion, which leads to cognitive deficits [149]. Moreover, insulin resistance promotes excessive A β aggregate accumulation and hyperphosphorylation of tau proteins, which results in cognitive impairment in patients with Alzheimer's disease [147,150]. Normal activity of insulin dependent signaling pathways facilitates A β aggregate clearance and inhibits senile plaque formation [151]. Insulin resistance has been found to accelerate A β formation in the vicinity of presynaptic neuronal cell membranes [152,153] and has been correlated with activation of JNK dependent signaling pathway, with subsequent inhibitory phosphorylation of insulin receptor substrate 1 (IRS-1) at S616 [154]. Thus, TLR4 activation by AGE or HMGB1 can increase the risk of dementia through promoting brain insulin resistance.

RAGE activation by AGE or by HMGB1 may exert a pro-inflammatory effect, because it activates intracellular signaling pathways stimulating NF- κ B activity [100,155], while AGE themselves can induce pro-inflammatory cytokine release, thus activating innate immunity dependent inflammatory response [156].

The sum of detrimental effects of dysglycemia towards BBB consists of its pro-oxidative effect promoting ROS production [77,78], which in turn inhibits glyceraldehyde 3 phosphate dehydrogenase (GADPH) activity [77], thus redirecting glucose metabolism to AGE [77,93]. Subsequently, the excess of AGE may stimulate RAGE as well as some TLR receptors (e.g., TLR4), which induces NF- κ B activity and promotes neuroinflammation [77,78,91,92]. Of note, pro-oxidative effects of dysglycemia can be neutralized with Nrf2 transcription factor induction [157].

Detrimental effects of dysglycemia towards BBB include: disrupting its integrity and increasing permeability, mainly through promoting oxidative stress [77,90], inducing inflammatory response through RAGE and TLR4 activation, redistribution of glucose transporters, such as GLUT-1, and related alteration of BBB permeability for choline and DHA [10,51]. These effects may in turn impair acetylcholine biosynthesis and antioxidative defense dependent on vitamin C.

As a result of the effects mentioned above, DM can increase the risk of neurodegeneration and dementia. Firstly, through disrupting BBB integrity. Secondly, through promoting brain insulin resistance with its detrimental effects on cognitive functions [121]. Thirdly, through inducing excessive production of some substances, such as AGE and HMGB1, which may promote neuroinflammation, thus abrogating the function of microglia and A β clearance [158,159]. Fourthly, through increased production of AGE and HMGB1 that can directly stimulate A β aggregate production [127,134,160], thus accelerating their accumulation in the CNS.

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Abbreviations

A β – beta-amyloid

ABC-transporter – ATP-binding cassette transporter (P-glycoprotein – P-gp)

AD – Alzheimer's disease

AGE – advanced glycation end products

ATP – adenosine triphosphate

BBB – blood-brain barrier

BiP – binding immunoglobulin protein

BRCP – breast cancer resistance protein

CCL-2 – C-C motif chemokine ligand 2 (also known as MCP-1 – monocyte chemoattractant protein-1)

CNS – central nervous system

CSF – cerebrospinal fluid

DHA – dehydroascorbic acid

DM – diabetes mellitus

FAD – flavin adenine dinucleotide

FADH – flavin adenine dinucleotide (reduced form)

GADPH – glyceraldehyde 3 phosphate dehydrogenase

GLUT-1, GLUT-3, GLUT-4 – glucose transporter 1, 3, and 4, respectively

GLUTs – glucose transporters

GSH - reduced L-Glutathione

GSK3 – glycogen synthase kinase-3

HAAF – hypoglycemia-associated autonomic failure

HBP – hexosamine biosynthetic pathway

HIF-1 α – hypoxia-inducible factor 1 alpha

HMGB1 – high mobility group box 1

ICAM-1 – intercellular adhesion molecule 1

IL-1, IL-4, IL-6 – interleukin 1, 4, and 6, respectively

iNOS – inducible nitric oxide synthase

IRS-1 – insulin receptor substrate 1

JAM-1 – junctional adhesion molecule-1

JNK – c-Jun N-terminal kinase

MAPK phosphatase-1 – mitogen activated protein kinase phosphatase 1

MARCKS – myristoylated alanine rich protein kinase C substrate

MMP-1, MMP-2, MMP-9 – matrix metalloproteinase-2

MMPs – matrix metalloproteinases

MRP-4 – multidrug resistance protein

MyD88 – myeloid differentiation primary response 88 (adapter protein)

NAD – nicotinamide-adenine dinucleotide

NADH – nicotinamide adenine dinucleotide (reduced form)

NADPH – nicotinamide adenine dinucleotide phosphate
NF-κB – nuclear factor kappa-light-chain-enhancer of activated B cells
NLRP3 inflammasome – leucine-rich repeat (LRR)-containing proteins (NLR) family member 3 inflammasome
Nrf2 – nuclear factor erythroid 2-related factor 2
p50/p65 – NF-κB heterodimer, member of the Rel family of transcription factors
P-gp – P-glycoprotein
PKC – protein kinase C
PKC β – β isoform of protein kinase C
PPP – pentose phosphate pathway
PRR – pattern-recognition receptors
RAGE – receptors for advanced glycation end-products
ROS – reactive oxygen species
SGLTs – sodium-glucose co-transporters
s-RAGE – soluble receptor for advanced glycation end-products
STAT-3 – signal transducer and activator of transcription 3
TCA – tricarboxylic acid
TIMP-1, TIMP-2 – tissue inhibitor of matrix metalloproteinase 1 and 2, respectively
TLR4 – toll-like receptor 4
TLRs – toll-like receptors
TNF- α – tumor necrosis factor alpha
UDP-GlcNAc – uridine diphosphate N-acetylglucosamine
VEGF – vascular endothelial growth factor
ZO-1 – zonula occludens-1 (also known as tight junction protein-1)

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