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PPAR α and PPAR γ as main regulators of fatty acid metabolism

PPAR α i PPAR γ jako główne czynniki regulujące przemiany kwasów tłuszczowych

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Summary

PPAR α and PPAR γ nuclear receptors are transcription factors responsible for regulating the expression of lipid, carbohydrate and protein metabolism genes. PPAR receptors are distinguished by their capability to efficiently bind a number of synthetic or natural ligands. Genes encoding PPAR α and PPAR γ are expressed in many tissues and organs, including in the adipose tissue. Changes in the expression of these genes are dependent on many factors, such as the impact of other genes and fatty acids. PPAR α and PPAR γ proteins are - due to their neuroprotective and anti-inflammatory properties, as well as the key role they play in lipid metabolism - important for human health. This is especially true with regards to neurological disorders such as Alzheimer's disease or those accompanied by a metabolic syndrome - type II diabetes, insulin resistance, obesity and hypertension.

Keywords: fatty acid • metabolism • nuclear receptor • PPAR α • PPAR γ

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Abbreviations: **CLA** – conjugated linoleic acid; **CRP** – C-reactive protein; **DHA** – docosahexaenoic acid; **DNA** – deoxyribonucleic acid; **EPA** – eicosapentaenoic acid; **FABP4** – fatty acid binding protein 4; **FFA** – free fatty acids; **GPR40** – G protein-coupled receptor 40; **IL** – interleukin; **LBD** – ligand binding domain; **MAPK** – mitogen-activated protein kinases; **NF- κ B** – nuclear factor kappa B; **PPAR** – peroxisome proliferator-activated receptor; **PPRE** – peroxisome proliferator response element; **PUFA** – polyunsaturated fatty acids; **RXR** – retinoid X receptor; **SCD** – stearoyl-CoA desaturase; **TNF- α** – tumor necrosis factor alpha; **UCP1** – uncoupling protein 1; **UFA** – unsaturated fatty acids; **WAT** – white adipose tissue

INTRODUCTION

Peroxisome proliferator-activated receptors (PPARs) are nuclear receptors of the steroid receptor superfamily responsible for regulating the transcription of genes involved in metabolic or inflammatory processes pathways [76]. Three PPAR isotypes - PPAR α , PPAR β/δ and PPAR γ - expressed in different tissues have been distinguished [42]. PPAR γ , like all nuclear receptors, is of a modular structure, and contains three major functional domains. The N-terminal region domain is responsible for the transcription activation function, while the C-terminal region domain is an LBD (ligand binding domain); between them is a conservative DNA-binding domain (deoxyribonucleic acid) [76].

Peroxisome proliferator-activated receptors alpha and gamma – genes

PPAR family proteins are encoded by genes of the same name: PPAR α , PPAR β/δ , and PPAR γ . In the case of PPAR γ , two isoforms of the gene, PPAR γ_1 and PPAR γ_2 have been identified. They are generated by an alternative action of the same gene promoter, resulting in four mRNA variants: PPAR γ_1 , PPAR γ_2 , PPAR γ_3 , PPAR γ_4 . All four mRNA variants encode the PPAR γ_1 protein. PPAR γ_2 encodes PPAR γ_2 protein that is identical to PPAR γ_1 but has an additional 30aa sequence at the N-terminus [17].

PPAR α is present in the heart, liver, kidneys, muscles and brown adipose tissue. The PPAR γ_1 gene is primarily expressed in white and brown adipose tissue, as well as the large and small intestines, in immune system cells, muscle tissue, pancreas, liver and kidneys. In contrast, PPAR γ_2 expression is observed mainly in adipose tissue [42].

PPAR α gene in humans consists of 13 exons and is located on chromosome 22. In mice, it is located on chromosome 15 and consists of 16 exons, while porcine PPAR α gene has 8 exons and is located on chromosome 5. In chickens, PPAR α is located on chromosome 1 and is made of 13 exons. In ruminants, the gene encoding peroxisome proliferator-activated receptors alpha is found on chromosome 3 for sheep, while in cattle it is located on chromosome 5. It is formed by, respectively, 9 (sheep) and 11 (cattle) exons.

In humans, PPAR γ contains 11 exons and is located on chromosome 3. In mice, the gene consists of 15 exons and is located on chromosome 6. In cattle, the gene is located on chromosome 22 and is made of 7 exons. In the case of sheep, this gene - formed by 17 exons - is located on chromosome 19. Porcine PPAR γ consists of 13 exons and is found on chromosome 13, while in chickens it contains 9 exons and is located on chromosome 12.

The main factors influencing the expression of genes encoding PPAR proteins are fatty acids. It was found that the presence of CLA (conjugated linoleic acid) acid

in the diet of mice with metabolic syndrome led to an increase in PPAR γ gene expression [73]. Also, the action of docosahexaenoic acid (DHA) stimulates increased expression level of this gene [56]. In contrast, long-term action of eicosapentaenoic acid (EPA) leads to a significant decrease in PPAR γ expression in rat adipocytes [40]. Introduction of n-3 PUFA (polyunsaturated fatty acids) into the diet of ob/ob (leptin-deficient) mice results in increased expression of the gene encoding peroxisome proliferator-activated receptors gamma [20]. However, studies demonstrating the negative effect of n-3 PUFA on the expression of this gene are available as well. Such an outcome was observed in adult obese patients on a 12-week diet rich in DHA, who experienced a significant decrease in PPAR γ expression [44]. The use of n-3 PUFA-enriched goat meat in mice nutrition resulted in a significant increase in expression of the analyzed gene [16]. In addition, also differentiated diet of mother ewes during pregnancy induces changes in the expression of PPAR α and γ in the fetus, causing changes in lipid metabolism [9]. The demonstrated significant increase in PPAR γ gene expression in skeletal muscle of fetuses of obese ewes can serve as confirmation of increased adipogenesis in muscle tissue [86].

Identification of polymorphisms of genes encoding peroxisome proliferator-activated receptors alpha and gamma in humans is relevant in studies focusing primarily on the nervous system, obesity, metabolic syndrome or, to an extent, cardiovascular diseases [49, 60, 71, 74]. In livestock, genetic variability in the PPAR α , γ is chiefly related to the quality of meat, due to the role these genes play in protein coding in adipogenesis and fatty tissue storage processes [36, 38].

Peroxisome proliferator-activated receptors alpha and gamma – proteins

In animal cells, PPAR proteins interact with nuclear proteins acting as co-activators or co-repressors [53]. Ligand-unbound PPAR, by interacting with a co-repressor, remains deacetylated and inhibits gene expression [55]. Ligand-bound PPAR after binding to the ligand forms a heterodimer with the retinoid X receptor (RXR) and a co-activator with intrinsic histone acetylase activity (for example PPAR γ 1- α or EP300 co-activator). It then joins - in the gene promoter region - the peroxisome proliferator response element (PPRE) gene sequence, regulating its transcription.

PPAR genes change their conformation depending on the bound ligand. In the case of binding an agonist, they may interact with the co-activator. After binding a PPAR antagonist, such an interaction is impossible [55]. Wang et al. [72] indicated yet another mechanism characteristic for PPAR γ . According to them, a PPAR γ ligand, such as for example rosiglitazone (a drug used to treat diabetes), is capable of activating the G protein-coupled receptor 40 (GPR40). This leads to phosphorylation of p38 MAPK (mitogen-activated protein kinases), which

sequentially phosphorylates and activates PPAR γ 1- α and EP300 co-activators, and causes conformational changes of PPAR γ [72].

What distinguishes PPAR proteins from other nuclear receptors is their capability to bind many different ligands [29, 63]. Because of the size of the ligand binding site, PPAR – especially PPAR γ – can bind a variety of natural or synthetic lipophilic acids, in particular fatty acids and their derivatives – eicosanoids [33].

Natural PPAR receptor agonists, i.e. molecules that bind to the receptor and stimulate initiation of a cellular reaction, include mainly fatty acids, such as α -linolenic (C18:3 n-3) and linoleic (C18:2 n-6) acids, which are essential for proper functioning of the body. These acids cannot be synthesized endogenously, and thus must be supplied with food. They serve as substrates for the desaturation and/or elongation reactions that result in long-chain PUFAs. It is also known that linoleic acid, a member of the omega-6 group, is one of PPAR γ ligands [77, 81]. A metabolite of this acid, 13-hydroxyoctadecadiene acid is also a PPAR γ receptor agonist; it inhibits cellular signaling of inflammation and lowers secretion of IL-8 (interleukin) in colon epithelial cells [2, 11]. Vangaveti et al. [70] observed proapoptotic action of 13-hydroxyoctadecadiene acid in monocytes, while linoleic acid bound to the PPAR γ receptor did not exhibit such properties [70]. In contrast, anti-inflammatory effect of lipoic acid, belonging to the omega-3 PUFA group, consists in the activation of PPAR γ receptors, which leads to a decrease in IL-1, -6, and TNF- α (tumor necrosis factor alpha) synthesis [84].

Another group of PPAR ligands are eicosanoids, organic compounds that are products of oxidation of UFAs (unsaturated fatty acids) such as arachidonic acid, linoleic acid and α -linoleic acid released from the phospholipid membrane. This group of compounds includes prostaglandins, prostacyclins, leukotrienes and lipoxins [42]. Previously mentioned rosiglitazone is a synthetic eicosanoid demonstrating neuroprotective action - in rodent brain after stroke it induces 5-lipoxygenase and lipoxin A4, which has an anti-inflammatory effect [67]. Neuroprotectine D1, another PPAR ligand, also has demonstrated neuroprotective effects, and has been shown to have beneficial effects in mice with Alzheimer's disease [85]. Beneficial effects of D1 protectine have also been shown in other tissues. White et al. [75] demonstrated a significant increase in expression of PPAR γ and RXR (retinoid X receptor) genes within the fatty tissue of fat-1 transgenic mice as a result of a fat-rich diet, which proves that D1 protectine may affect the adipogenesis process through PPAR γ receptors [75].

By activating PPAR α , conjugated linoleic acid prevents inflammation and neoplasm in the colon. Similar anti-inflammatory effects have been observed in macrophages and small and large intestine epithelium under CLA supplementation [8, 22, 30, 80, 82]. Also vaccenic

acid acts via both the PPAR γ and PPAR α . It has been shown that this acid, present in butter, meat or milk in human diet, inhibits cardiomyocyte hypertrophy [73]. Another anti-inflammatory ligand acting via PPAR γ are nitrated fatty acids generated in non-enzymatic reactions of fatty acids and nitric oxide [50].

Peroxisome proliferator-activated receptors alpha and gamma – functions

PPAR receptors have several important functions in the body. In the brain, they are involved in the differentiation of nerve cells, reduction of oxidative stress and improvement of memory. In skeletal muscles they are responsible for, *inter alia*, increased fatty acid oxidation, and, in the small intestine, for the reduction of inflammatory conditions. In fatty tissue, PPARs are responsible for the differentiation of adipocytes and triglyceride synthesis, and they stimulate insulin sensitivity [42].

ROLE OF PPAR IN LIPID METABOLISM

Polyunsaturated omega-3 fatty acids are among natural ligands bound by peroxisome proliferator-activated receptors [43]. PPAR α activation by n-3 PUFA in the liver leads to strengthened expression of lipid metabolizing genes, which in turn leads to the intensification of fat catabolic processes [18].

With PPAR α knock-down in mice, it was shown that the lack of functional protein encoded by this gene results in the impairment of cellular metabolism under fasting conditions. Under the experimental nutritional conditions, this led to the accumulation of lipids in the liver and heart, hypoglycaemia, hypothermia, ketonuria and elevated levels of free fatty acids, and eventually to the death of the individual [34]. In contrast, wild phenotype mice adapted to starvation conditions by activating increased expression of the PPAR α gene in the liver and heart, resulting in, as was found, increased FFA (free fatty acids) uptake and increased oxidation [27]. The gene also influences changes in the expression of cytochrome P450 4A11, leading to observed stimulation of β -oxidation and ω -hydroxylation [59, 64].

The paradox of PPAR α action consists in increasing the intensity of fatty acid oxidation, while simultaneously engaging receptors in changes of expression of genes encoding enzymes responsible for lipogenesis, including SCD (stearoyl-CoA desaturase) [13, 47]. PPAR α is also responsible for the regulation of the gluconeogenesis process in the liver by controlling the activity of enzymes involved in these processes [59, 78]. It has been shown that CLA acid, a strong PPAR α agonist contained in tomato juice, is responsible for a decrease in triglyceride levels in the liver of obese diabetic mice [28].

PPAR γ ligands, for example glitazones used as components of insulin sensitizers, are supplied to patients suffering from type 2 diabetes. Bogacka et al. [10] observed

that the administration of pioglitazone to patients with type 2 diabetes results in a change in the expression of genes involved in carbohydrate and lipid metabolism in the subcutaneous adipose tissue [10]. However, gli-tazones differ for the normal functioning of the body, as a number of side effects have a negative impact on the function of the liver and heart [12, 65].

In contrast, PPAR γ stimulates the formation and differ-entiation of new adipocytes [18, 69] due to increased transcriptional activity of the gene resulting from it ligand-binding DHA acid [48]. The presence of rosigli-tazone leads to the activation of the peroxisome prolif-erator-activated receptors gamma, which in the case of the WAT (white adipose tissue) precursor cells leads to the intensified creation of mitochondria and increased expression of the *UCP1* (uncoupling protein 1) gene [57]. Studies conducted on mice with PPAR γ knockdown have shown that it is essential for adipogenesis as well as fat tissue formation [6, 31, 61]. In mature adipocytes, ligand binding and PPAR γ activation lead to the increased expression of genes involved in lipid and glucose metab-olism, and strengthen fatty acid oxidation and increased insulin sensitivity. In addition, PPAR γ stimulates adipo-cytokine production [18]. Acting via PPAR γ receptors, EPA stimulates the synthesis and secretion of adipo-nectin in adipose tissue cells [56]. The increase in adipo-nectin secretion, as a result of PPAR γ , together with increased insulin sensitivity and reduced inflammation, is observed with omega-3 fatty acid supplementation in mice with impaired leptin synthesis [20]. Neschen et al. [51] observed that the increased production of adipo-nectin as a result of introduction of PUFA n-3 contained in fish oil into the diet is directly dependent on PPAR γ and not on PPAR α [51].

Role of PPAR in inflammation processes

PPAR α receptors exhibit anti-inflammatory action by inhibiting the expression of genes encoding acute phase proteins, such as CRP (C-reactive protein) and fibrino-gen. They contribute to reducing the risk of liver inflam-mation, cardiovascular disease and cancer [19].

PPAR γ and nuclear factor kappa B (NF- κ B) are also involved in the regulation of inflammatory processes: NF- κ B controls the expression of a number of genes involved in the inflammatory process, and PPAR γ inter-acts directly with NF- κ B [23]. Chen et al. [14], by admin-istrating hepatitis-inducing concanavalin (a mitogen stimulating T lymphocytes), observed a decrease in cyto-kine secretion and increased NF- κ B activity due to high PPAR γ concentration [14]. The use in the same experi-ment involving a 12-week n-3 PUFAs-rich diet showed an additional increase in PPAR γ gene expression in the liver and heightened T-cell counts. These results confirm the anti-inflammatory effect of PPAR γ [37]. PPAR γ receptor reducing the synthesis of Il-6, -8 is activated also to pre-vent inflammation in the intestine [41]. Studies in mice with induced colitis showed that a diet with low n-6:n-3

PUFA ratio effectively reduces inflammation via PPAR γ action [24]. Similar results have been obtained for breast cancer tumor cells [83].

Role of PPAR in cardiovascular diseases

Due to PPAR γ function in the regulation of inflamma-tion, lipid metabolism, oxidative stress and cellular apoptosis, these receptors interact with the cardiovas-cular system [26]. Activation of PPAR γ by vaccenic acid and of PPAR α , - γ by conjugated linoleic acid favorably impacts the development of cardiomyocytes, inhib-iting their hypertrophy [1, 73]. Beneficial effects of PPAR receptors on the cardiovascular system have also been demonstrated in patients following a diet rich in n-3 PUFA [52]. Anderson et al. [3] reported that the administration of a fluid enriched in DHA and EPA to patients three weeks prior to cardiac surgery increased PPAR γ activity in the myocardium atrium. In addition, increased fatty acid oxidation in the mitochondria and intensified activity of key enzymes catalyzing antioxi-dant and anti-inflammatory processes have both been demonstrated [3].

Tian et al. [68] observed that a decreased expression of the PPAR γ gene causes changes in expression of genes responsible for the development of pulmonary hyper-tension. Thus, this decline in gene expression is linked to the induction of hypertension in pulmonary circula-tion [68]. Changes in PPAR γ activity are not only associ-ated with changes in blood pressure, but also with the occurrence of vascular diseases [53, 54]. Studies in lambs have confirmed that irregularities in PPAR γ activity may lead to damage to blood vessels [66].

Protective role of PPAR

Krämer et al. [32] demonstrated that PPAR α expression is directly related to the quantity of type I fibers in skele-tal muscles [32]. These receptors are, through signaling pathways, responsible for increased resistance to dam-age caused by ischemia of the heart, liver, or muscle tis-sue [5]. Studies in post-liver transplant patients have demonstrated that polymorphisms within the donor's PPAR α gene increase susceptibility to metabolic distur-bances [39].

PPAR family receptors are a very important factor in the proper functioning of the nervous system and cell signaling. Active PPAR γ exhibits neuroprotective prop-erties – such action has been demonstrated in experi-ments where microglial cells were treated with a toxic lipopolysaccharide, as well as in studies focused on the action of neuroprotectin D1 [4, 67, 85]. Recent study findings also indicate the possible uses of PPAR γ in post-traumatic neuronal regeneration [35]. Attempts are made to make use of PPAR α and - γ neuroprotec-tive properties in interaction with various agonists in the treatment of degenerative neurological diseases – Parkinson's and Alzheimer's [7, 15, 58, 62]. In addition,

PPAR α activity has been shown to be associated with sleep modulation [46].

Studies focused on age-related macular degeneration in humans suggest that increased expression of the PPAR γ gene influences the development of this disease [21]. *In vivo* studies show that the use of n-3 PUFA-rich diet leads to an increased expression of this gene in the mouse model of age-dependent macular degeneration [79].

ROLE OF PPAR IN SHAPING FAT CONTENT AND QUALITY OF PRODUCTS OF ORIGIN PRODUCTS

Activation of PPAR γ receptors by chloroformic multiclorigide extract (*Lolium multiflorum*) induces cell proliferation and differentiation of adipocytes as well as fat deposition, which directly influences the quality of animal meat [25]. Also, a link between the expression level of the gene encoding peroxisome proliferator-activated receptors gamma and fat deposition has been observed [44]. Evaluation of transcriptome profile of

bovine muscle tissue has demonstrated that an increase in intramuscular fat content in beef [38] is connected to expression levels of 10 genes, including *SCD* and *FABP4* (fatty acid binding protein 4) that are regulators in the PPAR signaling pathway.

CONCLUSION

Peroxisome proliferator-activated receptors, due to their functions, seem to be one of the key factors responsible for the conversion of fatty acids and lipid metabolism, and thus also for the energy balance of the body. Particular emphasis should be placed on their effects in the case of the metabolic syndrome and its associated conditions: diabetes, obesity or hypertension, increasingly widespread in the twenty-first century. The significant number of studies and publications focusing on the role of expression and genetic variability of the genes, proteins and ligands of PPAR receptors in the metabolism of animal and human fat is thus a natural response to their significance for these processes.

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