



REVIEW

Peroxisome proliferator-activated receptors in the cardiovascular system

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Peroxisome proliferator-activated receptor (PPAR)s are a family of three nuclear hormone receptors, PPAR α , - δ , and - γ , which are members of the steroid receptor superfamily. The first member of the family (PPAR α) was originally discovered as the mediator by which a number of xenobiotic drugs cause peroxisome proliferation in the liver. Defined functions for all these receptors, until recently, mainly concerned their ability to regulate energy balance, with PPAR α being involved in β -oxidation pathways, and PPAR γ in the differentiation of adipocytes. Little is known about the functions of PPAR δ , though it is the most ubiquitously expressed. Since their discovery, PPARs have been shown to be expressed in monocytes/macrophages, the heart, vascular smooth muscle cells, endothelial cells, and in atherosclerotic lesions. Furthermore, PPARs can be activated by a vast number of compounds including synthetic drugs, of the clofibrate, and anti-diabetic thiazolidinedione classes, polyunsaturated fatty acids, and a number of eicosanoids, including prostaglandins, lipoxygenase products, and oxidized low density lipoprotein. This review will aim to introduce the field of PPAR nuclear hormone receptors, and discuss the discovery and actions of PPARs in the cardiovascular system, as well as the source of potential ligands.

British Journal of Pharmacology (2000) **129**, 823–834

Keywords: PPAR; macrophage; endothelial cells; vascular smooth muscle; angiogenesis; atherosclerosis

Abbreviations: COX-1, constitutive COX; COX-2, inducible COX; 15d-PGJ₂, 15-deoxy- $\Delta^{12,14}$ -PGJ₂; ETYA, eicosatetraenoic acid; HETE, hydroxyeicosatetraenoic acid; HODE, hydroxyoctadecadienoic acid; HPETE, hydroperoxyeicosatetraenoic acid; LDL, low density lipoprotein; NF- κ B, nuclear factor- κ B; OxLDL, oxidized LDL; PAI-1, plasminogen activator inhibitor-1; PPAR, peroxisome proliferator-activated receptor; PPRE, PPAR response element; RXR, retinoid X receptor

Introduction

The vascular endothelial cell layer plays a critical role in the homeostasis of vascular tissue and blood components. By producing mediators such as prostacyclin, nitric oxide and plasminogen activator, which promote vasodilatation and an anti-thrombotic surface, ECs stop platelets and inflammatory cells adhering and aggregating (see Drexler & Hornig, 1999). Similarly, vascular smooth muscle cells perform important homeostatic functions, providing structural support, and controlling vascular resistance, which ultimately controls blood pressure. If the endothelium or vessels are damaged or become dysfunctional, an inflammatory process can follow. Within the vessel wall itself, cytokines and growth factors are released which activate and modify smooth muscle cells, leading to vascular remodelling. Cytokine-activated smooth muscle cells can become a more synthetic or 'secretory' cell type, producing a wide variety of mediators (see Ross, 1999). In large blood vessels secretory smooth muscle cells divide faster, and become migratory, leading to medial hyperplasia and intimal thickening. Furthermore, secretory smooth muscle cells can take up large quantities of oxidized lipoproteins. The inflammatory response leads to the recruitment of monocytes within the vessel wall, by a mechanism involving both the upregulation of chemokines, and adhesion molecule expression

(see Ross, 1999). Once there, monocytes become phagocytic macrophages, taking up large quantities of oxidized LDL to become lipid laden foam cells (see Ross, 1999; Berliner & Heinecker, 1996).

Peroxisome proliferator-activated receptors (PPARs) are a family of at least three nuclear receptors (α , δ , also referred to as PPAR β FAAR or NUC1 and γ ; Issemann & Green, 1990; Dreyer *et al.*, 1992) Until relatively recently their actions were thought limited to specific tissue types, having roles in lipid catabolism, and peroxisome proliferation in the liver (PPAR α ; Issemann & Green, 1990), and adipogenesis (PPAR γ ; see Spiegelman & Flier, 1996; see Spiegelman, 1998). Although PPAR δ is almost ubiquitously expressed (Kliwer *et al.*, 1994; Braissant *et al.*, 1996; Mukherjee *et al.*, 1997), due to the lack of any selective agonists or antagonists, its roles have yet to be ascertained. For this reason discussion will be mainly limited to PPAR α and PPAR γ subtypes. Over the last 2 years however it has become apparent that different PPARs are present in a variety of different cell types. The finding that spleen contained relatively high expression of PPAR γ mRNA (Kliwer *et al.*, 1994; Braissant *et al.*, 1996), lead to the discovery that monocytes, and especially elicited macrophages, contained PPAR γ (Ricote *et al.*, 1998a). Subsequently, different PPARs have been shown to be expressed both in vascular endothelial cells and smooth muscle cells in culture, and in the atherosclerotic lesions. This review aims to discuss this data in the context of what is known about PPARs, and discuss how the modulation of the PPAR pathway may lead to novel

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therapeutic targets for vascular and other inflammatory diseases.

PPAR ligands

PPARs can be activated by a number of ligands (see Table 1; Issemann & Green, 1990; Yu *et al.*, 1995; Brun *et al.*, 1996; Forman *et al.*, 1997; Kliewer *et al.*, 1997; Nagy *et al.*, 1998), including the fatty acids docosahexaenoic acid, linoleic acid, WY-14643 (selective for PPAR α), the anti-diabetic thiazolidinediones (troglitzone, rosiglitzone, and ciglitzone), a number of eicosanoids, including, 5,8,11,14-eicosatetraenoic acid (ETYA), LTB $_4$; the prostanoids PGA $_1$, PGA $_2$, (dehydration products of PGE $_1$ and PGE $_2$ respectively) PGI $_2$, and PGD $_2$. A number of non-steroidal anti-inflammatory drugs also can activate PPAR α and PPAR γ (Lehmann *et al.*, 1997), albeit at

concentrations far exceeding their ability to inhibit COX activity (Mitchell *et al.*, 1993).

LTB $_4$ was one of the first endogenous ligands described for PPAR α . The activation of PPAR α by LTB $_4$ causes a negative feedback pathway, whereby the β -oxidation pathway is induced leading to its own degradation (Devchand *et al.*, 1996). Interestingly, the PGD $_2$ dehydration product 15-deoxy- $\Delta^{12,14}$ -PGJ $_2$ (15d-PGJ $_2$), was the first endogenous ligand for PPAR γ discovered (Forman *et al.*, 1995; Kliewer *et al.*, 1995). Since then, components of oxidized low density lipoprotein (OxLDL), including 9- and 13-hydroxyoctadecadienoic acid (HODE), as well as the further oxidized derivatives have been described as endogenous PPAR γ activators (Nagy *et al.*, 1998). 12- and 15-hydroxyeicosatetraenoic acid (HETE) and 13-HODE are also metabolites of 12/15-LOX, utilizing either arachidonic acid (12 or 15-HETE) or linoleic acid (13-HODE)

Table 1 The pharmacological activators of PPARs

Class	Ligand	PPAR α	Activation PPAR δ	PPAR γ
Synthetic	WY14,643	++	+	(+)
	clofibrates	+	(+)	(+)
	troglitzone		-	++
	pioglitzone		-	++
	rosiglitzone	-	(+)	+++
Fatty acids	ETYA	++		
	docahexanoic acid	+	+	+
	linoleic acid	+	+	-
	arachidonic acid	+	+	+
Saturated fatty acids	C6-C18	(+)	(+)	
Saturated fatty alcohols	C14-C16	(+)		
FA CoA synthetase inhibitor	Triacsin C	(++)		
Carnitine palmitoyl-transferase-1 inhibitor	2Br-C16	++	(++)	-
FA-CoA dehydrogenase inhibitors	tetradecylglycidic acid	++	(++)	
	nonythioacetic acid	+		
	tetradecythioacetate	+		-
Eicosanoids	15d-PGJ $_2$	++	++	++
	PGJ $_2$	++	++	++
	prostacyclin (PGI $_2$)	++	++	(+)
	PGA $_{1/2}$	++	++	+
	PGB $_2$	++	(+)	+
	PGEs	-	-	-
	PGFs	-	-	-
	8-HEPE	++		
	8-(R)HETE	(++)	-	(+)
	8-(S)HETE	+++	-	(+)
	12-HETE	(++)		
	15-HETE			(+)
Oxidized LDL	LTB $_4$	(++)		
Native LDL				++
Oxidized linoleic acid products	9-(R/S) HODE			+
	13-(R/S) HODE			+
	13-(S)HpODE			+
	9-oxoODE			+
	13oxoODE			++
Others	NSAIDs	+		+

Table lists a number of synthetic ligands, fatty acids, and eicosanoids that can activate PPAR receptors. This data is based on a number of studies (Yu *et al.*, 1995; Brun *et al.*, 1996; Forman *et al.*, 1997; Kliewer *et al.*, 1997; Nagy *et al.*, 1998) which utilize, transcriptional activation through either PPRES, or GAL4-fusion proteins containing PPAR ligand binding domains, but not functional assays. Agonists are classed as activators of PPAR at: + + +, nM; ++, low μ M; +, high μ M; concentrations; - no effect; and a blank is not yet determined. As in most cases full dose-response curves have not been described, effects shown are those in systems showing highest potency. Where brackets '(' are present, indicates where a agonist tested at a particular concentration, is a weak activator e.g. LTB $_4$ will activate PPAR α transcription responses at 10 μ M by only 2-3 fold, while under the same conditions WY-14,643 within a similar concentration range (5 μ M) will activate approximately 20 fold (Forman *et al.*, 1997). Therefore, I have arbitrarily rated LTB $_4$ as (+ +), while WY-14,643 being a potent agonist at the same concentration ++. Abbreviations used in table; ETYA, eicosatetraenoic acid; 2Br-C16, 2-bromopalmitate; 15d-PGJ $_2$, 15-deoxy- $\Delta^{12,14}$ -PGJ $_2$; HEPE, hydroxyeicosapentaenoic acid; HETE, hydroxyeicosatetraenoic acid; HODE, hydroxyoctadecadienoic acid; HpODE, hydroperoxyoctadecadienoic acid; oxoODE, oxidized octadecadienoic acid; NSAID, non-steroidal anti-inflammatory drugs; LDL, low density lipoprotein.

as substrate (see Funk, 1996). The support for 12-, 15-HETE, and 13-HODE, as endogenous PPAR γ ligands has recently been greatly strengthened by the finding that 12/15 lipoxygenase activity induced by IL-4 also leads to the release of PPAR γ ligands (Huang *et al.*, 1999). As many of these endogenous mediators activate PPARs in the high nM, though most commonly in the low μ M range, it is still debatable whether they are the true 'endogenous' ligands. Furthermore, although these are primarily described as PPAR γ ligands, compounds, such as 15d-PGJ₂ in some transcription assays are as effective an activator of PPAR α or - δ , as they are of PPAR γ (Brun *et al.*, 1996). Though, with many of these unstable lipid hydroperoxide mediators the amounts required to activate PPARs by exogenous application, may far exceed those required from endogenous sources. Local production of an agonist may be of sufficient concentration to activate PPARs, while many of the lipids or fatty acids applied exogenously may be metabolized before they come in to proximity with the receptor. One piece of evidence that supports this, is a study of the oestrogen induced peroxisome proliferation mediated by PPAR γ in the duck uropygial gland. In this tissue oestrogen induces production of a PPAR γ activator, which is a PGD₂ metabolite and indistinguishable from Δ^{12} -PGJ₂ (Ma *et al.*, 1998). In contrast to this report, NIH3T3 cells which over-express ADD1, the rat homologue of the sterol regulatory element binding protein-1, release a PPAR γ agonist which on initial characterization was not 15d-PGJ₂ (Kim *et al.*, 1998). The identification of highly potent endogenous ligands for PPARs therefore still remains elusive.

Recently the crystal structure of the ligand binding domains of PPAR γ (Nolte *et al.*, 1998; Uppenberg *et al.*, 1998), and PPAR δ (Xu *et al.*, 1999) have been solved, revealing a large ligand binding pocket, which has led to the suggestion that maybe PPARs act as general lipid sensors to a broad spectrum of ligands, none of which having a particular high affinity. Since however, many of these eicosanoids and fatty acids are produced and act in the vasculature, or have been discovered in conjunction with the recent findings demonstrating PPARs in vascular and inflammatory cells, there has been a great deal of interest in these pathways as potential new targets for cardiovascular diseases.

Expression of PPARs and retinoid X receptors

PPARs act as transcription factors upon ligand induced heterodimerization with the common nuclear receptor binding partner, the retinoid X receptor (RXR; Mangelsdorf *et al.*, 1990; see Mangelsdorf & Evans, 1995). Since the original discovery of PPAR α from mouse liver (Issemann & Green, 1990), as a nuclear receptor that responded to peroxisome proliferators such as WY-14,643, nafenopin, and clofibrilic acid, the related family members, PPAR δ (PPAR β ; Dreyer *et al.*, 1992), and PPAR γ (Dreyer *et al.*, 1992) were discovered, and since found to be expressed in a number of species including mice (Issemann & Green, 1990; Zhu *et al.*, 1993; Kliewer *et al.*, 1994), humans (Schmidt *et al.*, 1992; Sher *et al.*, 1993; Mukherjee *et al.*, 1997), *xenopus* (Dreyer *et al.*, 1992) and rats (Gottlicher *et al.*, 1992). In adult rodents or man, PPARs are differentially expressed (see Table 2; Kliewer *et al.*, 1994; Braissant *et al.*, 1996; Mukherjee *et al.*, 1997). PPAR α is found predominantly in the liver, heart, kidney, brown adipose and stomach mucosa. PPAR γ is found primarily in adipose tissue, where it plays a critical role in the differentiation of pre-adipocytes into adipocytes, but also in the large intestine, spleen and heart, while PPAR δ is almost ubiquitously expressed. RXR also exists in multiple isoforms, RXR α , - β ,

Table 2 Relative tissue expression of PPAR and RXR isoforms

Tissue	Expression					
	PPAR α	PPAR δ	PPAR γ	RXR α	RXR β	RXR γ
Heart	+++	++	++	+	+++	+++
Liver	+++	+	+	+++	+	+++
Lung	+	+++	+/-	++	+++	+/-
Spleen	+	++	++	++	+++	-
Kidney	+++	+++	++	++	++	+
Brain	-	+++	+	+/-	+++	++
Intestine	+	++	++	+	+	-
Adrenal	++	++	++	+	+++	+
SK Muscle	+++	++	++	++	+++	+++

Studies describe either levels of mRNA or description of relative intensity of immunostaining for different PPAR (Mukherjee *et al.*, 1997; Kliewer *et al.*, 1994; Auboeuf *et al.*, 1997; Braissant *et al.*, 1996) or RXR (Mangelsdorf *et al.*, 1992) isoforms. For more details of PPAR expression in different organs and tissue see Braissant *et al.* (1996).

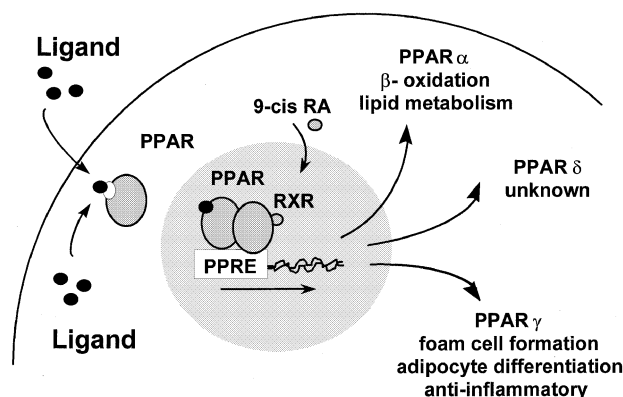


Figure 1 Activation of the PPAR receptors leads to an accumulation in the nucleus, where they heterodimerize with RXR. The PPAR:RXR heterodimer binds to DNA sequences called PPAR response elements (PPRE), leading to the transcription of the responsive gene. Activation of PPAR α can cause peroxisome proliferation, and increased lipid metabolism; PPAR γ causes cellular lipid accumulation in susceptible cells, and the down regulation of monocyte activation by cytokines; while the function of PPAR δ is not known.

and - γ (see Chambon, 1996), and like PPARs have a different tissue distribution (see Table 2; Mangelsdorf *et al.*, 1992). It is not known however, if any one of these particular RXR isoforms preferentially bind one or more of the PPAR isoforms.

Molecular mechanism of the PPARs

RXR isoforms are activated by 9-*cis* retinoic acid (Kliewer *et al.*, 1992), although other synthetic 'retinoid' ligands such as LG268 are also potent specific activators (Boehm *et al.*, 1995). As well as interacting with PPARs, RXR is a common binding partner for a number of other nuclear hormone receptors. RXR can homodimerize, or heterodimerize with thyroid hormone receptor, retinoic acid receptor, vitamin D receptor, as well as the orphan receptors liver X receptor, farnesol X receptor, and the pregnane X receptor (see Mangelsdorf & Evans, 1995). These receptors are activated by different ligands and lead to specific responses. In some combinations of RXR heterodimers (e.g.; RAR: retinoic acid receptor), the RXR ligand does not contribute to activation of a gene by the RXR heterodimer. However, when combined as a PPAR:RXR

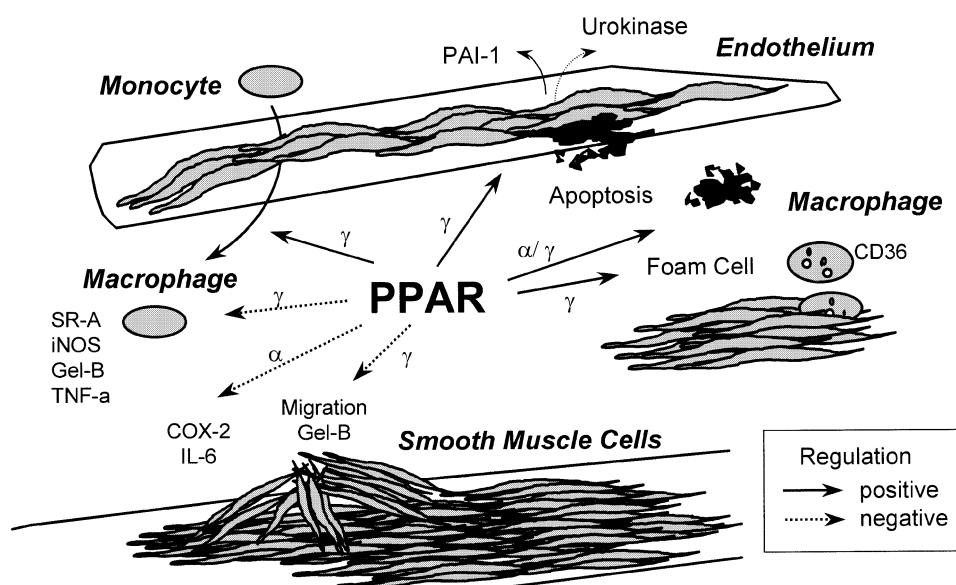


Figure 2 PPAR α and PPAR γ regulate vascular and inflammatory cell functions. Many different, often contradictory effects have been ascribed to PPAR ligands in vascular, and inflammatory cells. PPAR α , and γ are expressed in endothelial cells, monocytes/macrophages, and in the vascular smooth muscle cells of both medial and intimal layers. PPAR α ligands inhibit smooth muscle cell production of inflammatory products, and causes macrophage apoptosis. PPAR γ ligands (i) inhibit smooth muscle cell migration processes; (ii) inhibit monocyte/macrophage production of inflammatory enzymes (iNOS; inducible nitric oxide synthase; Gel-B, gelatinase B), cytokines, and scavenger receptor (SR)-A expression; (iii) induce monocyte/macrophages differentiation, and uptake of oxidized LDL; (iv) induce monocyte and endothelial cell apoptosis; (v), inhibit the expression of vascular endothelial cell growth factor receptors, endothelin-1 and urokinase expression in endothelial cells and, (vi) induce plasminogen activator inhibitor-1 expression in endothelial cells.

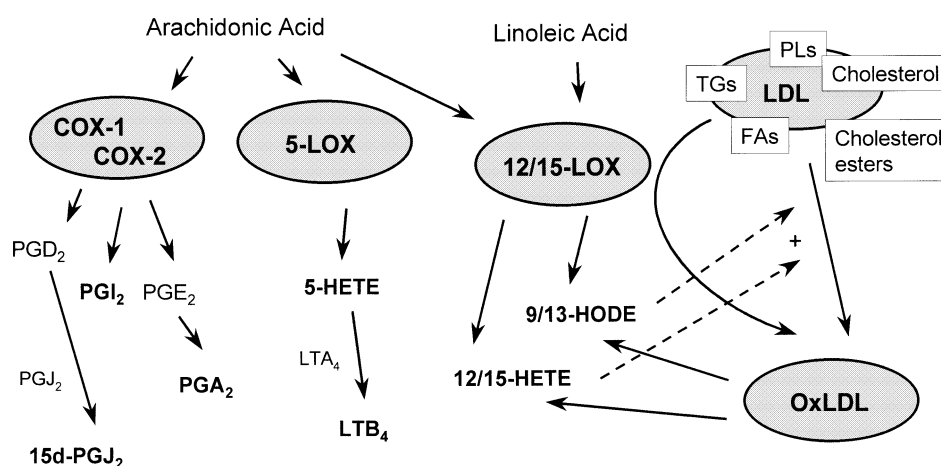


Figure 3 PPAR ligands as products of cyclo-oxygenase (COX), lipoxygenase (LO), or oxidized low density lipoprotein (oxLDL). Little is actually known about which mediators act as true endogenous ligands, therefore the figure describes some of the potential pathways for the production of PPAR ligands. Directly binding PPAR ligands are noted in bold text. Constitutive (COX-1), or inducible (COX-2), present in vascular and inflammatory cells, can utilize primarily arachidonic acid to form PGH₂. PGH₂ is metabolized further to a number of prostanoid mediators, some of which, 15-deoxy- $\Delta^{12,14}$ -PGJ₂ (15d-PGJ₂, a dehydration product of PGD₂), PGA₂ (dehydration product of PGE₂), and PGI₂ (prostacyclin) that activate PPAR receptors. 5-LO, present in leukocytes, forms from arachidonic acid 5-HpETE, the precursor for 5-HETE, which itself is the precursor for leukotrienes (LT)s, of which LTB₄ is also a PPAR(α) agonist. 12/15-LOs, whose presence is strongly implied in the pathogenesis of atherosclerosis can produce PPAR agonists from arachidonic acid, the 12- and 15-HETEs, and form linoleic acid, 9- and 13-HODE. Low density lipoprotein LDL, which contains a number of phospholipids (PL)s, triglycerides (TG)s, cholesterol, cholesterol esters, and fatty acids, can be directly oxidized by 12/15-LOs, to produce PPAR ligands including 9- and 13-HODE. Furthermore, these unstable lipid hydroperoxides, 9- and 13-HODE, and the 12- and 15-HETEs, may themselves directly contribute to oxidation of LDL through a non-enzymatic pathway.

heterodimer, PPAR ligands and 9-*cis* retinoic acid can act synergistically on PPAR responses (Kliewer *et al.*, 1992; see Figure 1). The way in which the different dimers of RXR allow specific responses to follow is by binding highly specific sequences in the promoter regions of the various genes they are to trans-activate (see Mangelsdorf & Evans, 1995). The specificity of these dimers in inducing different gene

transcription is due to subtle differences in the 'response element' promoter region they are able to bind. These response elements for nuclear hormone receptors have a general patterns of either direct repeats, palindromic or inverse palindromic sequences separated by one or more nucleotides (see Glass, 1994). For all the PPARs, the PPAR:RXR preferentially binds a direct repeat of the consensus sequence

'AGGTCA' separated by 1 nucleotide (or a DR1 sequence) called the PPAR response element (PPRE). For example the rat acyl Co oxidase gene which has the first characterized PPRE, has the sequence AGGaCA A AGGTCA (Tugwood *et al.*, 1992). Though, for all the RXR, nuclear receptors heterodimers, each has its own particular affinity to promoter regions of direct repeated sequences spaced by 1–5 nucleotides (called the 1–5 rule; see Mangelsdorf & Evans, 1995).

Although the PPAR:RXR dimer is the focus for determining specific gene transcription upon ligand activation, transactivation of a particular gene requires a large complex of proteins. Formation and components of such complexes have been reviewed extensively elsewhere (Torchia *et al.*, 1998; Bjorkland *et al.*, 1999). In the un-activated state the PPARs are thought to be in complexes bound with co-repressor proteins such as the silencing mediator for RXR and thyroid receptor or the nuclear receptor co-repressor. Furthermore, at this stage, in some but not all cell types, PPARs may also have a cytoplasmic rather than nuclear sub-cellular location (Chinetti *et al.*, 1998; Bishop-Bailey & Hla, 1999). Upon ligand activation, PPARs dissociated from co-repressors and recruit co-activators, including the PPAR-binding protein (Zhu *et al.*, 1997) and the steroid receptor co-activator-1 (Zhu *et al.*, 1996), and can translocate from the cytoplasm to the nucleus (Bishop-Bailey & Hla, 1999). Indeed, transcription factors act in large complexes, which facilitates their active recruitment to the correct site and orientation within the promoter region of the target gene. Many of the proteins known to be associated in such complexes also contain histone deacetylase activity, to allow for chromatin remodelling and access for the RNA polymerase (see Torchia *et al.*, 1998; Bjorkland *et al.*, 1999).

As the different PPARs have very specific yet diverse actions it is still uncertain how PPAR α , δ and γ all act on the same DR-1 consensus sequence. Apart from a difference in tissue expression, it is likely that PPARs gain selectivity in part by acting on slight differences in the PPREs from that of the consensus sequence. Indeed, *in vitro* study of the binding of PPAR α , δ and γ :RXR heterodimers to PPREs, indicates that although PPAR α , δ and γ equally bind the rat acyl CoA oxidase PPRE, the PPAR γ :RXR heterodimer preferentially binds aP2 (a gene involved in the adipogenesis gene activation program) PPREs (Brun *et al.*, 1996). Though this may still be a simplification of how specificity is achieved. Over-expression of PPAR δ can inhibit PPAR α , or thyroid hormone receptor mediated transcriptional activation, by competition for the RXR binding partner (Jow & Mukherjee, 1995). Furthermore, two high affinity partial agonists for PPAR γ have been recently described, MC-555 (Reginato *et al.*, 1998a), and GW0072 (Oberfield *et al.*, 1999). MC-555 and GW0072 are believed to act by virtue of causing a conformational change in the receptor that can only weakly recruit the co-activators; steroid receptor co-activator-1, PPAR binding protein (Oberfield *et al.*, 1999), or cyclic AMP response element binding protein, 'binding protein' (Reginato *et al.*, 1998a). Apart from being a partial agonist, GW0072, acts to antagonize full agonist (rosiglitazone) induced adipocyte differentiation (Oberfield *et al.*, 1999). A specific response through the PPAR pathway, may therefore be the result of a combination of tissue specificity, competition with other binding partners for the RXR, and the ability of the particular ligand to induce active receptor conformation, and recruit different co-activation complexes. This complexity when fully unravelled will however lead to the possibility of a number of targets for pharmacological or genetic intervention.

PPARs in cardiovascular disease

Over the last few years PPARs have been shown to play an important role in energy homeostasis, diabetes (see Spiegelman, 1998) and obesity (Ristow *et al.*, 1998), cellular differentiation, including considerable inhibitory effects, more recently described on tumour growth (Kubota *et al.*, 1998; Mueller *et al.*, 1998; Brockman *et al.*, 1998; Demetri *et al.*, 1999; Sarraf *et al.*, 1998). For interested readers, the cited reviews, and articles give a good overview of the current published data. In terms of this review, the recent data on the expression of PPARs and potential ligands in vascular, and inflammatory cells will be discussed in more detail.

Monocytes/macrophages Macrophages are the predominant inflammatory leukocyte in a number of chronic inflammatory diseases including atherosclerosis (see Ross, 1999). In the atherosclerotic lesion they are not only believed to contribute to the inflammation at the level of 'classical' mediator secretion, but they may also become lipid laden foam cells, and through the release of oxidants, or production of oxidizing enzymes, contribute to the formation of the highly damaging oxidized low density lipoproteins (OxLDL; see Berliner & Heinecke, 1996).

PPAR γ is induced in monocytes upon their differentiation *in vitro* or *in vivo* to macrophages (Ricote *et al.*, 1998a,b). Indeed, isolated monocytes or myeloid precursors activated by PPAR γ and RXR ligands, have increased macrophage/monocyte markers CD11b, CD18, and CD14 (Tontonoz *et al.*, 1998). Moreover, monocytes/macrophages treated with PPAR γ and RXR ligands increase lipid accumulation and express the OxLDL scavenger receptor CD36 (Tontonoz *et al.*, 1998). PPAR γ may therefore be involved in the differentiation from the myeloid precursors, of monocytes, to macrophages, further to foam cells (Tontonoz *et al.*, 1998). LDL particles are a rich potential source of fatty acids, and eicosanoid precursors, which may be potential PPAR ligands. Apart from the apoB core protein, LDL is rich in phospholipids, cholesterol, cholesterol esters, and triglycerides (see Steinberg, 1997). In a closely related article, these authors also demonstrated that OxLDL and the components 9- and 13-HODE, and oxidized derivatives were PPAR γ ligands, that induced macrophage foam cell formation (Nagy *et al.*, 1998). Furthermore, similar to the other PPAR γ ligands, OxLDL and its components induce the OxLDL scavenger receptor CD36 on macrophages, indicating a positive feedback for lipid accumulation (Nagy *et al.*, 1998), potentially explaining the ability of PPAR γ to induce foam cell formation. In slight contrast, in activated macrophages, the stimulation of PPAR γ by ligands also results in a global inhibition of inflammatory mediator production from these cells. PPAR γ ligands inhibit IFN- γ induced morphogenesis, inducible nitric oxide synthase induction, gelatinase B and scavenger receptor-A expression (Ricote *et al.*, 1998a), and the release of the pro-inflammatory cytokines TNF- α , IL-6, IL-1 β and IL-2 (Jiang *et al.*, 1998). Further analysis revealed that PPAR γ can actually trans-repress the activity of these inflammatory mediators at the level of transcription, by inhibiting nuclear factor (NF)- κ B, Stat1, and activation protein-1 signalling (Ricote *et al.*, 1998a); some of the central transcription factors implied in inflammatory responses.

In differentiated macrophages, at least *in vitro*, PPARs have yet a different function, in that both PPAR α and γ ligands induce macrophage apoptosis (Chinetti *et al.*, 1998). Little is known about how PPARs act in other inflammatory leukocytes, though murine T-helper₁ lymphocytes express at

least PPAR γ , activation of which leads to the inhibition of antigen or T-cell receptor induced proliferation (Clark *et al.*, 1999). Human neutrophils, and peripheral blood lymphocytes express a truncated PPAR γ message, which does not fully code the protein (Greene *et al.*, 1995).

PPAR γ and α are clearly expressed in monocytes/macrophages *in vitro*. *In vivo* a high expression of PPAR γ is also observed in experimental (Tontonoz *et al.*, 1998) and human atherosclerotic lesions (Ricote *et al.*, 1998b; Marx *et al.*, 1998b) predominantly in the macrophages, but also to the endothelial cell layer, and the vascular smooth muscle (more strongly in the intimal than medial layer). PPAR α and PPAR δ are also expressed in atherosclerotic lesions (D Bishop-Bailey, E Smith, C Haudenschild and T Hla, unpublished observations). PPAR α has a similar expression pattern to PPAR γ , while PPAR δ expression is much weaker, being sparsely expressed in the endothelial cell layer, intimal but not medial smooth muscle cells, and in adherent or infiltrated cells, presumably monocytes or macrophages.

Previous findings in the macrophage would appear to suggest opposing sides of PPAR γ activation, an anti-inflammatory effect, and a pro-atherogenic effect. Which of these many functions ascribed to PPARs in monocytes/macrophages predominates *in vivo* is yet to be ascertained. Though the effects subsequently demonstrated in vascular smooth muscle cells and endothelial cells, again show multiple sometimes opposing actions.

Vascular smooth muscle Vascular smooth muscle contains PPAR α (Staels *et al.*, 1998a; Marx *et al.*, 1998a), PPAR γ (Ricote *et al.*, 1998b; Marx *et al.*, 1998a) and PPAR δ (D Bishop-Bailey, unpublished observations in rat aortic vascular smooth muscle cells, by Northern blot analysis, and immunofluorescence). Before the realization of the existence of PPARs in vascular smooth muscle cells, and that thiazolidinediones were potent PPAR γ activators, troglitazone was shown to inhibit vascular smooth muscle proliferation and migration *in vitro*, and restenosis in a model of balloon angioplasty *in vivo* (Law *et al.*, 1996). Indeed, in this study troglitazone also inhibited *c-fos* induction, and serum response element induced transcriptional activation. These effects of troglitazone are presumably through PPAR γ , though direct effects on Ca²⁺ channels have also been noted (Zhang *et al.*, 1994; Song *et al.*, 1997). Subsequently, PPAR γ expression was demonstrated in human vascular smooth muscle cells, and PPAR γ , but not PPAR α agonists, inhibited platelet derived growth factor-BB induced migration (Marx *et al.*, 1998a). One of the mechanisms required for cellular migration is the degradation of extracellular matrix. Similar to macrophages (Ricote *et al.*, 1998a), troglitazone, and 15d-PGJ₂ selectively inhibited phorbol ester induced vascular smooth muscle cell expression of gelatinase-B (matrix metalloproteinase-9). The expression of matrix metalloproteinase-2, or tissue inhibitors of matrix metalloproteinase-1 or -2 was unaffected (Marx *et al.*, 1998a).

In contrast PPAR α , but not PPAR γ ligands, inhibit inflammatory responses in vascular smooth muscle cells by repressing NF- κ B signalling (Staels *et al.*, 1998a). Indeed, by this mechanism, fenofibrate, and WY-16,463 inhibits IL-1 induced IL-6 production, and inducible cyclo-oxygenase (COX-2) expression (Staels *et al.*, 1998a). These results are consistent with findings in the PPAR α knockout mouse which has increased inflammatory responses, though this is considered in part due to the reduced capacity to metabolize inflammatory mediators through β -oxidation (Devchand *et al.*, 1996). Moreover, hyperlipidaemic patients treated with fenofibrate also had reduced circulating IL-6 and acute phase

proteins (Staels *et al.*, 1998a). Although, it is unclear if fenofibrate is exclusively a PPAR α agonist *in vivo*, a number of beneficial cardiovascular effects may result from PPAR α activation in the liver, which is the central organ regulating lipid metabolism and lipoprotein expression. As recently reviewed by Staels *et al.* (1998b) PPAR α activation in the liver by the fibrate class of drugs leads to the increase in protective high density lipoproteins, while decreasing detrimental LDLs and VLDL synthesis, enhances lipolysis, and increases fatty acid catabolism.

Both PPAR α and γ appear to have protective effects on the activity of vascular smooth muscle cells. PPAR α ligands, reduce the inflammatory response within the vessel wall by at least interfering with NF- κ B signalling, while PPAR γ ligands seem to specifically inhibit smooth muscle migration, by interfering with processes involved in the degradation of the extracellular matrix.

Vascular endothelial cells Vascular endothelial cells contribute to the release of protective mediator in large blood vessels, where endothelial damage or dysfunction is considered one of the potential mechanisms for development of atherosclerosis and restenosis (see Drexler & Hornig, 1999). In the microvasculature, the endothelium contributes to chronic inflammatory processes, wound healing, and tumour formation by an angiogenic response, vascularizing the newly produced tissue (Folkman, 1995). Vascular endothelial cells contain PPAR α (Inoue *et al.*, 1998; Xin *et al.*, 1999; Bishop-Bailey & Hla, 1999), PPAR δ (Xin *et al.*, 1999; Bishop-Bailey & Hla, 1999), and PPAR γ (Marx *et al.*, 1999a; Xin *et al.*, 1999; Bishop-Bailey & Hla, 1999), though, as yet no definitive roles for PPAR δ have been described. PPAR α agonists, similar to their effects in vascular smooth muscle cells inhibit NF- κ B signalling, by which mechanism they can inhibit TNF- α induced expression of vascular cell adhesion molecule-1, and the subsequent adherence of monocytes (Marx *et al.*, 1999b). Like other cell types PPAR γ appears to have both potential protective and detrimental effects. In terms of large vessel disease, PPAR γ activators inhibit the endothelial cell release of endothelin-1 (Satoh *et al.*, 1999; Delerive *et al.*, 1999), a potent vasoconstrictor, and vascular smooth muscle cell mitogen (see Ruschitzka *et al.*, 1997). In contrast, PPAR γ agonists also increase plasminogen activator inhibitor (PAI)-1 expression (Marx *et al.*, 1999a; Xin *et al.*, 1999). PAI-1 is highly expressed in adipocytes, and levels correlate with obesity, furthermore, increased PAI-1 is associated with myocardial ischaemia, and thrombosis (Loskutoff & Samad, 1997). A role which is unsurprising since PAI-1 inhibits two of the enzymes which play a major role in fibrinolysis, tissue plasminogen activator and urokinase. The reverse side to the inhibition of PAI-1 in large vessels, is that in small vessels increasing PAI-1 (Xin *et al.*, 1999) may be one of the mechanisms responsible for inhibition of angiogenesis by PPAR γ agonists (Xin *et al.*, 1999; Bishop-Bailey & Hla, 1999). Furthermore, PPAR γ agonists also reduce the vascular endothelial cell growth factor receptors Flt-1, and Flk/KDR, and inhibit the expression of urokinase (Xin *et al.*, 1999). Alternatively, PPAR γ activators can inhibit endothelial cell angiogenesis by inducing apoptosis (Bishop-Bailey & Hla, 1999), similar to macrophages (Chinetti *et al.*, 1998), via a caspase-3 mediated process. Though again in contrast to the microvasculature, in a large vessel disease, inappropriate apoptosis may cause structural weakness in an atherosclerotic lesion, and may promote plaque rupture (Newby & Zaltsman, 1999), which may lead to embolism or stroke.

In the microvasculature PPAR γ may be a novel target for anti-angiogenic therapy for a number of tumours or chronic

inflammatory disorders. Like other cells, PPAR γ activation can inhibit cytokine induced mediator release, which may be beneficial in large vessel disease. Many of the effects described which would be beneficial in the microvasculature however, inhibiting endothelial cell function, may naturally be detrimental to protecting against atherogenic stimuli.

The heart Although, the heart contains very high levels of all PPAR and RXR receptors (see Table 2), very little is known about how PPAR ligands effect cardiac function. Certainly PPAR α appears to have an important role in mitochondrial fatty acid β -oxidation (Brandt *et al.*, 1998; Yu *et al.*, 1998). Indeed, in PPAR α knockout mice, once mitochondrial fatty acid import is blocked by the metabolic inhibitor etomoxir, the mice undergo a hypoglycaemia, and a massive increase in lipid accumulation in the heart and liver, resulting in 100% mortality of male mice (Djouadi *et al.*, 1998); females being largely (75%) protected by oestrogens (Djouadi *et al.*, 1998). Nothing however is known with regards to the potential role of PPARs in the pathologies of the heart, or cardiac tissue.

Potential sources of PPAR ligands in the cardiovascular system

Although PPARs are present in many different cell types, and are activated by a large variety of ligands, very little is known about the 'endogenous' pathways for PPAR activation. The homeostasis of the vasculature and the pathogenesis of vascular diseases such as atherosclerosis involves regulation of fatty acid metabolism, the uptake and accumulation of oxidized lipids, and the induction of inflammatory responses. With so many PPAR ligands now discovered (see Table 2), there are several potential pathways to form these agonists. While fatty acids are essential ubiquitous components of cell systems, other endogenous pathways for PPAR ligand production may be associated with other 'house-keeping' roles of cells, or subsequent disease processes. Indeed, vascular and inflammatory cells express, and can be induced to express cyclo-oxygenase and lipoxygenase enzymes, and many disease processes are associated with increases in oxidized lipids.

Cyclo-oxygenases A number of COX products are PPAR agonists (see Table 2), including 15d-PGJ $_2$, a dehydration product of PGD $_2$, as well as PGI $_2$, one of the major products of vascular COXs (see Bishop-Bailey *et al.*, 1999). Constitutive COX-1 is abundant in endothelial cells, while COX-2 can be readily induced in human endothelial cells (Hla & Nielson, 1992) and smooth muscle cells (Bishop-Bailey *et al.*, 1997; 1998) to produce large quantities of prostanoids, of which PGI $_2$ and PGE $_2$ have protective vascular functions (see Bishop-Bailey *et al.*, 1999). Similarly, monocytes/and macrophages express low levels of COX-1, and can also be readily induced to express high levels of COX-2 and produce large quantities of prostanoids (O'Banion *et al.*, 1992). Moreover, COX-2 is highly expressed in human atherosclerotic lesions (Baker *et al.*, 1999), and both COX-1 (Narko *et al.*, 1997) and COX-2 (Tsuji *et al.*, 1998) have been implicated in tumour formation, and angiogenesis. The pattern of prostanoid release depends on the presence of secondary metabolizing enzymes. However in the absence of secondary enzymes, PGF $_{2\alpha}$, PGD $_2$ and in particular PGE $_2$ can be formed non-enzymically (see Smith *et al.*, 1991). Vascular endothelial cells, and vascular smooth muscle contain large amounts of PGI $_2$ synthetase (Smith *et al.*, 1983), though in atherosclerotic lesions, synthetic smooth muscle cells also express the 'neuronal' lipocalin type PGD synthase (Eguchi *et al.*, 1997). Although, COX preferentially metabolizes arachi-

donic acid, it can also use a number of other free fatty acids such as 8,11,14-eicosatrienoic acid and linoleic acid as alternative substrates. For this reason COX can also form, under certain conditions, members of other biologically active lipid families, 11-, and 15-HETE; 9- and 13-HODE; 12-hydroxyheptadecatrienoic acid; and isoprostanes (see Bishop-Bailey *et al.*, 1999). Interestingly, a number of COX products which can activate the MAP kinase pathways, such as PGF $_{2\alpha}$ may also indirectly regulate PPAR pathways at the level of receptor phosphorylation (Reginato *et al.*, 1998b).

Lipoxygenase LOXs are a large family of enzymes which can metabolize mainly arachidonic acid, to a number of biologically active mediators (see Funk *et al.*, 1996); a number of which have since been demonstrated to activate PPARs. The three main classes of LOXs are those with 5-LOX, 12-LOX, or 15-LOX activity, which are expressed in a variety of primarily inflammatory cell types, platelets, epidermal tissue (see Funk *et al.*, 1996), and cardiomyocytes (Breitbart *et al.*, 1996). 5-LOX metabolize arachidonic acid to 5-HPETE, the precursor for LTs (LTB $_4$ of which is a PPAR α agonist), 12-LOX (rodent homologue of 15-LOX) metabolizes arachidonic acid to 12-HPETE >> 15-HPETE, while 15-LOX conversely metabolizes arachidonic acid to 15-HPETE >> 12-HPETE, precursors for their respective HETE moieties (see Funk, 1996). Arachidonic acid is not however the only substrate for 12/15-LOX, as both linoleic acid (metabolized to 13(S)HODE), and cholesteryl linoleate, one of the major lipid components of LDL can be utilized. Furthermore, it has also been suggested that LOX activity, may initiate oxidation of LDL, into toxic OxLDL (see Steinberg, 1997). 15-LOX is found upregulated, and co-localized to lipid rich atherosclerotic lesions (Yla-Herttuala *et al.*, 1990; 1991), both in the macrophage, and in the endothelial cell layer. Murine fibroblasts which over-express 15-LOX enhance lipoperoxides production with LDL as substrate (Benz *et al.*, 1995). Furthermore, apoE/ 12/15-LOX double knockout mice have reduced lesion formation (Cyrus *et al.*, 1999), compared to apoE knockouts alone. Though, LOXs can produce PPAR ligands *in vitro* (Huang *et al.*, 1999), and are strongly implicated in the progression of a number of diseases, including atherosclerosis, it is still uncertain if these 5-, 12-, or 15-LOX products serve as functional PPAR ligands in these processes.

Oxidized LDL and derivatives The atherosclerotic lesion is known to contain large amounts of oxidized esterified lipids, from OxLDL, 13-HODE (see Feinmark & Cornicelli, 1997), to 5-, 8-, 9-, 11-, 12- and 15-HETEs, 5,6-, 8,9-, 11,12-, and 14,15-epoxyeicosatetraenoic acids, F series isoprostanes (Mallat *et al.*, 1999). Sources of endogenous ligands LDL oxidized derivatives. It is as yet uncertain, how much of these lipids like 13-HODE are produced by enzymatic processes, 15-LOX produces specifically 13(S)-HODE, while 13-HODE found in lesions is primarily racemic, indicative of a non-enzymic mechanism. Such non-enzymic pathways include, oxidation by superoxide anion, by free Fe $^{2+}$ or Cu $^{2+}$ ions, or by the reactive intermediates of hydroperoxides, such as 15-HETE formed by LOXs (see Sigal *et al.*, 1994). Thereby, enzymes such as 15-LOX, myeloperoxidase or even cyclo-oxygenase can still contribute indirectly to lipid peroxidation, as opposed to directly utilizing LDL as substrate, by the mediators it releases from the cell. As 15-LOX is intracellular, while lipids largely remain extracellular indeed, this may be the primary mechanism by which 15-LOX causes oxidation of lipids within a lesion.

Conclusion

The early descriptions of PPARs limited their activity to specialist functions, such as β -oxidation and peroxisome proliferation in the liver (PPAR α), and adipogenesis (PPAR γ) in specialist cell types. It now appears that PPARs, are more widespread, albeit it is often at lower levels in vascular, and inflammatory cells. Furthermore, their actions apart from their classical roles in β -oxidation, and adipogenesis, seemed to have extended, where they now have important contributions in the control of inflammatory responses, cell growth and differentiation. Although the actions of PPAR α and PPAR γ are becoming clearer, it is still uncertain what roles PPAR δ has, especially in the vasculature. Though this may become clearer as PPAR δ selective ligands are developed and become available. Furthermore, PPAR γ seem to have a number of conflicting actions within inflammatory and vascular cells, appearing to both be able to have anti-inflammatory actions, as well as inducing foam cell formation, and causing apoptosis. Which of these roles dominates *in vivo* is still unsure, though very likely to be solved in the near future with the use of the potent selective agonists now available. Since now the crystal structures for PPAR γ and PPAR δ have been solved, it should

help to progress the development further of selective agonists, and may certainly help the development of antagonists for these receptors. PPARs seem to be able to respond to a vast number of ligands, from fatty acids, to eicosanoid and linoleic acid metabolites. One of the major prospectives in this field may be to find if these receptors are general lipid sensors, or in fact there are as yet unidentified highly potent endogenous ligands. PPARs have now been demonstrated in a number of diseases, including atherosclerosis, diabetes, inflammation and certain cancers, where the use of selective ligands has often led to beneficial effects. The newly discovered existence of PPARs in vascular and inflammatory cells, opens up the possibility that modulation of these nuclear hormone receptors in the cardiovascular system may have a large therapeutic potential.

D. Bishop-Bailey currently holds a British Heart Foundation Intermediate Fellowship. Many thanks to Dr Timothy Hla and Mr Philippe Delerive for useful discussions, and to Dr Nicolas Ancellin and Dr Robert Clark for their critical reading of the manuscript.

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(Received July 6, 1999
Revised December 8, 1999
Accepted December 9, 1999)