

Impact and Therapeutic Potential of PPARs in Alzheimer's Disease

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Abstract: Peroxisome proliferator activated receptors (PPARs) are well studied for their role of peripheral metabolism, but they also may be involved in the pathogenesis of various disorders of the central nervous system (CNS) including multiple sclerosis, amyotrophic lateral sclerosis, Alzheimer's and, Parkinson's disease. The observation that PPARs are able to suppress the inflammatory response in peripheral macrophages and in several models of human autoimmune diseases, lead to the idea that PPARs might be beneficial for CNS disorders possessing an inflammatory component. The neuroinflammatory response during the course of Alzheimer's disease (AD) is triggered by the deposition of the β -amyloid peptide in extracellular plaques and ongoing neurodegeneration. Non-steroidal anti-inflammatory drugs (NSAIDs) have been considered to delay the onset and reduce the risk to develop Alzheimer's disease, while they also directly activate PPAR γ . This led to the hypothesis that NSAID protection in AD may be partly mediated by PPAR γ . Several lines of evidence have supported this hypothesis, using AD related transgenic cellular and animal models. Stimulation of PPAR γ by synthetic agonist (thiazolidinediones) inducing anti-inflammatory, anti-amyloidogenic and insulin sensitizing effects may account for the observed effects. Several clinical trials already revealed promising results using PPAR γ agonists, therefore PPAR γ represents an attractive therapeutic target for the treatment of AD.

Keywords: Neuroinflammation, alzheimer's disease, PPAR, thiazolidinediones.

INTRODUCTION

The peroxisome proliferator activated receptors (PPARs) belong to the family of nuclear hormone receptors (NHR) that comprise 48 human ligand-inducible transcription factors which activity is regulated by steroids and lipid metabolites [reviewed in 1]. Three different PPAR genes (PPAR α , PPAR β , also called δ , and PPAR γ) have been identified in all metazoa, showing an unique spatio-temporal tissue-dependent expression pattern during fetal development in a variety of cell types deriving from the ecto-, meso- or endoderm in rodents. Functionally, PPARs are involved in adipocyte differentiation, lipid storage, and glucose homeostasis in all most all organs including the adipose tissue, brain, placenta and skin [reviewed in 2].

Functions of PPARs

PPARs act principally as lipid sensors and regulate whole body metabolism in response to dietary lipid intake and direct their subsequent metabolism and storage [3]. The prototypic member of the family, PPAR α , was initially reported to be induced by peroxisome proliferators, and now denotes the subfamily of three related receptors. The natural ligands of these receptors are dietary lipids and their metabolites. The specific ligands have been difficult to establish, owing to the relatively low affinity interactions and broad ligand specificity of the receptors.

PPAR α acts primarily to regulate energy homeostasis through its ability to stimulate the breakdown of fatty acids and cholesterol, driving gluconeogenesis and reduction in

serum triglyceride levels. This receptor acts as a lipid sensor, binding fatty acids and initiating their subsequent metabolism. PPAR γ binds a number of lipids including fatty acids, eicosanoids and other natural lipid ligands. Its dominant action is to stimulate adipocyte differentiation and to direct lipid metabolites to be deposited in this tissue. PPAR γ operates at the critical metabolic intersection of lipid and carbohydrate metabolism. PPAR γ activation is linked to reduction in serum glucose levels, likely as a secondary effect of its ability to regulate endocrine factors. It is this latter activity that has led to the development of specific PPAR γ agonists for the treatment of type II diabetes [4]. PPAR β/δ binds and responds to VLDL-derived fatty acids, eicosanoids including prostaglandin A1 [5] and appears to be primarily involved in fatty acid oxidation, particularly in the muscle.

PPARs regulate gene expression by forming heterodimers with retinoid-X-receptors (RXRs). Stimulation of target gene expression is controlled by specific PPAR-response elements in the promoter region (PPREs). Under unstimulated conditions these heterodimers are associated with corepressors, like N-CoR and SMRT, which suppress gene transcription [2]. Upon ligand binding to the nuclear receptor, the corepressors are displaced and transcriptional coactivators are recruited to the receptor. These coactivator:receptor complexes finally induce the formation of a much larger transcriptional complex which subsequently links the basal transcriptional apparatus and initiates transcription of specific target genes. In addition, activity of PPARs in general is also regulated by posttranslational modification such as phosphorylation and sumoylation [6,7].

There are several mechanisms involved in PPAR γ inactivation. Thus, phosphorylation can negatively or positively affect PPAR γ activity depending on which specific protein

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residue is modified. It has been shown that S82 (for PPAR γ 1) /S112 (PPAR γ 2) phosphorylation, by ERK and JNK pathways result in PPAR γ inhibition [8-11]. If this serine is substituted by alanine (S82A in mice, or S84A in human), MAPK mediated PPAR γ inhibition is lost [8,12]. Studies introducing a serine to aspartate (S112D) mutation suggest that the mechanism by which the N-terminus modulates ligand binding is caused by conformational changes of the unligated receptor and that the S112 phosphorylation status influences its conformation thereby decreasing its affinity for the ligand [9]. Serine (S82/S112) phosphorylation affects not only coactivators and co-repressors recruitment but also ubiquitination, proteasomal degradation and sumoylation [13]. For example PPAR γ activity is decreased *via* the ubiquitination degradation pathway [14]. Alternatively, PPAR γ sumoylation, which is enhanced by S112 phosphorylation, promotes the co-repressors recruitment and the repression of inflammatory or adipocyte differentiation genes [6,15]. In addition, SUMO-1 also affects PPAR γ stability but not the nuclear localization of PPAR γ [16]. One S82/S112 independent mechanism that affects the genomic actions of PPAR γ is its translocation to the cytoplasm by the AF-2/PPAR γ /MEKs-interaction after a mitogenic stimulus or PPAR γ ligand administration [17]. On the other hand the PPAR γ translocation to the nucleus induced by the ligand binding is blocked upon nitration of tyrosine residues [18].

Like other NHR, PPARs also inhibit proinflammatory gene expression by a controversial mechanism of *transcriptional transrepression*, which is not mediated by their binding to PPREs. PPAR γ is able to suppress expression of proinflammatory genes in myeloid lineage cells, such as microglia and macrophages, and in the vasculature, by suppressing the action of other transcription factors like NF κ B, AP-1 and STAT1 [19]. One mechanistic model, the corepressor-dependent transrepression, has recently been proposed: under basal conditions NF κ B-regulated inflammatory genes are maintained in a repressed state by N-Cor containing corepressor complexes. Upon exposure to proinflammatory stimuli this N-Cor containing complex is dismissed and gene expression is initiated. This dismissal can be prevented by sumoylated PPAR γ :PPAR γ agonist complexes that stabilizes NCor complexes at the promoters of NF κ B-regulated genes, thus preventing inflammatory gene expression [20,21].

Binding of PPARs to their specific ligands leads to conformational changes which allow co-repressor release and co-activator recruitment. Even though all PPARs can be attributed to a common ancestral nuclear receptor, each PPAR isotype has its own properties with regard to ligand binding. Synthetic thiazolidinediones (TZDs), which are commonly prescribed for the treatment of type II diabetes, are selective PPAR γ ligands. Naturally occurring PPAR γ ligands include eicosanoids and the prostaglandin 15d-PGJ2. The best characterized PPAR γ agonists are pioglitazone and rosiglitazone which are Food and Drug Administration (FDA) approved for treatment of type II diabetes and troglitazone, which has been withdrawn in 2000. PPAR α agonistic ligands include fibrates that are commonly used for the treatment of hypertriglyceridemia and WY14,643 and GW7647. PPAR β/δ agonists include the prostacyclin PGI2, and synthetic com-

pounds GW0742, GW501516, and GW7842. In addition, all PPARs can be activated by polyunsaturated fatty acids with different affinities [22]. An overview addressing the affinity of several natural and synthetic ligands has recently been summarized [23].

PPARs During Development

PPAR α and γ transcripts appear late during fetal development of rat and mouse (day 13.5 of gestation), with similar expression pattern to their adult distribution. PPAR α is found in the liver, the kidney, the intestine, the heart, the skeletal muscle, the adrenal gland and the pancreas. PPAR γ expression is restricted to the brown adipose tissue (day 18.5 of gestation), and to the CNS (day 13.5 to 15.5 of gestation). Compared to the two other isotypes, PPAR β/δ is expressed ubiquitously and earlier during fetal development [24]. In adult rodent organs, the distribution of PPAR α is similar to its fetal pattern of expression.

Not much is known about the expression of the PPARs during human development [25-27]. PPAR α is most highly expressed in tissues that catabolise fatty acids, such as the adult liver, heart, kidney, large intestine and skeletal muscle. PPAR β/δ mRNA is present ubiquitously, with a higher expression in the digestive tract and the placenta. PPAR γ is abundantly expressed in the white adipose tissue, and is present at lower levels in the skeletal muscle, the heart and the liver. Surprisingly, and in contrast to rodents, human PPAR γ seems to be absent from lymphoid tissues, even though PPAR γ has been shown to be present in macrophages in human atheroma.

PPARs in the Brain

All three PPAR isotypes are co-expressed in the nervous system during late rat embryogenesis, and PPAR β/δ is the prevalent isotype. The expression of the three PPAR isotypes peaks in the rat CNS between day 13.5. and 18.5 of gestation. Whereas PPAR β/δ remains highly expressed in this tissue, the expression of PPAR α and γ decreases postnatally in the brain [28]. While PPAR β/δ has been found in neurons of several brain areas, PPAR α and γ have been localized to more restricted brain regions [29,30]. Analysis of the expression of PPARs in different brain regions of adult mice revealed that PPAR β/δ mRNAs are preferentially found in the cerebellum, the brain stem and the cortex, whereas PPAR γ mRNAs are enriched in the olfactory bulb as well as in the cortex. Expression of all three isotypes was found to be low to moderate in the hippocampus. More detailed analysis of PPARs expression within the hippocampus by *in situ* hybridisation revealed an ubiquitous expression pattern for PPAR α , whereas PPAR β/δ was found to be enriched in the dentate gyrus/CA1 region and PPAR γ expression was restricted to the CA3 region [31].

Even though this pattern of expression, which is isotype specific and regulated during development, suggests that the PPARs may play a role during the formation of the CNS, their function in this tissue are still poorly understood. Both *in vitro* and *in vivo* observations show that PPAR β/δ is the prevalent isoform in the brain, and is found in all cell types, whereas PPAR α is expressed at very low levels predominantly in astrocytes [32]. Acyl-CoA synthetase 2, which is

crucial in fatty acid utilization, is regulated by PPAR β/δ at the transcriptional level, providing a facile measure of PPAR β/δ action. This observation strongly suggests that PPAR β/δ participates in the regulation of lipid metabolism in the brain. This hypothesis is further supported by the observation that PPAR β/δ null mice exhibit an altered myelination of the corpus callosum. Such a defect was not observed in other regions of the central nervous system, and the expression of mRNA encoding proteins involved in the myelination process remained unchanged in the brain.

Expression of all PPAR isoforms, including PPAR γ , has been confirmed in the adult brain. Furthermore, it has been suggested that PPAR activation in neurons may directly influence neuron cell viability and differentiation [33-37]. Of note, selective knockdown of PPAR γ renders neurons more vulnerable to oxygen-glucose deprivation *in vitro* as well as to ischemic brain damage *in vivo* [38]. Furthermore, neuronal PPAR γ seems to have, at least *in vitro*, an important function for neurite outgrowth [39].

The localization of PPARs has also been investigated in purified cultures of neural cells. PPAR β/δ is expressed in immature oligodendrocytes and its activation promotes differentiation, myelin maturation and turnover [40,41]. The PPAR γ is the dominant isoform in microglia. Astrocytes possess all three PPAR isotypes, although to different degrees depending on the brain area and animal age [42,43]. The role of PPARs in the CNS is mainly been related to lipid metabolism, however, these receptors, especially PPAR γ , have been implicated in neural cell differentiation and death as well as in inflammation and neurodegeneration [34]. PPAR α has been suggested to be involved in the acetylcholine metabolism [44] and to be related to excitatory amino acid neurotransmission and oxidative stress defence [29].

Inflammation and Alzheimer's Disease

The number of individuals with the Alzheimer's disease (AD) is dramatically increasing as a consequence of a longer life expectancy in our societies. The large number of affected individuals and the increasing prevalence of the AD presents a substantial challenge to health care systems and does so in the face of substantial economic costs. The pathological hallmarks of AD are the formation of extracellular plaques consisting of amyloid- β peptides and intracellular neurofibrillary tangles made up from hyperphosphorylated tau protein, causing neuronal death that is responsible for progressive memory loss and inexorable decline of cognitive functions [45,46]. Analysis of the genetic forms and animal models suggested a pivotal role for the amyloid β peptide (A β), nevertheless, the biological basis of AD, especially of the sporadic forms, is still poorly understood. Genetically, A β metabolism is closely linked to lipid metabolism as a certain allele of the lipid carrier protein ApoE is associated with significantly increased risk for AD [47]. Another key hallmark of AD brain is the presence of chronic neuroinflammation without any signs of leukocyte infiltration. Amyloid plaques within the brain are populated by abundant, activated microglia and astrocytes [48]. Microglial activation is accompanied by the secretion of inflammatory cytokines and chemokines including interleukin (IL)-1 β , IL-6, monocyte chemoattractant protein-1, (MCP-1) and tumor necrosis factor

(TNF)- α [49]. It was postulated that activation of microglia and the concurrent production of inflammatory molecules may deteriorate and accelerate the progression of AD and therefore directly contribute to neuronal loss [48,50]. Next to microglia, activation of astrocytes and glial derived inflammatory molecules may as well as neuronal expression of inflammatory enzyme systems, including iNOS, in significantly contribute the inflammatory component of AD [51-53]. Increasing evidence suggests that anti-inflammatory therapies may be beneficial for AD treatment see Fig. (1).

PPAR γ in Experimental Models of Alzheimer's Disease

PPAR γ is expressed in the brain at low levels under physiological conditions. Recently, a detailed gene expression analysis has demonstrated that mRNA levels are elevated in AD patients [54]. This suggests that PPAR γ could play a role in the modulation of the pathophysiology of AD. Currently used drugs are mainly targeted at symptomatic improvement of the patients. These agents have only modest therapeutic efficacy over rather short periods of time. Thus, the development of new therapeutic approaches is of critical importance.

The initial studies exploring the actions of PPAR γ in AD were based on the ability of non-steroidal anti-inflammatory drugs (NSAID) to activate this receptor. A number of epidemiological studies demonstrated that NSAID treatment reduces AD risk by as much as 80% and it was suggested that these effects arise from the ability of these drugs to stimulate PPAR γ and to inhibit inflammatory responses in the AD brain [55-59]. This hypothesis is supported by the finding that experimental expression of iNOS in neurons resulted in time dependent neuronal cell death which was prevented by activation of PPAR γ *in vitro* and *in vivo* [34,60]. In addition, PPAR γ activation in microglial cells suppressed inflammatory cytokine expression, iNOS expression and NO production as well as inhibited COX2 and therefore the generation of prostanoids [61]. These latter effects result from the ability of PPAR γ to suppress proinflammatory genes through antagonism of the transcription factor NF κ B, (and to a lesser extent, AP-1 and STATs) [19]. PPAR γ agonists have also been demonstrated to suppress the A β -mediated activation of microglia *in vitro* and prevented cortical or hippocampal neuronal cell death [61-63]. In a rat model of cortical A β injection, coinjection of ciglitazone and ibuprofen or oral pioglitazone administration potently suppressed A β -evoked microglial cytokine generation [64]. The effects of the PPAR γ agonists pioglitazone and ibuprofen have been investigated in animal models of AD (Tg2576) that overexpress human APP. Pioglitazone was selected as it passes the blood brain barrier, although with limited penetration [65]. 12 months old Tg2576 mice were treated orally for 4 months resulting in a significant reduction of SDS-soluble A β 40. A β 42 levels were only significantly lowered for ibuprofen treated animals, but a trend was observed for pioglitazone, too [66].

The modest effects of pioglitazone in this study were thought to be due to poor drug penetration into the brain. In a subsequent study treatment with larger doses of pioglitazone in aged APPV717I transgenic mice significantly decreased microglial and astroglial activation as well as A β plaque

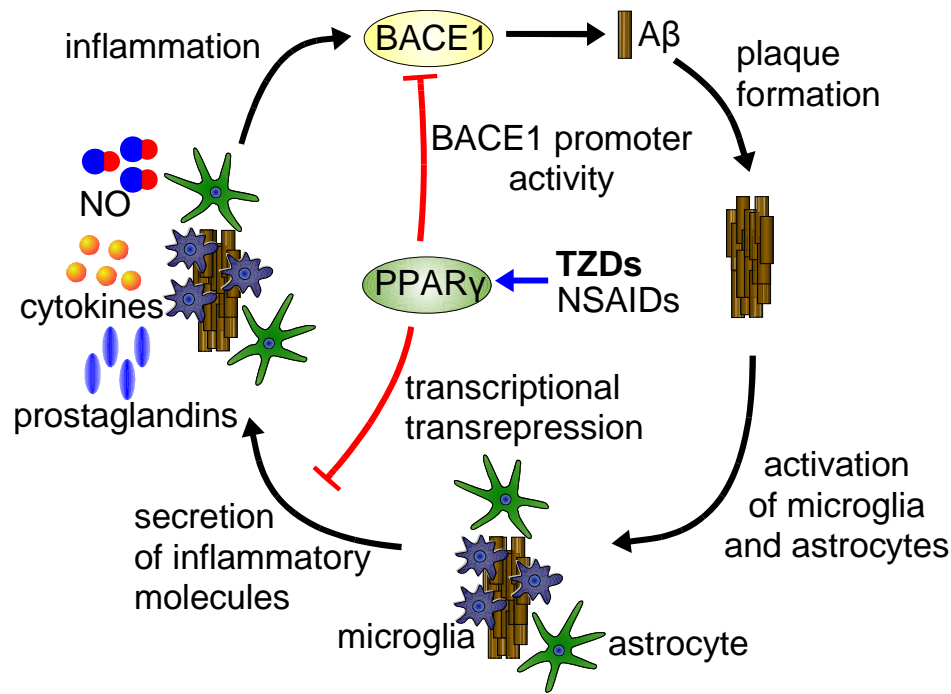


Fig. (1). Effects of PPAR γ on A β metabolism. Excessive production or insufficient clearance of A β results in its aggregation and finally in the formation of amyloid plaques. This process induces the activation of microglia as well as astrocytes which respond with the secretion of proinflammatory molecules like NO, cytokines and prostaglandins developing the inflammatory phenotype of AD. In addition, cytokines are able to increase BACE1 activity thereby stimulating A β production. PPAR γ agonists are able to abate both effects by either transrepress the production of proinflammatory molecules or directly interfere with the binding of PPAR γ to a PPRE in the BACE1 gene promoter.

burden [67]. The finding that PPAR γ agonists elicited a reduction in amyloid pathology may be the result of the ability of PPAR γ to affect A β homeostasis. According to this hypothesis, evidence has been provided that immunostimulated beta secretase 1 (BACE1) expression is silenced by a PPAR γ -dependent regulation of the BACE 1 gene promoter [68,69]. Similarly, oral pioglitazone treatment of APP transgenic mice reduced BACE1 transcription and expression. A recent study has found that PPAR γ is associated with enhanced A β clearance. PPAR γ activation, in both glia and neurons, led to a rapid and robust uptake and clearance of A β from the medium [70]. It has also been suggested that NSAIDs act directly on A β processing by the γ -secretase complex resulting in selective decrease of A β 1-42 production [71,72], even so this hypothesis has recently been challenged [73,74].

Additionally, modulation of the Wnt/ β -catenin signalling pathway may also account for some PPAR γ mediated beneficial effects in AD since recent findings show that PPAR γ mediated protection of hippocampal neurons against A β -induced toxicity directly correlates with β -catenin levels, inhibition of GSK 3 β activity and increased levels of Wnt-target genes [35,75]. Furthermore, recent evidence suggests that PPAR γ activation may also provide protection from excitotoxic stimuli [76] and positively influences neural stem cell proliferation and differentiation [77], both mechanisms that could potentially influence the overall salutary effects observed in models of neurodegenerative disease.

In a further animal study, Pedersen and colleagues have demonstrated that rosiglitazone treatment of Tg2576 mice resulted in behavioural improvement in these animals as well

as in reduction of A β 1-42 in the brain. Treatment with rosiglitazone for 7 months enhanced spatial working and reference memory [78]. Significantly, drug treatment was associated with a 25 % reduction in A β 1-42 levels, however A β 1-40 levels remained unchanged. Similar results were obtained in a recently published study in 10 month of J20 mice, treated with rosiglitazone for 4 weeks [79]. This reduction of A β 1-42 was argued to arise from increased levels of insulin degrading enzyme (IDE) in rosiglitazone treated transgenic mice. In line with this, it has been suggested that IDE is positively regulated by PPAR γ in primary neurons [80]. IDE is a A β degrading metalloprotease, that has been genetically linked to AD [81]. Similarly, chronic treatment of hAPP mice with rosiglitazone reverted memory decline and hippocampal glucocorticoid receptor down-regulation [81]. In addition, prevention of cognitive decline in an intracerebroventricular infusion model of A β 1-40 by telmisartan, a partial PPAR γ agonist, was abolished when mice were treated with the PPAR γ antagonist GW9662, further supporting a role of PPAR γ for neuroprotection [82]. Interestingly, infusion of the same drug into the fourth ventricle of APPS1 transgenic mice increased A β levels and gliosis within the cerebellum. Consequently, these mice did show a reduction of IDE expression and impaired motor function [83].

PPAR γ and Alzheimer's Disease

The influence of genetic mutations on the course or overall risk of Alzheimer's disease have almost only be addressed for the PPAR γ 2^{Pro12Ala} polymorphism, albeit a change in PPAR γ activity by this mutation will most likely affect the adipose tissue. However, a recent study revealed a

significant overrepresentation of the Ala12 allele in octogenarian AD patients [84], suggesting that carrying this polymorphism increased the AD risk in this population by nearly twofold. In contrast to the above, another study showed that the Ala12 polymorphism protected from AD in females but not in males [85]. Two further studies, investigating a German and a Finnish population failed to detect any significant association between the Ala 12 variant and the genetic risk of AD [86,87]. However, the study by Koivisto and colleagues, who analyzed the Pro12Ala as well as the C478T polymorphisms suggests that the carriers of both alleles have a lower age of onset compared to Pro12Pro/478CC carriers [86]. Importantly, this effect was independent of the ApoE4 status and various other factors. This finding has been partly reproduced in a recent study of a Chinese population, that found that in a subgroup of ApoE4 non-carriers, the Pro12Ala polymorphism was associated with an earlier disease onset [88]. In diabetics, however, Ala12 allele carriers show an increased risk of dementia or cognitive impairment in general when compared to non-carriers [89,90]. Exceeding these previous studies and looking at further single nucleotide polymorphisms (SNPs) in the PPAR γ gene, Helisalmi and colleagues failed to find any association between AD and their study groups in a Finnish population [91]. Therefore, a strong influence of PPAR γ gene polymorphisms on AD risk seems to be rather unlikely. Conducting a more detailed SNP-analysis may settle this contradiction. However, it may be important to gain deeper mechanistic understanding of the Pro12Ala mutations in peripheral tissues, thereby potentially revealing further insight on the interplay of obesity, insulin sensitivity and cholesterol metabolism in the context of AD.

Clinical investigations of insulin-sensitizing TZDs that are in clinical use for type II diabetes are currently ongoing. A small study of 30 patients with mild AD or MCI found that 6 months of treatment with rosiglitazone resulted in improved memory and selective attention. A larger trial of rosiglitazone in AD patients has recently been reported [92]. More than 500 patients with mild to moderate AD were treated for 6 months with rosiglitazone, resulting in a statistically significant improvement in cognition in those patients that did not possess an ApoE4 allele [93]. Patients with ApoE4 did not respond to the drug and showed no improvement in standard cognitive tests. As an explanation it was suggested that rosiglitazone acts on mitochondria in the brain, increasing their metabolic efficiency and number. This hypothesis is supported by the observation that rosiglitazone induces neuronal mitochondrial DNA expression, enhances glucose utilization by inducing transcription of glucose metabolism and mitochondrial biogenesis genes leading to improved cellular function in mice. Noteworthy, these effects were also observed in animals expressing the ApoE4 allele. Determination of the amount of rosiglitazone in the brain revealed that 9-14 % of the blood rosiglitazone crossed the blood brain barrier after oral treatment [94]. The actions of TZDs on mitochondria occur through both PPAR γ dependent and independent mechanisms [95]. The basis of the differential effects of rosiglitazone in individuals depending on their ApoE genotype is unexplained. The outcome of this clinical trial is, however, consistent with previous findings with respect to the influence of the ApoE4 genotype [96-98]. A

recently published single center clinical trial using pioglitazone for the first time in type II diabetic AD patients showed a significant improvement concerning neuropsychological tests, regional cerebral blood flow as well as plasma A β levels in response to pioglitazone treatment. In strong contrast, most of these parameters worsened in the control population without pioglitazone treatment [99]. While this study is limited by its small number of recruited patients and an open- but not placebo controlled trial design, it strongly calls for a more elaborated study.

CONCLUSION

PPARs exhibit a wide range of activities to positively influence the pathology of Alzheimer's disease. Beside the ameliorating effect of PPAR γ agonists on the inflammatory status of the AD brain by repressing the secretion of proinflammatory molecules and the enhancement of mitochondrial function, a direct involvement in the processing of the A β peptide has been demonstrated Fig. (1). The compelling results from animal models of Alzheimer's disease underline the beneficial effects of PPAR γ agonists for future therapies. The importance of these activities for the disease altering actions of PPAR γ agonist as well as the underlying molecular mechanisms have to be elucidated in future research.

ABBREVIATIONS

A β	=	Amyloid β
AD	=	Alzheimer's disease
BACE	=	β -secretase
IDE	=	Insulin degrading enzyme
IL	=	Interleukin
MCI	=	Mild cognitive impairment
NF κ B	=	Nuclear factor κ B
NSAID	=	Non steroidal anti-inflammatory drug
PPAR	=	Peroxisome proliferator activated receptor
SNP	=	Single nucleotide polymorphism
TNF α	=	Tumor necrosis factor- α
TZD	=	Thiazolidinedione

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