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Influence of melatonin on glutathione system in rats skeletal muscle under alloxan induced diabetes

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The aim was to determine the influence of melatonin on basal levels of glucose (BG), the levels of protein carbonyl content, HbA_{1c} and thiobarbituric acid reactive compounds (TBCRC), reduced glutathione (GSH), activities of glutathione reductase [EC 1.6.4.2] (GR), glutathione peroxidase [EC 1.11.1.9] (GPx), glucose-6-phosphate dehydrogenase [EC 1.1.1.49] (G-6-PhD) in the muscles of alloxan diabetic rats (DMIT). BG levels in blood of rats with DMIT increased on 139%, while in group of alloxan diabetic rats with impaired glucose tolerance (IGT) – were not differ from control. HbA_{1c} levels in the blood of animals with DMIT and IGT exceeded control by 219% and 123%, the level of protein carbonyl groups – by 76% and 36%, the level of TBCRC – by 58% and 36% respectively. Activities of GR, GPx, G-6-PhD and the level of GSH were decreased on 25%, 18%, 50% and 42% in rats with DMIT while in rats with IGT these indexes (besides GSH) were increased on 37%, 22%, 35% respectively than control. Melatonin lowered the BG level on 56% in DMIT rats in comparison to initial levels. It normalized activities of GR, GPx, G-6-PhD and lipid peroxidation, and protein carbonylation as well as hemoglobin glycosylation, so these indexes did not differ from control. Thus, melatonin has strong potential to regulate glucose homeostasis through enhanced glucose consumption, decreased oxidative stress by activation of the glutathione protection system.

Key words: melatonin; alloxan diabetes; glutathione system; skeletal muscle; rats.

INTRODUCTION

Diabetes mellitus can damage eyes, kidneys, nerves and heart. Microvascular and macrovascular disorders are the leading causes of morbidity and mortality in diabetic patients [1]. Hyperglycemia can increase the indicators of lipid peroxidation and oxidative stress in which free radicals fulfill the main role in the pathogenesis of these complications.

Alloxan diabetes was reported to induce oxidative stress and generates reactive oxygen species (ROS) [2]. In the presence of intracellular thiols, especially glutathione, alloxan generates ROS in a cyclic redox reaction with its reduction product, dialuric acid. Autoxidation of dialuric acid generates superoxide radicals, hydrogen peroxide and, in a final iron-catalysed reaction step, hydroxyl radicals. These hydroxyl radicals are ultimately responsible for the death of the beta cells, which have a particularly low

antioxidative defence capacity, and the ensuing state of insulin-dependent ‘alloxan diabetes’.

Type 1 diabetes mellitus (DMIT) is characterized by autoimmunity against pancreatic β cells, resulting in their destruction and the patients’ subsequent dependency on lifelong insulin replacement. DMIT patients have many complications, including cardiovascular, renal, and retinal disorders. Among them, skeletal muscle is a major target tissue of diabetic damage. Skeletal muscle is one of the largest organs in the human body and is, quantitatively, the most important tissue involved in maintaining glucose homeostasis under insulin-stimulated conditions. Type 1 diabetic subjects without insulin treatment display a dramatic loss of muscle, which leads to a higher blood glucose concentration, resulting in a vicious cycle [3].

Melatonin (N-acetyl-5-methoxytryptamine) is the major product of the pineal gland, which functions as a regulator of sleep, circadian

rhythm, and immune function. Melatonin and its metabolites have potent antioxidant/anti-inflammatory properties, and they have proven to be highly effective in a variety of disorders linked to inflammation and oxidative stress [4,5]. Melatonin not only neutralizes reactive oxygen species (ROS), but also acts through the stimulation of several antioxidative enzymatic systems and stabilizing cell membranes [6].

The influence of melatonin on glutathione system in muscles of alloxan-induced diabetic rats is poorly understood. The present study evaluated whether melatonin administration would have protective effect against muscle alloxan-induced changes in glutathione turnover.

The aim was to determine the influence of melatonin on basal levels of glucose (BG), the levels of protein carbonyl content, HbA_{1c} and thiobarbituric acid reactive compounds (TB-CRC), reduced glutathione (GSH), activities of glutathione reductase [EC 1.6.4.2] (GR), glutathione peroxidase [EC 1.11.1.9] (GPx), glucose-6-phosphate dehydrogenase [EC 1.1.1.49] (G-6-PhD) in the muscles of alloxan diabetic rats.

METHODS

Research performed in compliance with the Rules of the work using experimental animals (1977) and the Council of Europe Convention on the Protection of Vertebrate Animals used in experiments and other scientific purposes (Strasbourg, 1986), according to directions of International Committee of Medical Journals Editors (ICMJE), as well as “Bioethical expertise of preclinical and other scientific research conducted on animals” (Kyiv, 2006). Diabetes was induced in male Wistar rats by single i.p. injection of alloxan (170 mg/kg) [7]. Four days after diabetes induction, rats were divided into diabetic (untreated) and melatonin-diabetic group (10 mg/kg «Sigma» USA, daily and orally for 42 days starting from 5th day) [8]. Among diabetic rats were rats with preserved normoglycemia (impaired glucose tolerance – IGT) and rats with diabetes mellitus (DM1T) BG ≥

8.0 mmol/l. Animals belonged to the group with IGT when they detected a statistically significant increase in the level of postprandial glycemia > 9.0 mmol/l (2 hours after food intake) as compared to the control rats, in which this score was within (7.6-9.0) mmol/l. Blood was taken from the tail vein on 4th and 47th day evaluate the BG level with the use of OneTouchUltra (LifeScan, USA). Animals were sacrificed at the 47-th day from the beginning of the experiment accordance with the ethical treatment of animals. The rectus femoris muscle tissue was quickly removed, rinsed in saline, blotted, weighed and homogenized. The homogenate, 5% in ice-cold 0,25 mM tris-HCl-buffer (pH 7.4), was made using a homogenizer. The supernatant of the homogenate, prepared by ultracentrifugation for 10 min at 3000g was used for measurement of enzyme activities. Oxidant status was assessed by measuring of protein carbonyl contents, TB-CRC, GSH levels, GR, GPx, G-6-PhD activities and in addition the level of HbA_{1c} in the blood was determined. Determinations of the enzymes activities were by standard methods, concentration was measurement by spectrophotometer SP-46 / photocolorimeter KFK3 [9].

In the process of oxidative modification of proteins in the radicals of the aliphatic amino acid residues, aldehyde and ketone groups are formed. They interact with 2,4-dinitrophenylhydrazine (2,4-DNPH) to form 2,4-dinitrophenylhydrazones with a specific absorption spectrum. Aldehyde- and keto-derivatives which are neutral in nature are determined at a wavelength of 370 nm, alkali – at 430 nm [10].

The composition of the incubation medium consisted of 0.8 ml of 0.9% NaCl, 0.2 ml of 5% supernatant of muscle homogenate, 1ml of 1 M 2,4-DNPH, 1 ml of 10% trichloroacetic acid (TCA) solution. After incubation (1 hour at 37°C) samples were centrifugated at 1800 g for 10 minutes. Formed sediment was washed three times by 5% TCA solution. To the resulting precipitate was added 5 ml of 8 M urea solution, samples were kept in a boiling water bath until the precipitate dissolved. The level of protein

carbonyl contents was calculated by using an absorption coefficient of $2.1 \times 10^4 \cdot \text{M}^{-1} \text{cm}^{-1}$. The results were expressed in terms of optical density per gram of tissue (OD / g tissue).

The method of TBCRC determination [11] is based on a spectrophotometric determination of the trimetinic colored complex formed from the TBCRC interaction with thiobarbituric acid. The composition of the reaction medium was: 1 ml of 10% post-native supernatant of muscle homogenate; 1.5 ml distilled H_2O ; 0.2 ml of $20 \mu\text{M}$ FeSO_4 ; 1 ml of 0.8% TBC solution and 0.3 ml of 60% TCA solution. The optical density of the colored solution was measured on a photo-electrocolorimeter at a wavelength of 532 nm. TBCRC content ($\mu\text{mol/g}$) is calculated based on the molar absorption coefficient = $1.56 \times 10^5 \text{M}^{-1} \cdot \text{cm}^{-1}$.

The spectrophotometric/microplate reader assay method for GSH involves oxidation of GSH by the sulfhydryl reagent 5,5'-dithio-bis(2-nitrobenzoic acid) (DTNB) to form the yellow derivative 5'-thio-2-nitrobenzoic acid (TNB), measurable at 412 nm [12].

The composition of the incubation medium of the experimental test consisted of: 0.1 ml of 0.001 M solution of the Elman reagent (4 mg DTNB in 10 ml of 0.2 M potassium sodium phosphate buffer, pH 8.0), 3 ml of 0.2 M potassium sodium phosphate buffer (pH 8.0), 0.1 mM GSH solution, 0.3 ml of supernatant. The reaction was stopped by the addition of 10% TCA solution in 10 minutes. Measurements were made against a control sample (without a supernatant) at a wavelength of 412 nm. Values were expressed in micromol per 1 g of tissue.

The activity of GR was determined [12] by the rate of glutathione recovery in the presence of NADPH. The GR activity was determined in a supernatant (1500 g, 10 min) by decreasing the amount of NADPH in an incubation medium (3 ml) of the following composition: 50 mM Tris-HCl buffer (pH 7.5), 1 mM EDTA, 0.16 mM NADPH, 1 mM GSSG and 0.1 ml of 5% supernatant. The activity of GR was expressed in micromol NADPH used in response to 1 mg protein in 1 min.

The activity of GPx was determined [12] by the amount of oxidized glutathione formed from reduced glutathione in the detoxification of hydrogen peroxide in the glutathione peroxidase reaction.

The composition of the incubation medium consisted of 2.7 ml tris-HCl buffer (50 mM pH 7.4, sodium azide 12 mM, EDTA 6 mM), 0.1 ml of 2.5 mM reduced glutathione, 0.1 ml of 5% post-native supernatant of the muscles. The reaction was started by adding of 0.1 ml 0.5 mM hydrogen peroxide to the sample and stopping after 5 minutes by adding 1 ml of 10% TCA solution. Control sample was stoped before the reaction has begun. After centrifugation at 1800 g for 15 minutes in a supernatant of muscle homogenate, the optical density of oxidized glutathione at 262 nm was measured on a spectrophotometer. The activity of the enzyme was expressed in nanomol of the formed oxidized glutathione for 1 minute per 1 mg of protein.

The investigation of G-6-PhD activity was made [13] spectrophotometrically according to increase of the optical density at 340 nm, which is due to the rise in the number of NADPH in the process of enzymatic reaction.

Final concentrations of the components of the reaction mixture: 6.22 mM NADP, 10 mM G-6-Ph. To the centrifuge tube: 2.6 ml of 50 mM Tris-HCl buffer (pH = 7.4) containing 0.1 ml of magnesium sulfate (3 g of magnesium sulfate diluted in 25 ml of distilled water); 0.1 ml NADP; 0.1 ml of G-6-Ph; 0.1 ml of 5% muscle homogenate prepared on 50 mM Tris-HCl buffer (pH = 7.4). Samples incubated for 15 minutes in a thermostat at 37°C. The reaction stopped by the addition of 1 ml of 10% TCA solution. Samples are centrifuged for 10 minutes (3000 g). The activity of G-6-PhD in the muscles of rats was expressed in nanomol in 1 min per 1 mg of protein.

The blood samples were collected in EDTA anticoagulation bottles and sent cooled at 4°C. Determinations of HbA_{1c} were performed within three days using a kit Bio-Rad Laboratoria Inc., France.

Total protein determination (according to Lowry). It is performed according to the process described by V. Gudumac and coauthors [11].

Statistical analysis was performed using Statistica 10 StatSoft Inc. To determine an adequate method of statistical estimation of the average difference between the study groups held preliminary check distribution quantities in samples. According to the criteria Shapiro-Wilk, which is used to assess the normality of distribution in the sample volume $n \leq 50$, all samples not received data on deviation of the distribution of samples from normal ($P > 0.05$). Given these data, the use of Mann-Whitney test was considered sufficient for valid conclusions. Differences were considered to be statistically significant at $P \leq 0.05$ [14, 15].

RESULTS AND DISCUSSION

The blood glucose level (fig.1) of diabetic rats increased significantly by

139% ($P=0.000079$) compared with control value throughout the experimental period (42 days). Melatonin injections caused a sharp decrease by 56% ($P=0.00009$) in the elevated serum glucose level in DM1T group of rats compared with BG level before treatment.

ROS reacts with some amino acid, producing anything from modified, denatured and non-functioning proteins that in further may be responsible for oxidative stress. Diabetic

hyperglycemia, by the process of free radical production, causes protein glycation and oxidative degeneration. The degree of such protein glycation is estimated by using some biomarkers such as glycated hemoglobin. Alteration in function and structure of antioxidant protein enzymes may also be due to nonenzymatic glycation [16].

According to our own investigations (fig.2), HbA_{1c} levels in the blood of animals with DM1T and IGT exceeded control by 219% ($P=0.00003$) and 123% ($P=0.00026$) respectively. In rats with alloxan diabetes, the administration of melatonin during 42 days resulted in a decrease of HbA_{1c} level: in animals with DM1T by 45% ($P=0.00011$) compared to group of animals that did not receive correction; in rats with IGT, the normalization of this indicator was observed.

Possible link between melatonin and insulin interaction may be in its protective effect against free radical attack of β -cells Langergans islets in pancreas. In witness of such think we did the investigations of Langergans islets in our previous studies [14]. Histomorphological alterations in Langergans islets of pancreas in diabetic rats were recorded: their share reliable decreased by 55%, numbers of beta-cells decreased by 90%, percentage of beta-cells with necrosis was 97% respectively compared with the indices of control animals. Melatonin treatment caused a sharp decrease in the elevated serum glucose and partial regeneration/proliferation of β -cells of islets. It is concluded that the hypoglycemic

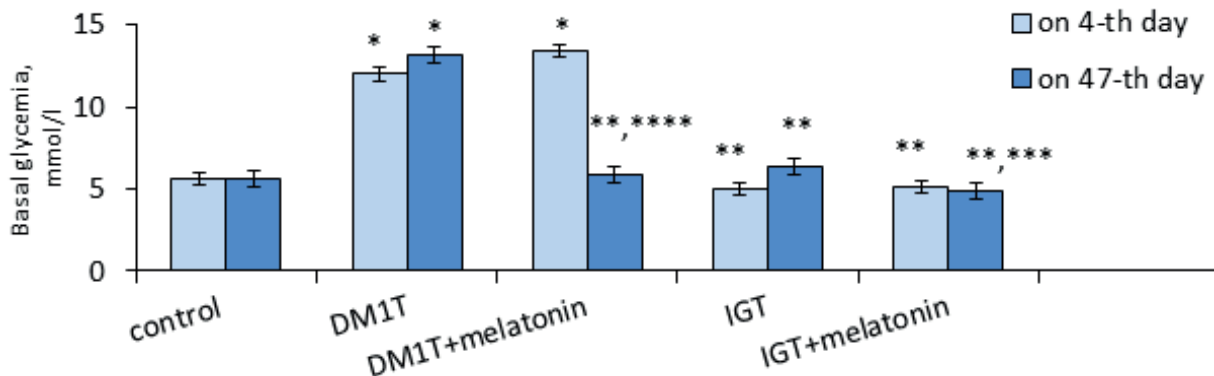


Fig. 1. The level of basal glycemia (mmol/l) in blood of rats, ($n=6$, $\bar{x} \pm Sx$): 1. *, **, ***, **** – changes are reliable ($P \leq 0.05$). 2. * – concerning control; ** – concerning rats with DM1T; *** – concerning rats with IGT; **** – concerning indices on 4-th day

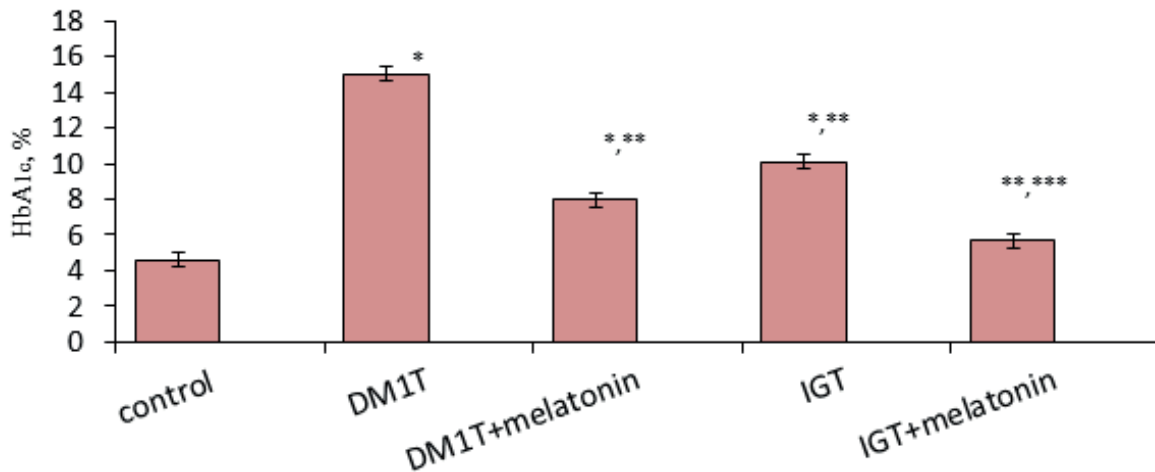


Fig. 2. The level of HbA_{1c} (%) in blood of rats, (n=6, x±Sx): 1. *, **, ***, **** – changes are reliable (P≤0.05). 2. * – concerning control; ** – concerning rats with DM1T; *** – concerning rats with IGT

action of melatonin could be partly due to amelioration in the beta-cells of pancreatic islets.

Results [17] indicate that melatonin treatment suppresses autoimmune recurrence by inhibiting the proliferation of Th1 cells in non obese diabetic mice and thus prolongs the survival of syngeneic islet grafts.

To access the protein oxidation mediated by glycation process, the levels of protein carbonyl content (fig. 3) were measured for the duration of 42 days.

The level of protein carbonyl groups was significantly increased in DM1T by 76% and

in IGT by 36% compared with control, whereas melatonin treatment significantly suppressed an increase in protein carbonyl content. When comparing with index of diabetic rats, the percentage reduction of carbonyl content by melatonin was found to be 42% (P=0.00044) and was not differ from control.

According to the results obtained, it may be postulated that melatonin inhibits glycation by reducing the generation of reactive carbonyl or dicarbonyl groups either from fructosamine or glucose, probably due to stimulation of glucose transport to skeletal muscle cells [18]

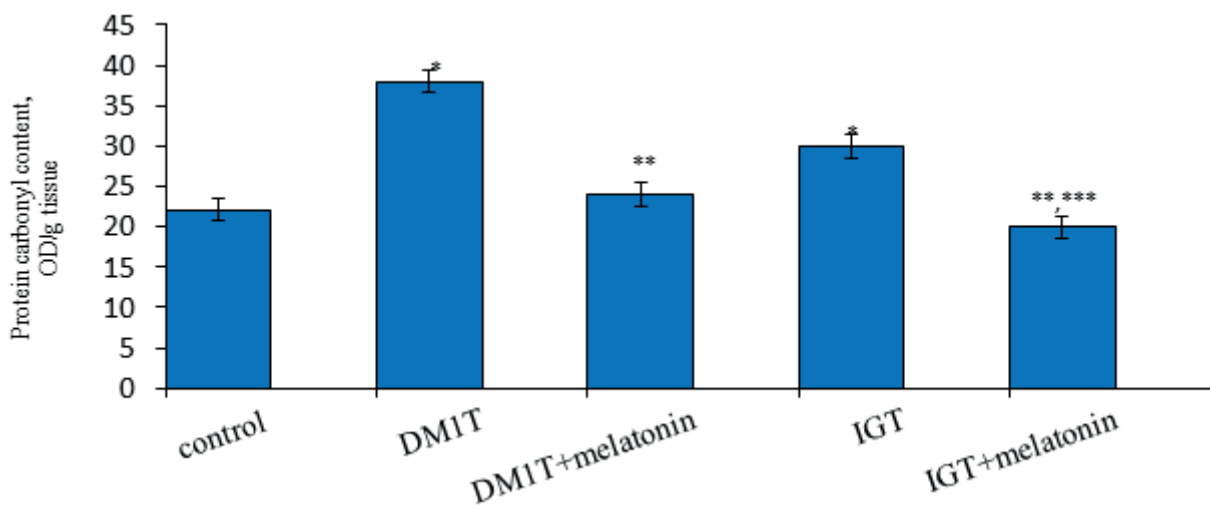


Fig. 3. The levels of protein carbonyl content in muscles of rats (OD / g tissue), (n=6, x±Sx): 1. *, **, ***, **** – changes are reliable (P≤0.05). 2. * – concerning control; ** – concerning rats with DM1T; *** – concerning rats with IGT

and preventing of ROS formation in conditions of hyperglycemia [16]. Diabetes mellitus produces disturbances in the lipid profile of body making the cells more susceptible to lipid peroxidation. Experimental studies [16] show that polyunsaturated fatty acids in cell membrane are extremely prone to attack by free radicals due to the presence of double bonds. Lipid peroxides through intermediate radical reactions produce such fatty acids that generate highly reactive and toxic lipid radicals that form new LHP. A critical biomarker of oxidative stress is Lipid peroxidation which is the most explored area of research when it comes to ROS. TBCRC are formed as a result of lipid peroxidation that can be used to measure lipid peroxides after reacting it with thiobarbituric acid. The level (fig.4) of TBCRC was found to be higher on 58% ($P=0.00089$) in DMIT group and on 36% ($P=0.0053$) in IGT group respectively than in control on 47th day of experiment.

So, the lipid peroxidation was increased in diabetic muscles. Melatonin partly prevented diabetes-induced increase in TBCRC in muscles.

Correspondingly, changes in the normal fiber type distribution are accompanied by changes in fuel oxidation and metabolic capacity of the muscle. Due to the reduced ability of skeletal muscle to access carbohydrates in times of inadequate/low insulin, diabetic skeletal muscle

must promote the use of other fuel sources. Skeletal muscle of individuals with DMIT is associated with the excessive deposition of intramyocellular lipids (IMCL). This high level of IMCLs is noted in the muscle following food consumption, and very low levels in the fasted state, as this fuel source is heavily relied upon. Muscle from the streptozotocin DMIT mouse model also demonstrates increased acetyl CoA/CoA ratio, hypothesized to be due to increased fatty acid oxidation, as well as increased fat utilization and mobilization, as the muscle tries to deal with the increased fat content. The alloxan-induced DMIT model similarly demonstrates an increase in free fatty acid levels in cardiac and skeletal muscle tissues. It is believed that as the levels of IMCL deposition increase, lipotoxicity ensues, enhancing stress to the tissue [19].

Diabetes induces alterations in activity of enzymes glutathione peroxidase and glutathione reductase (table). These enzymes are found in cell that metabolizes peroxide to water and converting glutathione disulfide back into glutathione. Any alteration in their levels will make the cells prone to oxidative stress and hence cell injury. So, the biochemical function of glutathione peroxidase is to reduce lipid hydroperoxides to their corresponding alcohols and to reduce free hydrogen peroxide to water.

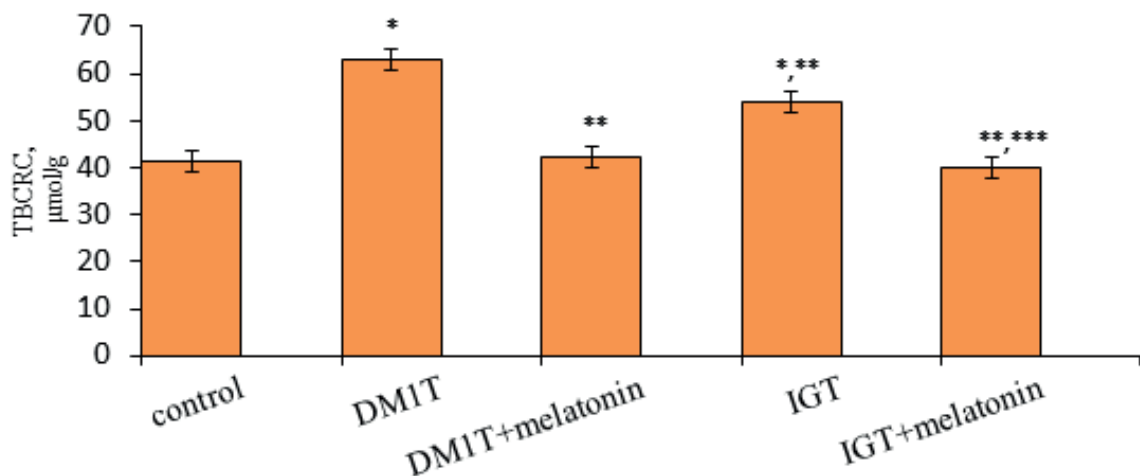


Fig. 4. The levels of TBCRC in muscles of rats ($\mu\text{mol/g}$), ($n=6$, $\bar{x}\pm Sx$): 1. *, **, ***, **** – changes are reliable ($P\leq 0.05$). 2. * – concerning control; ** – concerning rats with DMIT; *** – concerning rats with IGT

Table. Changes of the antioxidant defence in muscles of diabetic rats, (n=6, x±S)

Indexes Groups	GPx, nmol/min×mg	G-6-PhD, nmol/min×mg	GR, nmol/min×mg	GSH, μmol/g
Control group	40.4±2.12	3.6±0.05	0.5±0.04	0.14±0.01
DM1T	32.3±3.10 ^a	1.8±0.06 ^a	0.3±0.06 ^a	0.08±0.02 ^a
DM1T + melatonin	43.2±3.04 ^b	3.8±0.08 ^b	0.6±0.06 ^b	0.15±0.02 ^b
IGT	49.3±3.01 ^{a,b}	4.9±0.04 ^{a,b}	0.7±0.04 ^{a,b}	0.16±0.03 ^b
IGT + melatonin	45.0±2.08 ^b	3.9±0.08 ^{b,c}	0.5±0.04 ^{b,c}	0.15±0.02 ^b

Note: 1. a, b, c - changes are reliable ($P \leq 0.05$).

2. a – concerning control (intact rats);

b – concerning rats with diabetes mellitus;

c – concerning rats with IGT

G6PD reduces NADP to NADPH while oxidizing glucose-6-phosphate. The NADPH in turn maintains the level of GSH in cells that helps protect cells against oxidative damage from compounds like peroxides. On the other hand GR, GPx, G-6-PhD activities also depend on the presents of hyperglycemia. In DM1T group of rats activities of GR, GPx, G-6-PhD were decreased on 25% ($P=0.0077$), 18% ($P=0.036$), 50% ($P=0.000014$) respectively compare with control rats. These results are consistent with the degenerative role of hyperglycemia on cellular reducing equivalent homeostasis and antioxidant defense, and provide further evidence that pharmacological intervention of antioxidants may have significant implications in the prevention of the prooxidant feature of diabetes and protects redox status of the cells.

In the group of rats with preserved normoglycemia (IGT) activities of GR, GPx, G-6-PhD were increased on 37% ($P=0.0076$), 22% ($P=0.02$), 35% ($P=0.000022$) respectively compare with control rats. Increase of G6PhD activity in condition of diabetes with IGT is probably a compensatory reaction aimed to reduce of ROS.

NADPH reducing equivalents (that are produced in this reaction) are used for regeneration of glutathione from its oxidized form due to action of NADPH-dependent glutathione reductase. Glutathione neutralizes ROS, both directly and through GPx. We have found the level of GSH lower by 42% ($P=0.01$)

in DM1T group of animals compared with control. Melatonin injections was helpful for normalization this index under study.

It is well known, that muscle tissue is dependent on the presence of insulin. In the absence of insulin glucose cannot even enter muscle cells. In the conditions of low glucose influx to muscles there is no substrates for glycolysis, hexose monophosphate shunt and glycogenesis. That's why the activity of G6PD is decreased. We know [18] that pinealectomy, same as its hypofunction caused by permanent lighting, leading to decreased synthesis and secretion of melatonin, which causes insulin resistance and reduce the gene expression of glucose transporter GLUT 4, 2, 1. Overexpression of GLUT4 in skeletal muscle improves glucose homeostasis in animal models of diabetes mellitus and protects against the development of diabetes mellitus. Thus, GLUT4 is an attractive target for pharmacological intervention strategies to control glucose homeostasis. It was detected, that melatonin stimulates glucose transport to skeletal muscle cells via insulin receptor substrate-1 / phosphoinositide 3-kinase (IRS-1/PI-3-kinase) pathway, which implies, at the molecular level, its role in glucose homeostasis and possibly in diabetes. Additionally, exposure to light at night and aging, both of which lower endogenous melatonin levels may contribute to the incidence and/or development of diabetes [17] It is logical that the activity of G6PD is

reduced under diabetes mellitus, whether an administration of melatonin leads to increase its activity, probably due to direct activation [18].

So, enhanced skeletal muscle and fundamental understanding of the biological process are critical for effective glucose homeostasis in metabolic disorders.

Melatonin injections helped to normalize parameters of antioxidant body defense. Physiologically, the key to effective glucose homeostasis is the hormone insulin and insulin sensitivity of target tissues. Enhanced skeletal muscle, by either intrinsic mechanism or physical activity, offers great advantages and benefits in facilitating glucose regulation.

Melatonin besides being safe, lowered the blood glucose significantly without any hypoglycemic effect on their normoglycemic counterparts. It increased glutathione peroxidase, glutathione reductase activities significantly and reduced lipid peroxidation (32%) and protein carbonylation (42%) as well as hemoglobin glycosylation (45%). It also increased the glucose-6-phosphate dehydrogenase activity in muscles of diabetic rats which plays a critical role in glucose homeostasis. Thus, melatonin has strong potential to regulate glucose homeostasis through enhanced glucose consumption, decreased oxidative stress by activation of the glutathione protection system.

The authors of this study confirm that the research and publication of the results were not associated with any conflicts regarding commercial or financial relations, relations with organizations and/or individuals who may have been related to the study, and interrelations of coauthors of the article.

О.Ю. Кушнір, І.М. Яремій, В.І. Швець, Н.В. Швець

ВПЛИВ МЕЛАТОНІНУ НА СИСТЕМУ ГЛУТАТІОНУ В СКЕЛЕТНИХ М'ЯЗАХ ЩУРІВ З АЛОКСАНОВИМ ДІАБЕТОМ

Вивчали вплив мелатоніну на базальний вміст глюкози, реактивних сполук тіобарбітурової кислоти (ТБКРС), карбонільних похідних білків, НbA_{1c}, глутатіону (GSH), ак-

тивність глутатіонредуктази [КФ1.6.4.2] (ГР), глутатіонпероксидази [КФ1.11.1.9] (ГП), глюкозо-6-фосфатдегідрогенази [КФ1.1.1.49] (Г-6-фДГ) у м'язах щурів з алоксановим діабетом (ЦД1Т). Слід відзначити, що базальний вміст глюкози БГ в крові щурів із ЦД1Т збільшився на 139%, тоді як у групі щурів з діабетом і порушенням толерантності до глюкози (ПТГ) - не відрізнявся від контрольних. Вміст НbA_{1c} у крові тварин з ЦД1Т та ПТГ перевищував контроль на 219 та 123%, карбонільних груп - на 76 та 36%, ТБКРС - на 58 та 36% відповідно. Активність ГР, ГП, Г-6-фДГ та вміст GSH були знижені відповідно на 25, 18, 50 та 42% у щурів з ЦД1Т, тоді як у щурів з ПТГ ці показники (крім GSH) були збільшені на 37, 22, 35% відповідно порівняно з контролем. Введення мелатоніну знизило базальний вміст глюкози на 56% у щурів ЦД1Т порівняно з вихідним значенням, що супроводжувалося нормалізуванням активності ГР, ГП, Г-6-фДГ, процесів перекисного окислення ліпідів, карбонілювання білка, глікозилювання гемоглобіну, оскільки показники не відрізнялися від контролю. Таким чином, мелатонін має видимий потенціал для регулювання гомеостазу глюкози через посилення її споживання, зниження окисного стресу через активацію глутатіонової системи антиоксидантного захисту.

Ключові слова: мелатонін; алоксановий діабет; система глутатіону; скелетні м'язи; щури.

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ВЛИЯНИЕ МЕЛАТОНИНА НА СИСТЕМУ ГЛУТАТИОНА В СКЕЛЕТНЫХ МЫШЦАХ АЛЛОКСАНДИАБЕТИЧЕСКИХ КРЫС

Изучали влияние мелатонина на базальное содержание глюкозы, НbA_{1c}, карбонильных производных белков и реактивных соединений тиобарбитуровой кислоты (ТБКРС), содержание восстановленного глутатиона (GSH), активность глутатионредуктазы [КФ1.6.4.2] (ГР), глутатионпероксидазы [КФ1.11.1.9] (ГП), глюкозо-6-фосфатдегидрогеназы [КФ1.1.1.49] (Г-6-фДГ) в мышцах крыс с аллоксановым диабетом (СД1Т). Базальное содержание глюкозы в крови крыс с СД1Т увеличилось на 139%, тогда как в группе крыс с диабетом и нарушением толерантности к глюкозе (НТГ) - не отличался в сравнении с контролем. Содержание НbA_{1c} в крови животных с СД1Т и НТГ превышало контроль на 219 и 123%, карбонильных групп - на 76 и 36%, ТБКРС - на 58 и 36% соответственно. Активность ГР, ГП, Г-6-фДГ и содержание GSH были снижены на 25, 18, 50 и 42% соответственно у крыс с СД1Т, тогда как у крыс с НТГ эти показатели (кроме GSH) были увеличены на 37, 22, 35% соответственно по сравнению с контролем. Введение мелатонина снизило базальное содержание глюкозы на 56% у крыс СД1Т по

сравнению с исходным уровнем, что сопровождалось нормализацией активности ГР, ГП, Г-6-ФДГ, процессов перекисного окисления липидов, карбонилирования белка, гликозилирования гемоглобина, так как показатели не отличались от контроля. Таким образом, мелатонин обладает сильным потенциалом для регулирования гомеостаза глюкозы за счет увеличения ее потребления, а также снижения окислительного стресса за счет активации глутатионовой системы защиты.

Ключевые слова: мелатонин; аллоксановый диабет; система глутатиона; скелетные мышцы; крысы.

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