

Molecular and biochemical mechanisms of diabetic encephalopathy

Igor Belenichev¹, Olena Aliyeva²✉, Olena Popazova³ and Nina Bukhtiyarova⁴

¹Department of Pharmacology and Medical Formulation with Course of Normal Physiology, Zaporizhzhia State Medical and Pharmaceutical University, Zaporizhzhia, Ukraine; ²Department of Medical Biology, Parasitology and Genetics, Zaporizhzhia State Medical and Pharmaceutical University, Zaporizhzhia, Ukraine; ³Department of Histology, Cytology and Embryology, Zaporizhzhia State Medical and Pharmaceutical University, Zaporizhzhia, Ukraine; ⁴Department of Clinical Laboratory Diagnostics, Zaporizhzhia State Medical and Pharmaceutical University, Zaporizhzhia, Ukraine

Diabetes mellitus is one of the important independent risk factors for the development of neurological disorders such as ischemic stroke, transient ischemic attacks, vascular dementia and neurodegenerative processes. Hyperglycemia plays a crucial role as a trigger in the pathogenesis of these disorders. In this review, we summarize the existing data on the molecular mechanisms of diabetic encephalopathy development, consider the features of oxidative and nitrosative stresses, changes in the thiol-disulfide system, as well as mitochondrial and endothelial dysfunction in diabetes. We focus on the role of HSP 70 in cellular responses in diabetic encephalopathy. HSP70 protein is an important component of the endogenous system of neuroprotection. It acts as an intracellular chaperone, providing the folding, retention, and transport of synthesized proteins, as well as their degradation under both normoxic and stress-induced denaturation conditions. HSP70 can be considered a molecular marker and a promising therapeutic target in the treatment of diabetes mellitus.

Keywords: diabetes mellitus, diabetic encephalopathy, thiol-disulfide system, mitochondrial dysfunction, HSP70, HIF-1 α

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✉ e-mail: aliyeva1eg@gmail.com

Abbreviations: AMPA, α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid; BDNF, brain-derived neurotrophic factor; DE, diabetic encephalopathy; DM, diabetes mellitus; eNOS, endothelial nitric oxide synthase; GABA, Gamma-aminobutyric acid; GLUT4, glucose transporter type 4; GR, glutathione reductase; GSH, reduced glutathione; GSSG, glutathione disulphide; GST, glutathione S-transferase; HIF-1, hypoxia-Inducible Factor 1; HSP70, 70 kilodalton heat shock protein; IGF, insulin-like growth factor; IL-1 β , interleukin-1 β ; IL-6, interleukin 6; iNOS, inducible nitric oxide synthase; IRAK-1, interleukin-1 receptor-associated kinase; JNK, N-terminal kinase; MD, mitochondrial dysfunction; NMDA, N-methyl-D-aspartate; ROS, reactive oxygen species; TLR, toll-like receptors; TNF- α , tumour necrosis factor α

INTRODUCTION

Diabetes mellitus (DM) is recognized by numerous studies as an independent risk factor for ischemic stroke, transient ischemic attacks and vascular dementia (Tun *et al.*, 2017; Maida *et al.*, 2022). In some cases, it can be associated not only with cerebral vascular disease but also with neurodegenerative processes, in particular Alzheimer's disease. At the same time, diabetic neuropathy is the most common complication of diabetes mellitus and the leading factor among the causes of reduced quality of life in patients with diabetes (Feldman *et al.*, 2019;

Aleidan *et al.*, 2020). One of the most difficult parts of diabetes mellitus treatment is the correction of its late neurological complications. Diabetic lesions of the nervous system inevitably occur even against the background of many years of compensation for the disease, achieved through the use of modern effective and affordable antidiabetic drugs. For example, diabetes increases the risk of acute cerebrovascular events 6 times, with a nearly three-fold increase in mortality from them (Maida *et al.*, 2022; Ergul *et al.*, 2012). The development of cerebral circulatory disorders in such patients is severe, and carbohydrate metabolism disorders are associated with high mortality and disability (Lin *et al.*, 2020; Hill-Briggs *et al.*, 2021). Chronic lesions of the brain in diabetes are called diabetic encephalopathy, which leads to a decrease in cognitive and mental functions, loss of performance and quality of life in this category of patients. The discovery of new mechanisms of many cerebrovascular diseases, in particular, disturbances of the functional state of the endothelium, inflammation of the vascular wall, and programmed cell death, has opened up opportunities for the development of effective pathogenetic correction measures. In the pathogenesis of diabetic encephalopathy, the trigger link is hyperglycemia (Shi *et al.*, 2016). An important mechanism of vascular complications against the background of hyperglycemia is the activation of the polyol pathway of glucose oxidation under the influence of the enzyme aldose reductase. As a result, glucose is converted into sorbitol under the influence of aldose reductase, which leads to depletion of NADPH and, subsequently, to depletion of the glutathione link of the thiol-disulfide system, reduction of endothelial NO synthase expression, necessary for NO synthesis. A special role in the formation of vascular complications in DM belongs to the activation of protein kinase C, and subsequently to the increase in the concentration of endothelin-1 and the production of growth factors: vascular endothelial growth factor VEGF, epidermal growth factor EGF and transformed growth factor TGF- β . Also, hyperglycemia leads to increased incorporation of glucose into the hexose substitutable pathway, resulting in increased transcription of inflammatory cytokine genes and hyperproduction of reactive oxygen species and NO. Currently, oxidative and nitrosative stress are considered a universal mechanism of development of all complications in DM, including neurodegeneration and endothelial dysfunction (Tota *et al.*, 2021; Pitocco *et al.*, 2013). Recently, experimental studies have discovered a new mechanism that explains many aspects of endothelial dysfunction and neurodegradation in DM (Sivitz &

Yorek, 2010; Teodoro *et al.*, 2019; Cheng *et al.*, 2020). This is mitochondrial dysfunction, which is one of the causes of the increase in oxidative stress. Endogenous mechanisms, limiting the harmful effects of cytotoxic derivatives of NO, are provided by the thiol-disulfide system, the derivatives of which have transport properties with respect to NO, thus increasing its bioavailability. In addition, many thiols (glutathione, cysteine, methionine) can significantly limit the cytotoxicity of nitrosative stress, increasing the chance of cell survival (Kükürt *et al.*, 2021; Ren *et al.*, 2017; Aoyama, 2021). A significant role in the mechanisms of endogenous cyto- and neuroprotection is attributed to the proteins shaperones (HSP) especially, with mM 70 kDa (Belenichev *et al.*, 2023; Zhang *et al.*, 2022). However, their involvement in the molecular and biochemical mechanisms of the damage cascade mechanism in DM has not been fully identified; experimental data are scarce and sometimes contradictory. All this has prompted us to analyze and systematize world achievements in this direction, taking into account our modest results as well.

DIABETIC ENCEPHALOPATHY

The term “diabetic encephalopathy” (DE) was proposed by R. de Jong in 1950 and represents a persistent cerebral pathology resulting from the effects of acute, subacute and chronic vascular disorders, which are clinically manifested as neurosis-like and psychosis-like defects, organic, mild to moderate cognitive deficit (Dejong, 1950). It has been established that the most significant pathogenic factors initiating the development of DE are the duration of the disease, degree of DM, level of glycosylated hemoglobin, diastolic BP and total cholesterol (Li *et al.*, 2023; Wang *et al.*, 2020). In clinical observations and experimental equivalents of diabetes in animals, the duration of DM is associated with pathological changes in the CNS, characterized by cognitive and emotional deficits, which can be considered a factor in the development of dementia, as well as the risk of vascular brain complications (Feldman *et al.*, 2019; Aleidan *et al.*, 2020; Ergul *et al.*, 2012; Li *et al.*, 2023). A probable correlation between DM and cognitive function was established as early as 1922 (Miles & Root, 1922). Over the past 20 years, a number of studies evaluating the relationship between type 2 diabetes and cognitive function have been completed (Moheet *et al.*, 2015; Alkethiri *et al.*, 2021; Antal *et al.*, 2022; Kinattungal *et al.*, 2023). In DM, memory and attention are the most frequently impaired cognitive functions. Hypoglycemic conditions have a pronounced effect on the development of mnemonic disorders. In some cases, the development of dementia is possible. Under the term dementia, we understand a diffuse disorder of mental functions as a result of organic brain damage, manifested by primary disorders of thinking and memory, as well as secondary emotional and behavioral disorders. A diagnosis of dementia can be made when impairments of memory and other cognitive functions are pronounced to an extent that significantly interferes with the performance of professional and social activities in previous amounts and quality (Hugo & Ganguli, 2014). In its development, DE passes through several stages of its formation, which fully depend on the age of the disease: subclinical (coinciding with the debut of DM), clinical (corresponding to the age of the disease – from 2 to 5 years), subcompensation (over 5 years), severe (10–15 years) and decompensation (with the age

of the disease over 20 years) (Cheon & Song, 2021; Popruga *et al.*, 2021). The clinical picture of DE is characterized by a typical triad of symptoms: headaches, dizziness, and memory impairment, which in general unite DE with other types of encephalopathies. However, DE has its own specific features: progressive cognitive decline, which is sometimes referred to as cognitive aging, and sometimes is considered as a pre-stage of Alzheimer’s disease (Jayaraj *et al.*, 2020; Falvo *et al.*, 2023). Moreover, it has been established that the leading domains of cognitive impairment in type 2 DM patients are a decrease in short-term verbal memory and attention, correlating with atrophic changes in the cerebral cortex, which appear already in the early stages of diabetes and are not associated with vascular factors (Moheet *et al.*, 2015; Antal *et al.*, 2022). In addition, the clinical course of DE is characterized by frequent episodes of acute impairment of cerebral circulation, transient ischemic attacks and cerebral strokes. In particular, cerebral strokes are 6 times more frequently registered in DM patients (Maida *et al.*, 2022; Ergul *et al.*, 2012). Furthermore, chronic cerebral circulatory disorders are observed to a greater extent in patients with diabetes, leading to chronic cerebrovascular insufficiency syndrome, and the mechanisms of progressive memory decline are associated with chronic hypoxia, cerebral tissue ischemia, and neurometabolic disorders (Ergul *et al.*, 2012; Pitocco *et al.*, 2013). The concept of “brain insulin resistance”, according to which insulin receptors exist in the limbic system along with neurotransmitter receptors, is of certain interest. Their role in the mechanisms of synaptic plasticity in the hippocampus has now been established. In particular, it has been shown that insulin rapidly mobilizes functional GABA-A receptors on postsynaptic membranes of hippocampal neurons and improves synaptic transmission. In addition, a regulatory role of insulin in the functioning of AMPA and NMDA receptors of hippocampal neuronal membranes has been established (Spinelli *et al.*, 2019). Insulin can act as a mediator by accelerating the synthesis and synaptic trafficking of acetylcholine, dopamine, and other mediators. With time there is a depletion of insulin receptors in the CNS and a weakening of the function of other neurotransport systems, which to some extent explains the processes of cognitive aging (Kleinridders *et al.*, 2014). In addition, it has been experimentally established that insulin also has a neuroprotective effect under conditions of oxidative stress or DM (Soto *et al.*, 2019). Pathomorphological studies have determined a decrease in the expression of insulin and insulin-like growth factor-1 (IGF) in the hippocampus, cerebellum, pons, and basal ganglia, as well as neuronal losses in the hippocampus and frontal neocortex (Jafferli *et al.*, 2000). In the brain, insulin and IGF-1 mediate numerous effects, including glucose utilization and energy metabolism, oxidative stress, genetic regulation of other neurotrophic factors and their receptors, cholinergic gene expression and tau-protein phosphorylation, and formation regulation. They also exhibit anti-inflammatory and anti-apoptotic effects (Dandona *et al.*, 2007). Decrease of insulin regulation inhibits early response genes c-fos and c-jun with subsequent expression of IGF1 and IGF2, nerve growth factor, neurotrophin-3 and their receptors (Griffiths *et al.*, 1998; Zhang & Li, 2017). Also, insulin and IGF provide neurotrophic support in the hippocampus. It is known that diabetic neurovascular pathology is a metabolic disorder whose pathogenesis is based on the lack of insulin requirement for glucose to enter both nerve

tissue and the vascular wall. Hyperglycemia contributes to a significant (up to fourfold) increase in neuronal glucose levels with subsequent disruption of intracellular glucose metabolism and neuronal damage (Barrett *et al.*, 2017). This results in the development of secondary processes in the form of oxidative stress and protein glycosylation. At the same time, it was found that insulin activates the movement of GLUT4 to the plasma membrane in hippocampal neurons by mechanisms similar to those observed in peripheral tissues. It is considered that due to this, hippocampal neurons can significantly increase glucose utilization during neuronal activity growth (Cisternas *et al.*, 2019). But it was found that elevated plasma glucose levels of up to 6.1 mmol/L were associated with greater atrophy of structures associated with aging and neurodegeneration processes, in particular, the hippocampus and amygdalin. This indicates the relevance of glycemic control and correction in the subclinical course of DM or its absence. Disregarding the interdependence and similarity of pathogenesis, damages to the neuronal and vascular systems are fundamentally different, including the therapeutic approaches. Studies have established the role of hippocampal dysfunction in diabetes mellitus and its role in the development of DE (Spinelli *et al.*, 2019). In particular, it was electrophysiologically discovered that behavioral and mnemonic deviations in diabetic animals are associated with a deficit of long-term potentiality in the CA1 area of the hippocampus (Kumar, 2011). This factor, which reflects the synaptic plasticity of the hippocampus, was prevented by insulin therapy, while interventional treatment to normalize hyperglycemia had only a partial effect on long-term potentiality (Ho *et al.*, 2013). The role of hyperglycemia in the activation of damage mechanisms and activation of apoptosis in the rat hippocampus has also been established (Chen *et al.*, 2019). Induction of apoptosis in the hippocampus may be associated with an increase in the Bax/Bcl-2 and Bax/Bcl-xl ratio as well as caspase-3 activity (Liu *et al.*, 2013). It was experimentally determined that streptozotocin-induced diabetes led to cholinergic receptor dysfunction and reduced the neuroprotective activity of the GABAergic system in the hippocampus, indicating a high vulnerability of neurons of this brain formation and a relationship between the development of cognitive impairment and deficit (Sherin *et al.*, 2012).

Processes of systemic neuronal inflammation in the CNS also correlate with manifestations of cognitive deficits and are associated with increased levels of inflammatory cytokines (IL-1 β , TNF- α and IL-6) against a decrease in BDNF in the hippocampus (Fourrier *et al.*, 2019; Dugue *et al.*, 2017). Long-term hyperglycemia leads to the activation of cyclooxygenase-2 expression that in turn results in the increase in biosynthesis of prostaglandin E₂, which inhibits glucose-stimulated insulin secretion that disturbs cell tolerance to glucose and plays a significant role in the pathogenesis of DM. It has been experimentally established that streptozotocin-induced diabetes leads to an increase in the immunoreactivity of the inducible form of cyclooxygenase-2 (Cox-2) in the dentate gyrus and CA3-zone of the hippocampus and can thereby influence the synaptic plasticity processes in this structure (Nam *et al.*, 2011).

In general, such disorders can be the cause of rapid development of neurodegenerative changes in DM. The leading role in the pathogenesis of these complications belongs to the negative impact of oxidative stress on the function of the cells of the central nervous system,

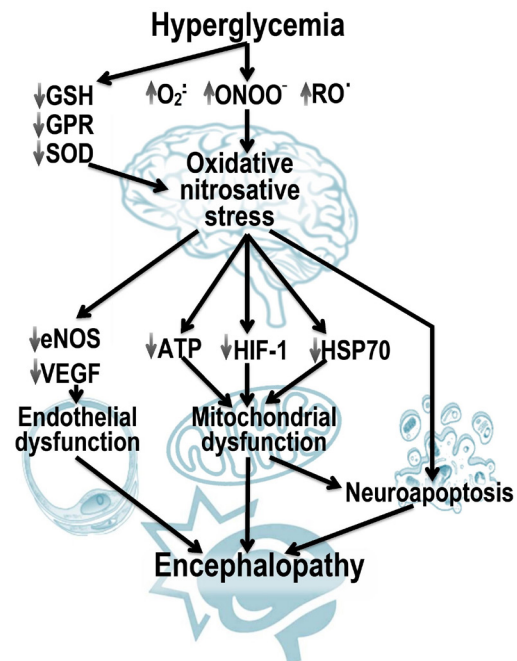


Figure 1. Molecular and biochemical cascade of links of diabetic encephalopathy pathogenesis.

and in particular the hippocampus, provided both high intracellular glucose levels and impaired microcirculation (Kükürt *et al.*, 2021; Li *et al.*, 2023). (Fig. 1).

OXIDATIVE STRESS

The main source of free-radical oxidation processes in DM is a state of chronic hyperglycemia. The development of neuronal dysfunction and the appearance of signs of cognitive deficit are based on disorders of carbohydrate metabolism, which lead to self-oxidation of glucose, to activation of the polyol (sorbitol) and hexoamine pathways of glucose metabolism with intracellular accumulation of their reaction end products, to an increase in intracellular glutathione and ascorbate redox systems, as well as disorders of the metabolism of NO and prostaglandins, non-enzymatic glycosylation of proteins and formation of glycosylation end products with the subsequent development of neuroinflammatory processes and cytotoxic edema of neuronal tissue (Kükürt *et al.*, 2021; Belenichev, 2013). Hypoxia observed in DM is an additional factor contributing to the increased formation of reactive oxidants. The accumulation of peroxidation products under conditions of hyperglycemia leads to the interaction of glucose with amino groups of proteins, increasing their glycosylation and oxidation (auto-oxidative oxidation). Non-enzymatic glycosylation of antioxidant defense enzymes leads to a decrease in their activity and even complete inactivation (Belenichev *et al.*, 2015). The greatest number of free radicals in the organism refers to combinations of reactive oxygen with a very short lifetime: the superoxide oxygen radical anion (O_2^-) and alkoxy radical (RO^*) – 10^{-6} s, hydroxyl radical (OH^-) – 10^{-9} s, peroxy radical (ROO) – 10^{-12} s (Collin, 2019; Edge & Truscott, 2021).

NITROXIDERGIC SYSTEM AND NITROSATIVE STRESS

The unique chemical nature and large number of intracellular targets for NO and its physiologically active oxidative-redox forms leave open the question of the way and the specificity in which the damaging effect of nitric oxide is mediated in the neuron under ischemic conditions. Numerous studies have shown the direct involvement of NO in the process of neuronal destruction in ischemia, arterial hypertension, and DM. A slight increase in NO concentration activates the synthesis of chaperone proteins and NO-dependent activation of HSP70 may constitute an important endogenous cellular defense mechanism. However, iNOS hyperexpression is suppressed by HSP70 by reducing the activation of the iNOS transcription factor (NF κ B), which leads to the limitation of nitrosative stress and neuroapoptosis (Belenichev *et al.*, 2015). Now, there is an active study of NO targets and whether NO is sufficiently cytotoxic, or whether its derivatives are more active (Aquilano *et al.*, 2011; Liu *et al.*, 2019). Studies in recent years have established that NO, and especially the products of its conversion, such as peroxynitrite (ONOO⁻), nitrosonium ion (NO⁺), nitroxyl (NO⁻) and diazotrioxide (N₂O₃), are major factors in the realization of nitrosative stress, which results in direct interaction of NO with metals (hem iron of hemoglobin, myoglobin, iron-containing enzymes, as well as non-heme iron of iron-sulfur proteins and DNA, copper and zinc of active enzyme centers), and indirect interaction of NO⁺ (S-, N-, O-nitrosation) with thiol, phenolic, hydroxyl and amino groups of proteins and DNA. Such interaction leads to receptor desensitization, inhibition of mitochondrial enzyme activity and fragmentation of nucleic acids (Belenichev *et al.*, 2015; Liu *et al.*, 2019).

GLUTATHIONE LINK OF THE THIOL-DISULFIDE SYSTEM

Hyperglycemia promotes the activation of the sorbitol pathway of glucose metabolism, which, together with activation of NADPH oxidase, leads to depletion of the NADPH cytosolic level and, consequently, of the reduced glutathione (GSH) level (Yan, 2018). (Fig. 2). A decrease in GSH levels below normal values can serve as an indicator of impaired cellular redox status and changes in the redox-dependent regulation of genes. The consequence of this disturbance is a significant change in the mechanism of cellular redox-dependent signaling, controlled both non-enzymatically and enzymatically with the participation of glutathione transferase and glutaredoxin isoforms (Luo *et al.*, 2016; Ohiagu *et al.*, 2021). It is known that GSH is a neurotransmitter and neuromodulator (in micromolar concentrations it is an agonist of glutamate receptors; in millimolar concentrations, it modulates the SH groups of NMDA receptors). Oxidized forms of glutathione in concentrations above 200 μ M decrease the expression of early response genes, and in concentrations of 5 mM or more, it activates p53-dependent apoptosis and reduces HSP levels (Belenichev *et al.*, 2020). GSH, competitively binding to nitric oxide, forms a complex in the form of S-nitrosoglutathione, which forms a depot of endogenous NO (further NO release is catalyzed by the thioredoxin system). Also, the release of NO from S-nitrosoglutathione occurs with glutamyltranspeptidase to form S-nitrosocysteinylglycine as a producer of NO. Cystine, which is

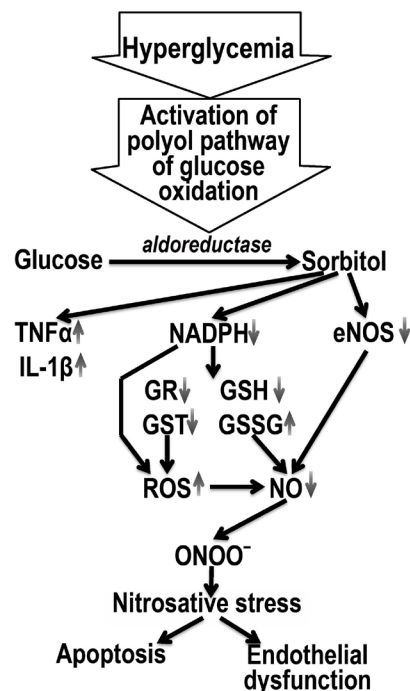


Figure 2. Disturbance in the coupling of NO/SH system in diabetes mellitus and nitrosative stress activation.

reduced to cysteine, takes part in the transport of S-nitrosoglutathione. These reactions are controlled by glutathione reductase and glutathione transferase. Nitrosative stress results in oxidative modification of low molecular weight thiols, formation of homocysteine and its cytotoxic derivatives which enhance thiol oxidation (Ren *et al.*, 2017; Aoyama, 2021; Belenichev *et al.*, 2015). It should be noted that the increased intracellular levels of cytotoxic forms of NO and decreased levels of reduced glutathione may be associated with deprivation of the heat shock protein (HSP70) level (Belenichev *et al.*, 2020). The action of NO formed with the participation of mtNOS results in the opening of mitochondrial pores and the release of pro-apoptotic proteins into the cytosol. The opening of the pores is due to oxidation or nitrosylation of the thiol groups of the cysteine-dependent portion of the mitochondrial inner membrane protein (ATP/ADP-antiporter), and this converts it into a permeable nonspecific pore channel (Pavlov *et al.*, 2017). The interaction of NO with members of the Bcl-2 superfamily is also reflected in the fact that the action of nitric oxide in the cell decreases significantly the level of intracellular Bcl-2 protein, possibly through caspase-induced splitting or p53-dependent inhibition of its expression (Shamas-Din *et al.*, 2013; Fricker *et al.*, 2018; Török *et al.*, 2002). The proapoptotic effect of nitric oxide is also expressed in its induced increase in the expression of apoptogenic Bax proteins. GSH and its precursor N-acetylcysteine can modulate NF- κ B, inhibit IL-1 β expression, and exhibit anti-inflammatory effects (Belenichev *et al.*, 2020; Pavlov *et al.*, 2017). It is known that increased production of TNF- α , IL-1 β , IL-6, and iNOS occurs against a background of GSH deficiency (Skelly *et al.*, 2013).

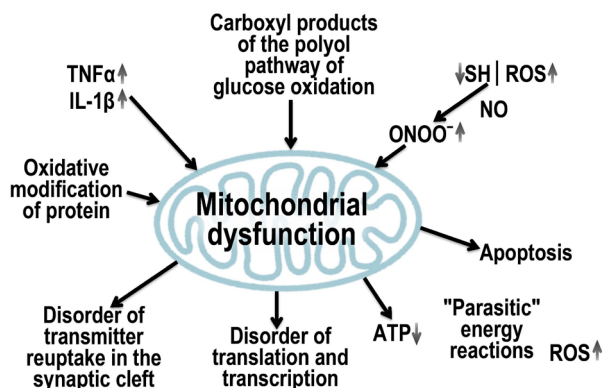


Figure 3. Formation of mitochondrial dysfunction in diabetes mellitus and its consequences.

MITOCHONDRIAL DYSFUNCTION

Mitochondrial dysfunction (MD) has no etiological and nosological specificity and is a typical pathological process. However, it leads to disruption of mediator reuptake (noradrenaline, dopamine, serotonin), ion transport, impulse generation and conduction, protein synthesis, processes of transcription and translation, and activation of “parasitic” energy-producing reactions, resulting in significant energy expenditure in the neuronal cell. The role of MD in the development of various pathological conditions, including neurodegenerative ones, has also been confirmed in DM (Norat *et al.*, 2020; Wang *et al.*, 2020; Pessoa & Duarte, 2023).

It was experimentally determined that diabetes mellitus leads to a significant decrease in the membrane potential of rat brain mitochondria. These changes were accompanied by a decrease in the ATP content and ATP/ADP ratio in brain synaptosomes, which indicates diabetes-induced disorders in the functioning of the electron-transport chain and energy coupling of the electron transfer process with ATP synthesis (Belenichev *et al.*, 2015; Pinti *et al.*, 2019; Singh *et al.*, 2021).

In conditions of hypoperfusion of brain tissues, compensatory mechanisms are depleted and energy deficiency develops, which leads to an increase in Ca^{2+} levels in the cell cytoplasm because energy-dependent pumps that “load” Ca^{2+} into the cisterns of the endoplasmic reticulum or “unload” it from the cell are blocked. These processes activate Ca^{2+} -dependent phospholipases. One of the defense mechanisms preventing the accumulation of calcium ions in the cytoplasm is their capture by mitochondria. However, it increases their metabolic activity aimed at maintaining intramitochondrial charge and proton pumping. This is accompanied by an increase in ATP expenditure. In general, a vicious circle occurs when oxygen deficiency disrupts energy exchange and stimulates the formation of ROS damaging mitochondrial and lysosome membranes, which can lead to mitochondrial dysfunction. In turn, mitochondrial dysfunction leads to the initiation of apoptosis and irreversible damage and death of the neuron (Norat *et al.*, 2020). In addition, the key enzyme of the Krebs cycle, aconitate hydratase (aconitase), is highly sensitive to the effects of oxidative and nitrosative stress (Pessoa & Duarte, 2023). In DM its activity is significantly reduced, and this leads to impaired mitochondrial glucose oxidation, and ATP deficiency (Sivitz & Yorek, 2010; Kuretu, 2023) (Fig. 3).

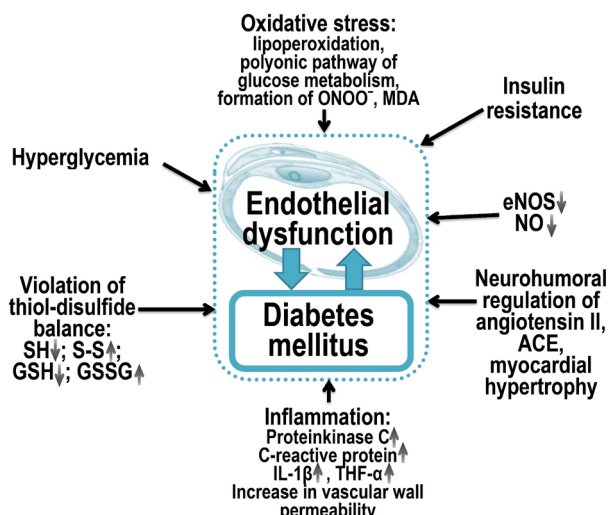


Figure 4. Pathogenesis of endothelial dysfunction in diabetes mellitus.

ENDOTHELIAL DYSFUNCTION

Hyperglycemia is a trigger in the pathogenesis of diabetic encephalopathy. As a result of protein kinase C activation, there is an increase in the concentration of endothelin-1 and production of growth factors: vascular endothelial growth factor VEGF, epidermal growth factor EGF and transformed growth factor TGF- β (Ergul, 2011; Heydarpour *et al.*, 2020; Wu & Derynck, 2009; Chen *et al.*, 2020). Also in hyperglycemia, there is an increased incorporation of glucose into the hexose-substituted pathway, resulting in increased transcription of inflammatory cytokine genes, which also contributes to the formation of vascular inflammation and proatherogenic state (Shi *et al.*, 2016; Wang *et al.*, 2020). Currently, oxidative stress is considered the main universal mechanism of development of all complications in DM, in particular, as a result of endothelial dysfunction. Moreover, hyperglycemia-induced oxidative stress triggers damage reactions to protein structures of ion channels and receptors, as well as activation and phosphorylation of cytosolic phospholipase A2 (cPLA2) with increased formation of arachidonate and prostaglandin E2, which leads to changes in vascular permeability (Wang & Hsiao, 2020; Sun *et al.*, 2021). As a result of oxidative stress, DNA damage occurs, an obligatory stimulus for the activation of the nuclease enzyme poly(ADP-ribose) polymerase, which depletes the intracellular concentration of NAD^+ , reducing the level of glycolysis, slowing electron transport and ATP formation, blocking glyceraldehyde-3-phosphate activity, which leads to endothelial dysfunction and development of diabetic complications (Liu *et al.*, 2017; Pacher & Szabó, 2005) (Fig. 4).

HEAT SHOCK PROTEINS

Clinical studies of HSP70 levels in diabetes are limited. It is known that patients with DM1 had elevated blood levels of HSP72, which decreased significantly after treatment (Ludwig *et al.*, 2014). However, another case-control study reported reduced serum HSP70 levels in type 1 diabetic patients with and without microvascular complications (Atalay *et al.*, 2004). Increased

serum HSP70 levels were also found in patients with type 2 DM not treated with insulin (Nakhjavani *et al.*, 2010). A decrease in iHSP70 expression and an increase in eHSP70 expression are found in patients with obesity and metabolic diseases, including DM2. HSF-1, which is one of the HSP72 transcription factors, is also repressed in subjects with DM2 (Seibert *et al.*, 2022). HSF-1 expression in skeletal muscle was found to be five times lower in obese and DM2 patients than in a control obese group.

The family of heat shock proteins HSPs (Heat shock proteins) is considered to be one of the most studied cytoprotective factors (Kim *et al.*, 2020; Deka & Saha, 2018; Belenichev *et al.*, 2022). There is a class of proteins (chaperones) whose main function is to restore the correct tertiary structure of damaged proteins, as well as to form and dissociate protein complexes. Many chaperones are heat shock proteins, that is, proteins whose expression is initiated in response to increased temperature or other cellular stresses (Belenichev, 2013; Ortan *et al.*, 2018). HSPs act as intracellular chaperones that maintain cell proteostasis in normal and under various stress conditions (hyperthermia, hypoxia, oxidative stress, radiation, etc.).

The most interesting is the HSP70 protein as an important component of the system of endogenous cyto- and neuroprotection, which, first of all, performs the function of intracellular chaperones and provides the processes of folding, holding and transport of synthesized proteins, as well as their degradation, both in normoxia and under stress-induced denaturation (Turturici *et al.*, 2011). It is known that the heat shock protein family 70 includes: inducible/stress-inducible block HSP72/HSP70i, constitutive/physiological protein HSP73/HSC70, constitutive glucose-regulating mitochondrial protein GRP75, constitutive heme oxygenase-1 (HO⁻¹) participates in bilirubin metabolism (Belenichev *et al.*, 2023). The constitutive form of HSP70 is still present in all subcellular compartments and participates in the functioning of cell life support systems in normoxia. On the contrary, the inducible form of HSP70 appears in cells in response to stress, including ischemic stroke (Turturici *et al.*, 2011). In response to stress, ischemia, hypoxia, etc., a sharp increase in the level of HSP70 is registered, and its highest concentration is observed in vital parts of cells: nuclear, perinuclear space, mitochondria, endoplasmic reticulum, which indicates the importance of chaperone70 in protecting cells from death. As nuclear pre-ribosomes resume functioning, the concentration of HSP70 in the nucleus decreases and increases in the cell cytoplasm. Thus, the level of HSP70 can be considered as a marker of cellular and tissue damage. Hyperproduction of HSP70 in cells inhibits the development of autophagy as an alternative, more “radical” mechanism of cellular stress response (Belenichev, 2013; Ortan *et al.*, 2018). Recent studies have established a direct cytoprotective effect of HSP70, which is realized by regulating the processes of apoptosis and cell necrosis. HSP70 inhibits mitochondrial and cytoplasmic pathways of apoptosis. Thus, HSP70 inhibits the transition of procaspase 9 into active caspase 9 and disrupts apoptosome formation in the cytoplasm of cells. Against the background of HSP70 hyperexpression the level of anti-apoptotic protein Bcl-2 increases, which prevents the release of cytochrome c from mitochondria and translocation of apoptosis-inducing factor (AIF) into the nucleus preventing cell apoptosis. HSP70 protein inhibits

TNF α -induced apoptosis and also effectively inhibits the development of Fas- and TRAIL- (TNF-related apoptosis-inducing ligand) mediated apoptosis in different cell types. Accumulation of HSP70 in cells increases their resistance to staurosporine, and doxorubicin, known as apoptosis inducers (Naka *et al.*, 2014). Ubiquitination of the insulin-like growth factor-1 receptor is inhibited by HSP10 and HSP60 which results in insulin-like growth factor-1 receptor signaling in cardiac muscle in streptozotacin-induced DM (Shan *et al.*, 2003). It is known that ischemia and hyperglycemia lead to the development of local inflammation. In this situation, HSP70 blocks the activation of the inflammatory transcription factor NF- κ B and inhibits its cytokine-mediated translocation to the nucleus (Belenichev, 2013). HSP70 inhibits the production of proinflammatory cytokines (TNF- α , IL-6), inhibits the activity of matrix metalloproteinases (MMPs) and inducible nitric oxide synthase (iNOS) in models of ischemia *in vitro* and *in vivo* (Belenichev *et al.*, 2015). It was found that HSP70 in astrocyte culture under ischemia inhibits proapoptotic Jun N-terminal kinases (JNK) and p38 mitogen activating protein kinase (MAPK), disrupting the apoptosis signaling pathway (Giffard *et al.*, 2008). The studies demonstrated a direct relationship between HSP70 and TNF α expression levels, iNOS in microglia, and MMP-9 in astrocytes. When IL-1b interacts with receptors, the nuclear transcription factors AP-1 and NF- κ B are activated, which alters the behavior of target cells and leads to the development of an acute cellular response, to the expression of other pro-inflammatory factors, to the stimulation of iNOS and cytotoxic NO produced by astrocytes, to an increase in mitochondrial pore permeability and to the initiation of neuroapoptosis. The IL-1b signaling pathway that enhances mechanisms of delayed neuronal death may be regulated by HSP70 (Kim *et al.*, 2020). An increase in HSP70 concentration within the physiological norm leads to an increase in IL-1 β to levels necessary for its participation in cyto- and neuroprotection; while HSP70 deficiency can lead to a significant increase in IL-1. Overexpression of HSP70 attenuates IL-1 β expression by inhibiting the C/EBP β and C/EBP δ transcription factors (Senf *et al.*, 2008). HSP70 can prevent the production of inflammatory cytokines by inhibiting NF- κ B-dependent transcription (Lyu *et al.*, 2020; Ferat-Osorio *et al.*, 2014). Many studies investigating the mechanisms of endogenous neuroprotection in ischemia show that the glutathione link of the thiol-disulfide system is reactivated against the background of an increase in the level of HSP70, and the introduction of exogenous HSP70 leads to an increase in the functional activity of the glutathione system (Belenichev *et al.*, 2020). HSP70 proteins mobilize antioxidant resources in neurons by increasing the level of both cytosolic and mitochondrial pools of reduced glutathione (Belenichev, 2013; Belenichev *et al.*, 2015; Zhang *et al.*, 2022). The data on the role of HSP in the stabilization of hypoxia-inducible factor (HIF-1a), which in conditions of ischemia is responsible for providing proliferation, apoptosis, angiogenesis, stabilization of protein molecules under oxidative stress, have recently emerged (Belenichev *et al.*, 2023). Under hypoxia conditions, HSP 70 is displaced from the complex with HIF-1a, thus, during 20–30 min of hypoxia, protecting the factor structure from targeted proteolysis. It is likely that HSP 70 is able to increase the lifetime of factor HIF-1a under pre- and post-hypoxia conditions and

is required for cells to respond appropriately to oxygen deprivation under ischemic conditions (Kim *et al.*, 2020). HIF-1 α determines the ability to activate the compensatory energy shunt and HSP 70 determines the ability of its long-term function. This statement is supported by the works of other researchers. It was found that one of the chaperones, the HSP 90 protein, is able to bind to the PAS domain of B-factor and stabilize it. Another cellular chaperone, HSP 70, recognizes a different structural motif of the HIF-1 α molecule, the so-called oxygen-dependent degradation (ODD) domain (Dery *et al.*, 2005). It should be noted that the role of these inter-protein interactions is unclear; it is assumed that they are required for the stabilization of HIF-1 α under normoxia conditions. Under hypoxia conditions, at least one of the chaperones (HSP 70) is displaced from the complex with HIF-1 α by the ARNT protein, which protects the factor structure from targeted proteolysis during 20–30 min of hypoxia. Thus, HSP 70 is able to increase the lifetime of the HIF-1 α factor under pre- and post-hypoxia conditions and is required for cells for an appropriate response to oxygen deprivation (Belenichev *et al.*, 2022; Belenichev *et al.*, 2023; Kim *et al.*, 2019).

Under DM, HSP levels are higher in some tissues and lower in other tissues. Defects in response to heat shock are seen in diabetic wounds. Both chaperones play important roles in cardiac defense. HSP60 expression is impaired in the heart of diabetic animals and this may contribute to diabetic cardiomyopathy (Atalay *et al.*, 2004). Also, expression of HSP72 is suppressed in streptozotacin-induced DM. A decrease of HSP70 was found in the mitochondria and cytosol of neurons in the CA1 hippocampus and sensorimotor cortex of rats with streptozotacin-induced DM (Chen *et al.*, 2013)15. Lower levels of HSP70 expression have been reported in insulin-sensitive tissues such as muscle and heart. There is evidence of decreased HSP70 protein levels in exercising diabetic animals with increased mRNA expression. Insulin resistance may contribute to a decrease in HSP70 levels in the heart and brain. It is hypothesized that increased expression of HSP70 in the brain is found in insulin sensitivity, in the brain as well as in peripheral tissues (Moin *et al.*, 2021).

HSP70 demonstrates diverse mechanisms of action on the development of inflammation and insulin resistance. Extracellular eHSP70 plays a role as a ligand for TLR2 and TLR4 in surrounding cells, which activates JNK *via* MEKK4/7 and suppresses NF- κ B through IRAK4 activation (Mulyani *et al.*, 2020; Zhang *et al.*, 2013). eHSP70 also regulates the expression of eNOS, iNOS and other cytokines such as TNF α and IL1 β (Belenichev *et al.*, 2023; Kim *et al.*, 2020; Deka & Saha, 2018; Giffard *et al.*, 2008). Activation of JNK leads to increased inflammation, triggers insulin resistance mechanisms and enhances the formation of mitochondrial dysfunction. In contrast, intracellular iHSP70 inhibits JNK activation and suppresses pathological mechanisms associated with its activation such as inflammation, insulin resistance, mitochondrial dysfunction, and ROS production (Bironaite *et al.*, 2012; Nagai & Kaji, 2023). The effect of HSP70 action depends on the iHSP70/eHSP70 ratio. A higher level of iHSP70 compared to eHSP70 leads to a decrease in inflammation processes (Krause *et al.*, 2015; Oliveira *et al.*, 2022; Seibert *et al.*, 2022; Alemi *et al.*, 2019).

LABORATORY DIAGNOSTICS

As a molecular marker in DM, it is necessary to determine the iHSP70/eHSP70 ratio in patients' serum or lymphocytes instead of total HSP70, which leads to a departure from the real picture of the pathological process. In healthy people, it is about 1 (Krause *et al.*, 2015; Seibert *et al.*, 2022). An [eHSP70]/[iHSP70] ratio of more than 5 indicates a significant pathological process – inflammation, insulin resistance, and endothelial dysfunction in DM (Mulyani *et al.*, 2020). The lower the ratio or the dynamics are more negative, the better the outcome of the disease.

PERSPECTIVES ON THE DEVELOPMENT OF PHARMACEUTICAL PRODUCTS

Such a role of chaperone proteins in cellular responses in pathological conditions raises the question of developing new drugs capable of providing modulation/protection of the genes encoding the synthesis of HSP 70 and HIF-1 α proteins. Strategies aimed at modulating HSP70 in DM (prolonged exercise, pharmacological modulation of HSP70) require adherence to and maintenance of iHSP70 expression, thus preventing progression toward more severe DM and allowing restoration of insulin sensitivity. Thus, HSP70 is attractive as a therapeutic target in the treatment of DM.

Declarations

Author contributions. IB designed the research. IB and OA wrote the original manuscript. OP and NB performed the literature search and revised the manuscript. All authors contributed to the article and approved the submitted version.

Conflict of interest. The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed a potential conflict of interest.

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