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Molecular Mechanisms Regulating Vascular Tone by PPARγ

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Abstract

Purpose of review—This review summarizes recent findings on the regulation of vascular tone by the nuclear receptor transcription factor, peroxisome proliferator-activated receptor (PPAR) γ . Much of the recent work utilize genetic tools to interrogate the significance of PPARγ in endothelial and smooth muscle cells and novel $PPAR_{\gamma}$ target genes have been identified.

Recent findings—Endothelial PPARγ prevents inflammation and oxidative stress, while promoting vasodilation by controlling the regulation of NADPH oxidase, catalase and superoxide dismutase gene expression. Moreover, the protective functions of endothelial PPARγ appear more prominent during disease conditions. Novel findings also suggest a role for endothelial PPARγ as a mediator of whole body metabolism. In smooth muscle cells, $PPAR_Y$ regulates vascular tone by targeting genes involved with contraction and relaxation signaling cascades, some of which via transcriptional activation, and some through novel mechanisms regulating protein turnover. Furthermore, aberrant changes in renin-angiotensin system components and exacerbated responses to angiotensin II-induced vascular dysfunction are observed when PPARγ function is lost in smooth muscle cells.

Summary—With these recent advances based partially on lessons from patients with PPARγ mutants, we conclude that vascular $PPAR\gamma$ is protective and plays an important role in regulation of vascular tone.

Keywords

PPARγ; endothelial cells; smooth muscle cells; vasculature; vessel tone

Introduction

The purpose of this review is to highlight mechanisms by which peroxisome proliferatoractivated receptor (PPAR) γ regulates vascular tone. Vascular tone is a key determinant of systemic vascular resistance and local tissue perfusion. It is dependent on the function of the endothelium, smooth muscle cells and to some extent on the perivascular adipose tissue (PVAT). As vascular cell type-specific functions of $PPAR\gamma$ have just started to emerge, the authors will discuss key findings from previous studies that provide insights into the mechanisms of PPAR_Y action. We will focus the review on the function of PPAR_Y in

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endothelial and smooth muscle cells, particularly mechanisms uncovered by the use of genetically engineered mice as models to overcome the limitations of systemic thiazolidinedione (TZD) administration. Overall, numerous reports consistently support the concept that vascular PPARγ is protective.

PPARγ **and Mechanism of Actions**

PPAR_{γ} is a nuclear hormone receptor transcription factor. Two human PPAR γ isoforms have been identified, PPAR γ 1 and PPAR γ 2, which differ by a 30-amino acid extension at the N terminus. PPAR γ 2 (the longer form) is highly expressed in adipose tissue, where its function is a prerequisite for adipogenesis [1]. PPAR γ 1 is ubiquitously expressed at a low level in other tissues where it exerts cell-specific functions. PPARγ in macrophages regulates immune defense through the regulation of a distinct set of genes, with little overlap with PPARγ-target genes in adipocytes [2].

PPAR_Y regulates gene expression by forming a heterodimeric complex with the retinoid X receptor (RXR) at the PPAR response elements (PPRE). Without ligand, the transcription of PPAR_{γ} target genes is repressed due to the interaction of PPAR_{γ} with co-repressor complexes. Upon ligand binding, PPARγ undergoes a conformational change in the ligand binding domain and promotes gene expression through recruitment of a co-activator complex [1, 3].

Clues to PPARγ**-Regulated Blood Pressure and Vascular Function**

The evidence that implicates $PPAR_{\gamma}$ in the regulation of cardiovascular homeostasis comes primarily from the observations that: 1) TZD, potent PPARγ agonists, lower blood pressure and alleviate cardiovascular diseases [4] and 2) patients with some PPARγ mutations develop severe early-onset hypertension [5]. However, the molecular mechanism and the site of action for PPARγ-mediated blood pressure regulation requires further investigation.

TZDs are high affinity synthetic ligands of $PPAR_Y$, previously used to treat patients with type 2 diabetes. The effectiveness of the TZD class in regulating glycemic control occurs through multiple mechanisms including augmented β-cell function and enhanced insulin sensitivity in skeletal muscle, adipocyte and liver [6**]. Induction of adiponectin, fibroblast growth factor family (FGF) 1 and FGF21 by TZD were reported to be important mediators for the insulin sensitizing effect of a TZD [3, 7-9]. In combination, the activation of these mechanisms by TZD provides a long lasting glycemic control compared to sulfonylureas and metformin [6*]. However, reports of serious adverse events including weight gain, congestive heart failure, fluid retention and osteoporosis have limited the clinical usefulness of TZD. The controversy was sparked in 2007 by a meta-analysis which reported the association of rosiglitazone with increased risk of myocardial infarction and cardiovascular deaths [10]. Despite the limitations of meta-analysis, this study raised serious public concerns and the prescription of rosiglitazone was subsequently restricted. As of last year, the FDA recommended that this restriction be removed after an independent re-adjudication showed no cardiovascular harms associated with the use of rosiglitazone [11**]. Although the results were not significantly different, fewer deaths from a cardiovascular cause were observed in patients received rosiglitazone compared to those treated with metformin and

sulfonylurea. Another member of the TZD class, pioglitazone has been shown to reduce cardiovascular outcomes in diabetic patients [11**]. It has been speculated that the improvement of lipid profile by pioglitazone might account for these differences [6**]. However, it is important to note that safety concerns reporting increased risk of bladder cancer with pioglitazone have also been raised [12**].

Interestingly, multiple studies in human and animal models have shown that TZDs cause a modest decrease in both diastolic and systolic blood pressure, an effect that is not observed with other insulin sensitizer drugs [4]. It is interesting to note that TZD-mediated blood pressure lowering occurs despite edema and plasma volume expansion, implicating robust non-renal actions of PPAR γ to lower blood pressure. Consistent with this is the observation that patients with some PPAR γ mutations exhibit severe early onset hypertension [5, 13^{**}, 14, 15]. This suggests the plausible hypothesis that there is a direct non-renal action of PPARγ, perhaps in the vasculature. That the same mutations also cause lipodystrophy, dyslipidemia, insulin resistance and type 2 diabetes argues either that the effects of the mutations are pleiotropic, that is, they target multiple tissues, or the hypertension might be a secondary consequence from metabolic changes. A knock-in mouse model carrying the mouse equivalent to one of the PPARγ mutations (P465L) originally identified in patients, exhibited hypertension and cerebral arterial dysfunction despite normal insulin sensitivity, thus arguing against the latter [16]. This is further supported by studies where PPARγ activity was manipulated by deletion or over-expression of PPARγ mutants in a vascular cell type-specific manner which resulted in blood vessel dysfunction, independent of metabolic abnormalities [17-21**]. Moreover, transfection of vascular cells *in vitro* with two different mutants in PPARγ (R165T and L339X), which were reported to cause severe hypertension in patients, recapitulated a robust induction of the renin-angiotensin system (RAS) and increased inflammation, and phenocopied what occurred in cells isolated from patients [13^{**}]. Taken together, these studies reinforce the concept of the direct actions of PPAR_{γ} in vascular cells.

Roles of PPARγ **in Endothelium**

Many studies have shown that endothelial PPARγ has anti-inflammatory and anti-oxidant actions, while promoting vasodilatation (Figure 1). TZD treatment of cultured vascular endothelial cells increases nitric oxide (NO) production via post-translational modification of eNOS [22] and preserves NO bioavailability through suppressing NADPH oxidase expression [23]. Activation of PPAR γ in the vasculature counteracts endothelin-1-induced constriction by induction of endothelin receptor type B expression in endothelial layers [24]. It was reported that endothelial PPAR γ is required for the blood pressure lowering effect of TZD [25, 26]. Endothelial-specific disruption of a conditional allele of PPAR γ (PPAR γ ^{flox}) with Tie2-promoter driven Cre-recombinase (Tie2Cre) resulted in mild hypertension and endothelial dysfunction that was associated with reduced NO production, increased reactive oxygen species, and enhanced NFκB activity [27]. In contrast, another study with Tie2Cremediated PPARγ disruption reported no change in systemic blood pressure at baseline but disrupted diurnal variations of blood pressure and heart rate [28]. Femoral arterial reactivity to phenylephrine, angiotensin II and KCI were significantly increased in Tie2Cre/PPAR $\gamma^{\rm{flow}}$ mice [26]. Other phenotypes outside the vasculature were reported in studies using Tie2Cre,

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most likely because the promoter is active in cells other than endothelium [29, 30]. For this reason, other investigators have utilized mice with vascular endothelial-cadherin (cdh5)-Cre recombinase-driven PPARγ deletion. These mice showed aggravated ischemia-induced blood-brain barrier disruption due to cerebrovascular permeability, with no change in systemic blood pressure [31**]. Mechanistically, Kruppel-like factor (KLF)-11 has been identified as a PPAR γ gene target and was reported to function as a PPAR γ co-regulator in cerebral vascular endothelial cells. Pioglitazone-mediated vascular protection following middle cerebral artery occlusion was significantly lost in KLF11 null mice [31**].

Emerging evidence highlights the prominent role of endothelial PPAR γ during stress conditions. Transgenic mice expressing a dominant negative PPARγ mutant (V290M or P467L) specifically in endothelium exhibited reduced vasodilation to acetylcholine in basilar artery and aorta after prolonged high fat diet treatment [20] or during dyslipidemia induced by disruption of Apolipoprotein E, but not at baseline [32**]. This impairment was restored by superoxide scavenger, suggesting increased oxidative stress caused by the loss of PPARγ function. A significant increase in transcription of pro-oxidant genes such as p22phox, Noxo2, and NoxA2, concomitant with a reduction of mRNA of anti-oxidant catalase and Cu/Zn SOD was observed in vascular endothelium from these mice [20]. This is consistent with catalase and Cu/Zn SOD being PPARγ target genes [33, 34]. It is entirely possible that endogenous PPARγ ligands, perhaps derived from free fatty acids, induce upregulation of these genes during high fat feeding in normal mice, and this provides a protective mechanism. The downregulation of catalase and Cu/Zn SOD transcripts in the transgenic mice is potentially a direct consequence of the failure of endogenous PPARγ ligands to increase the transcriptional activity of the V290M and P467L mutants of PPARγ which are located in the ligand binding domain [15].

Endothelial dysfunction is often superimposed with obesity and type 2 diabetes. A novel role for endothelial cells in energy storage through fatty acid release in and out of the peripheral tissues has started to emerge [35]. Indeed, classic PPARγ transcriptional targets involved with fatty acid metabolism such as CD36, aP2 and RBP7 were shown to be expressed in endothelial cells [18]. Activation of PPARγ in microvascular endothelial cells markedly increased expression of aP2 and CD36 [36*]. Down-regulation of these fatty acid transporters mRNA in endothelium from endothelial-specific PPARγ knockout mice was associated with dyslipidemia after high fat overload [18,36*]. Unexpectedly, these mice manifested reduced white adipose tissue mass and improved insulin sensitivity following high fat diet, effects which persisted even after reconstitution of hematopoietic cells in the endothelial-specific PPAR_{γ} knockout, suggesting that endothelial PPAR_{γ} is critical for metabolic balance and lipid accumulation in periphery [18]. Despite improving insulin sensitivity, impaired vasorelaxation and hypertension following high fat diet was observed [18]. These findings suggest that endothelial PPARγ might regulate vascular function by targeting a set of genes distinct from those responsible for metabolism. This study also enforces the hypothesis that endothelial PPAR γ is required for vascular protection during fatty acid overload.

Roles of PPARγ **in Smooth Muscle**

The net signaling in smooth muscle initiated by neurohumoral factors as well as intrinsic properties of the vessels determines the tone of the vasculature (Figure 2). Specific expression of dominant negative mutant PPARγ (P467L) in smooth muscle cells of transgenic mice (termed S-P467L) resulted in hypertension, vascular dysfunction, cerebral arteriole remodeling, tachycardia and baroreflex dysfunction [19, 37**]. Aberrant vascular reactivity is observed in both conduit and resistance arteries, emphasizing the critical role of PPARγ in regulation of the blood vessel tone. In aorta and cerebral arteries, interference of PPAR_Y contributed to loss of responsiveness to NO-dependent vasorelaxation [38, 39^{**}]. Strikingly, the hypercontractile phenotype in response to receptor-dependent agonists is apparent in these transgenic mice. Inhibition of Rho kinase not only blunted the augmented constriction, but it also restored the impaired NO responsiveness in aorta and cerebral arteries from these transgenic mice [38, 39**], which is consistent with aberrantly increased Rho kinase activity. Mechanistically, mutant PPARγ led to a defect in RhoA degradation caused by a reduced expression of Cullin-3, a component of an E3-ubiquitin ligase complexmediating proteasomal degradation of different substrates including RhoA. In smooth muscle cells, down-regulated Cullin-3 expression significantly increased RhoA [38]. Preincubation of isolated aorta from a normal mouse with a pan-Cullin inhibitor resulted in augmented vasoconstriction that was dependent on Rho kinase, resembling the phenotype observed in S-P467L aorta. RhoBTB1, a potential Cullin-3 adapter was identified as a transcriptional target of PPARγ and its expression was dramatically decreased by P467L mutant PPAR γ [38]. It remains unclear if loss of this specific adapter contributes to suppression of Cullin-3 activity and reduced RhoA degradation.

The role of PPAR γ in regulation of myogenic tone has also been reported. Myogenic tone is an intrinsic property of smooth muscle in resistance arterial beds. This mechanism allows precise control of local blood flow despite the fluctuation of systemic blood pressure. Inhibition of PPARγ function mediated by dominant negative expression in smooth muscle (S-P467L) resulted in a marked increase in myogenic constriction in small mesenteric arteries. Augmented basal tone is attributable to a robust downregulation of the regulator of G protein signaling 5 (RGS5) mRNA and subsequent increase in protein kinase C activity. Loss of RGS5 transcript in S-P467L mesenteric arteries is also associated with a selective increase in angiotensin II-induced vasoconstriction and ERK1/2 activation [40, 41*], consistent with findings from RGS5 null mice [42*]. Several lines of investigation reveal that RGS5 is a novel PPAR_Y target. For example, PPAR_Y bound to a PPRE in the RGS5 locus and TZD increased RGS5 transcript, an effect that was completely lost in resistance vessels with dominant negative PPARγ expression [40].

Down-regulation of PPARγ expression or activity has often been correlated with increased AT_1 receptor (AT_1R) expression, and a recent study implicated hypoxia-inducible factor (HIF)-1α [43*]. Loss of HIF1α in smooth muscle resulted in suppressed PPARγ expression and subsequent upregulation of AT_1R ; and smooth muscle-specific HIF1 α knockout mice exhibited augmented vasoconstriction to angiotensin-II and increased systolic blood pressure. Recent clinical evidence also highlights the interaction between PPARγ and the renin-angiotensin system (RAS). Patients with mutations in the DNA binding domain

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(R165T) or ligand binding domain (L339X, causing a truncation) of PPARγ exhibited lipodystrophy, insulin resistance and severe hypertension. It is surprising to note that in addition to AT_1R , other RAS components including renin and angiotensinogen were markedly increased in fibroblasts or blood mononuclear cells isolated from these patients [13**]. The authors reported that the effect of the mutations was reproduced in transfected vascular smooth muscle cells, which lead them to conclude the induction of the RAS was due to a direct effect of loss of PPAR γ function. AT_1R antagonist reportedly improved blood pressure control in these patients [13**], but blood vessel function was not studied. It is notable that AT_1R has been recently shown to act as a mechanosensor $[44^*]$. Although the mechanism whereby reduced PPAR γ activity leads to increased AT₁R remains unclear, it is possible that increased AT_1R in these patients may have contributed to increased myogenic tone, vascular dysfunction, and oxidative stress.

Mice with lifelong PPAR γ deficiency specifically in smooth muscle (SM22Cre/PPAR γ^{flox}) exhibited slightly increased systemic blood pressure and impaired circadian variations of mean arterial pressure [28]. Another study with inducible smooth muscle specific PPAR γ inactivation in adult mice using smooth muscle myosin heavy chain promoter reported no difference in baseline blood pressure before or after angiotensin-II infusion compared to control mice [21**]. Interestingly, the mesenteric arteries from these mice exhibited exacerbated angiotensin-II-induced endothelial dysfunction, vascular remodeling and inflammation [21**]. These findings are consistent with the hypothesis that smooth muscle PPAR γ is protective against hypertension and vascular dysfunction. In contrast to these studies, another line of smooth muscle PPARγ deficient mice generated by crossing $PPAR\gamma^{flow}$ mice with knock-in mice expressing Cre-recombinase in the endogenous SM22 gene locus manifested decreased blood pressure and enhanced vasodilation to β-adrenergic receptor [17]. The surprising and inconsistent results were potentially reconciled when the same author later reported that these mice completely lack perivascular adipose tissue (PVAT) resulting from the transient activation of SM22 promoter in PVAT during development [45]. Whereas it is entirely possible that loss of PVAT might contribute to discrepant results compared with other studies, it also underscores the potential importance of PVAT in regulating vascular tone.

New Surprises

The progressive work on the function of $PPAR_Y$ in the vasculature consistently support the notion that PPAR γ is required to maintain homeostasis of the blood vessel (Figures 1 and 2). Emerging evidence has recently revealed that the complexity of $PPAR_Y$ extends far beyond the classic view as a regulator of gene transcription. For example, PPARγ was reported to act as E3 ligase that promotes ubiquitination of p65, a subunit of NFκB, through a direct PPARγ-p65 interaction to regulate inflammatory responses [46]. PPARγs activity as an E3 ligase for p65 would be functionally analogous, but mechanistically distinct from its better characterized trans-repression of inflammatory gene transcription [47]. Other studies demonstrated that the activity of PPARγ can be regulated by post-translational modifications. This has become a very provocative topic of discussion and investigation. With relevance to anti-inflammatory activities of PPARγ, ligand dependent-sumoylation of PPAR γ inhibits pro-inflammatory gene targets of NF κ B or AP1 through a trans-repression

mechanism [48]. The transcriptional activity of $PPAR_Y$ can also be modulated by phosphorylation and acetylation. Phosphorylation of PPARγ at Ser273 selectively decreases expression of a subset of genes involved with insulin sensitivity [49]. Deacetylation of PPAR γ at Lys268 and Lys293 is involved with browning of white adipose tissue, a phenomenon that promotes enhanced metabolism [50]. Thus, PPARγ activity may not only be impaired by mutation, but also under conditions which promote its post-translation modification. Accordingly, it is entirely possible that the mediators often correlated with vascular diseases and hypertension such as oxidative stress, angiotensin II and free fatty acids are able to modify and impair PPARγ activity via these mechanisms which further promotes vascular dysfunction.

Conclusion

Understanding the complexity of $PPAR_Y$ function, particularly in specific tissues or cell types will help develop new classes of drugs that can either target PPARγ or the physiological pathways it regulates more selectively. With new data suggesting that activation of PPAR_Y in the brain and bone $[12**]$ can contributes to weight gain and bone loss, respectively suggests the need for a new therapy that provides less accessibility to these tissues. Although it would be technically challenging to chemically activate PPARγ selectively in the vasculature it may be possible to selectively activate or inhibit some of the final common pathways regulated by PPARγ in endothelium or smooth muscle.

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Key points

- **•** PPARγ, a nuclear receptor plays an important role in the vasculature but the mechanisms remain under investigation.
- **•** Recent studies from genetically modified mouse models have emphasized the specific functions of PPARγ in endothelial and smooth muscle cells, where it provides protective mechanism against cardiovascular diseases.
- **•** Endothelium PPARγ modulates target genes expression involved with oxidative stress, inflammation and fatty acid transporters.
- **•** PPARγ regulates genes in signaling cascades related to constriction and relaxation in smooth muscle.
- **•** PPARγ may control the expression and activity of the renin-angiotensin in smooth muscle cells.

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In endothelial cells, PPARγ modulates target gene expression involved with oxidative stress, inflammation, cell survival and fatty acid transporters. Endothelial PPARγ promotes vasodilation through enhanced nitric oxide (NO) production, increased NO bioavailability and decreased reactive oxygen species (ROS). Activation of PPARγ in the vasculature counteracts endothelin-1-induced vasoconstriction by induction of ET_BR expression. PPAR_{γ} ameliorates inflammation in endothelial cells, perhaps through trans-repression of NFκB. It has been shown that KLF11 is necessary for pioglitazone-mediated vascular protection following middle cerebral artery occlusion. KLF11 is a direct target of PPAR γ and also functions as a PPARγ co-regulator in cerebral vascular endothelial cells. Other evidence also suggests that PPARγ in endothelial cells is important for metabolic balance and lipid accumulation in the periphery, possibly by regulating the fatty acid transporters, Fatty acid translocase (CD36), Fatty acid binding protein 4 (aP2) and retinol binding protein (RBP)-7. Retinoid X receptor (RXR); Cu/Zn superoxide dismutase (SOD); Endothelin receptor type B (ETBR); Nuclear factor kappa-light-chain-enhancer of activated B cells (NF-κB); Kruppellike factor (KLF)-11.

Figure 2. Role of PPARγ **in Vascular Smooth Muscle Cells**

PPARγ in smooth muscle cells regulates vascular tone via transcriptional activation, and some through novel mechanisms regulating protein turnover (RhoBTB1/Cullin 3). Downregulation of RhoBTB1/Cullin 3 in the blood vessel caused by smooth muscle PPARγ mutant is associated with increased RhoA/Rho kinase activity, leading to increased contraction. Smooth muscle PPAR γ is also important for normal myogenic tone via regulating Regulator of G-protein signaling (RGS)-5 transcript. Furthermore, exacerbated responses to angiotensin II either by up-regulated Angiotensin II receptor, type 1 (AT_1R) or decreased RGS5 are observed when PPARγ function is lost in smooth muscle cells and that potentially contributes to increased oxidative stress and vascular hypertrophy. In addition to AT_1R , other evidence from newly identified PPAR γ mutants indicates that PPAR γ is critical in regulating renin and angiotensinogen expression. Transcription of β2 adrenergic receptor has also been reported to be regulated by smooth muscle PPARγ, thereby promoting vasodilation. Retinoid X receptor (RXR); Rho-related BTB domain-containing protein (Rho BTB)-1; Ras homolog family member A (RhoA); Rho-associated protein kinase (ROCK); Inositol trisphosphate (IP3); Diacylglycerol (DAG); Phospholipase C (PLC); Protein kinase C (PKC); Mitogen-activated protein kinase (ERK1/2).