

Cross-talk between heme oxygenase and peroxisome proliferator-activated receptors in the regulation of physiological functions

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1. ABSTRACT

Peroxisome-proliferator-activated-receptors (PPARs) are transcription factors belonging to the superfamily of nuclear receptors. The isoforms of PPAR include PPAR alpha, PPAR gamma and PPAR delta (also known as PPAR beta). Generally, PPARs potentiate insulin sensitivity, improve glucose/lipid metabolism, suppress inflammation/oxidative stress, attenuate excessive immune responses, regulate cell-growth and differentiation. Interestingly, agonists of PPAR gamma and PPAR alpha have been shown to upregulate the heme-oxygenase (HO)-system. Conversely, the HO-system also enhances PPAR alpha, and potentiates the expression and activity of PPAR γ . Moreover, the HO-system and related products including bilirubin, biliverdin, carbon monoxide and ferritin have been shown to increase insulin sensitivity, improve glucose/lipid metabolism, suppress inflammation/oxidative stress, abate immune response, and modulate cell-growth/differentiation. Therefore, an intimate, reciprocal, stimulatory and synergistic relationship between PPAR-signaling and the HO-system can be envisaged in the regulation of physiological functions. Thus, both the HO-system and PPARs-signaling participate in fine-tuning similar physiological functions, so novel pharmacological agents capable of optimizing this interaction should be sought. The coordinated regulation of PPAR-signaling and the HO-system may constitute the basis for future drug design.

2. INTRODUCTION

Transcription factors are biosensors that modulate physiological processes by converting signals generated by cells into gene expression after binding to specific DNA sequence in gene promoters. Interestingly, a specific signaling pathway can activate multiple transcription factors, and conversely, the expression of a specific gene may be controlled by a wide spectrum of different transcription factors.

Peroxisome-proliferator-activated-receptors (PPARs) are transcription factors belonging to the superfamily of nuclear receptors and are activated upon binding to specific ligands which are generally small lipophilic molecules (1-11). The superfamily of nuclear receptors include: (i) steroid hormones like progesterone, estrogen, glucocorticoids and mineralocorticoids, (ii) thyroxine, retinoic acid and vitamin D, and (iii) PPARs agonists (1-11). The three commonly described isoforms or subtypes of PPAR includes PPAR alpha, PPAR gamma and PPAR delta (sometimes referred to as PPAR beta) (1-5, 11). PPARs have different tissue distribution and are important regulators of physiological events (1-5, 12-68).

PPARs are intracellular receptors that upon interaction with agonists or ligands like fatty acids or their derivatives translocate to the nucleus to modulate gene transcription. Within the nucleus, PPARs exist as obligate

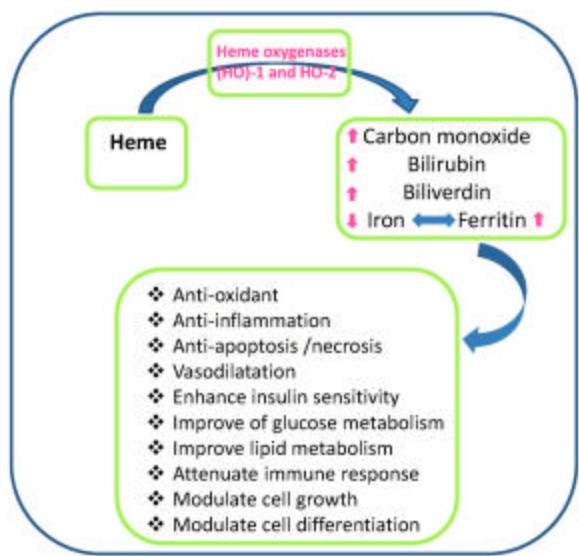


Figure 1. Summary of diverse role of PPARs. The three isoforms of PPAR, namely PPAR alpha, PPAR gamma and PPAR delta (sometimes referred to as PPAR beta) are encoded by distinct genes and are implicated in the regulation of several physiological functions including the potentiation of glucose/lipid metabolism, suppression of inflammatory/oxidative insults, attenuation of immune response, abrogation of apoptosis/necrosis, modulation of vascular tone and the regulation of cell growth and differentiation.

heterodimers with retinoid-X-receptor (RXR) and are anchored to DNA with co-repressor molecules. Upon activation by ligands, the heterodimers undergo conformational changes that cause the dissociation of co-repressors and the recruitment of different transcriptional co-activators and/or co-activator-related proteins, with subsequent modulation of gene transcription, protein synthesis and specific cellular functions (6, 25, 40, 69-75). Generally, PPARs are implicated in a wide range of physiological functions including enhancement of insulin sensitivity, improvement of glucose and lipid metabolism, suppression of inflammatory/oxidative insults, modulation of the immune response and the regulation of cell growth and differentiation (1-5, 11, 29-38).

There is an intimate reciprocal relationship between many transcription factors and the heme oxygenase (HO) system (76-88). The HO system and related products including carbon monoxide, bilirubin, biliverdin and ferritin have gained significant recognition in the regulation of several physiological functions such as insulin sensitivity, glucose metabolism, lipid metabolism, inflammation/oxidative stress, immune response, apoptosis, cell growth and differentiation (89-136). Emerging evidence indicates that many of the effects modulated by the HO system are mediated through different transcription factors, amongst which peroxisome proliferator-activated receptor (PPARs) is of particular interest given its reciprocal, stimulatory and synergistic relationship with the HO system (76-88). HO is a microsomal enzyme that

catalyzes the breakdown of the pro-oxidant heme, generating cytoprotective products including biliverdin, bilirubin and carbon monoxide (89, 137-139). The two principal isoforms of HO are HO-1 (inducible) and HO-2 (constitutive), while the third isoform HO-3, is a pseudotranscript of HO-2, so HO activity is largely determined by inputs from HO-1 and HO-2 isoforms (140-142).

Recent studies indicate that HO-1 gene promoter also contains motifs for PPARs response element (82, 84, 85), suggesting that the HO system may regulate PPAR. This notion is further strengthened by reports indicating that the HO system potentiates PPAR alpha (81, 82, 84, 85). On the other hand, the PPAR alpha agonist and lipid-lowering drug, fibrates, have been shown to enhance HO-1 (76). The interaction between the HO system and PPARs may constitute the basis of many beneficial effects against insulin resistant diabetes, hypertension, obesity and related cardiometabolic complications. Thus, the coordinated activity of HO system and PPARs may be important for fine tuning physiological functions and the maintenance of cellular homeostasis.

3. PEROXISOME PROLIFERATOR-ACTIVATED RECEPTOR (PPAR)

PPARs are transcription factors belonging to the superfamily of nuclear receptors, and are known to enhance glucose/lipid metabolism, abate inflammatory/oxidative insults, attenuate immune response, suppress apoptosis/necrosis, increase vasodilatation and regulate cell growth and differentiation (Figure 1) (1-11, 29-38, 43-47, 63-68, 143-163).

Like other nuclear receptors, PPARs are activated upon binding to specific ligands or agonists which are generally small lipophilic molecules (1-11) including: (i) steroids like progesterone, estrogen, glucocorticoids and mineralocorticoids, (ii) thyroxine, retinoic acid and vitamin D, and (iii) PPARs agonists (1-11). Upon activation, PPARs undergo conformational changes, and recruit co-activators or release co-repressors, which determines the functional state of the receptor-ligand transcriptional complex and the expression of distinct target genes (1-11). The activities of PPARs are regulated by complex mechanisms involving: (i) ligand-selective and cell-specific interactions with PPAR-binding proteins, (ii) cyclic guanosine monophosphate (cAMP)-response element-binding protein, (iii) different cofactors, (iv) heterodimerization with members of the RXR family, and (v) the activation of kinases and phosphatases to phosphorylate and dephosphorylate a wide variety of substances such as mitogen-activated protein kinases (MAPK), protein kinase A (PKA), protein kinase C (PKC), adenosine monophosphate-activated protein kinase (AMPK), glycogen synthase kinase-3 and other co-factors (6, 25, 40, 69-75).

The three commonly described isoforms of PPAR includes PPAR alpha, PPAR gamma and PPAR delta (sometimes referred to as PPAR beta) (1-6, 11). The

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different isoforms are encoded by distinct genes and their distribution pattern in tissues is different (39-47). For example, PPAR alpha is highly expressed in metabolically active tissue like the skeletal muscle, liver, kidneys, heart, brown adipose tissue and intestinal mucosa (39, 41). Similarly, PPAR gamma is widely expressed in tissues like the large intestines, white adipose tissue, spleen and brown adipose tissue (39, 42), although the expression of PPAR gamma is particularly high in adipose tissue (40). On the other hand, the distribution of PPAR delta is more ubiquitous than PPAR alpha, PPAR gamma (43, 44, 46, 47), and has been reported in a variety of tissues and/or cells across different organs and systems including nervous, hematopoietic, reproductive, immune, cardiovascular, urinary, respiratory, digestive, endocrine, musculoskeletal, sensory and the islet of Langerhans systems (43-47). All PPAR isoforms are composed of a C-terminal ligand-binding domain and an N-terminal DNA-binding domain linked to each other by a hinge region for the interaction of specific PPAR response elements in the promoter region of PPAR-regulated target genes (1-6, 11). Given that each PPAR isoform controls the expression of multiple genes, the activation of PPAR alpha, PPAR gamma and PPAR delta by a given upstream signal, whether a synthetic agonist or a natural ligand, would ultimately lead to the modulation of several genes and corresponding proteins. Accordingly, PPARs are implicated in a wide range of physiological functions including enhancement of insulin sensitivity, improvement of glucose and lipid metabolism, suppression of inflammatory/oxidative insults, modulation of the immune response and the regulation of cell growth and differentiation (1-5, 29-38).

Since transcriptional PPAR responses are triggered by ligand-induced recruitment and/or the release of small accessory molecules which may be co-activators and co-repressors, PPARs can positively regulate target genes as observed in the potentiation of insulin signaling and lipid metabolism or repress the transcription of genes as seen during the suppression of inflammation (164-170). Interestingly, the modulation of PPAR activity by cofactors has been explored to design many pharmacological agents like agonists or ligands for PPAR alpha and PPAR gamma (71). In general, PPAR gamma agonists such as thiazolidinediones improve insulin resistance and glucose metabolism, while PPAR alpha agonists such as fibrates improve dyslipidemia (12-25, 171). The concomitant activation of receptors of PPAR alpha and PPAR gamma may yield a synergistic effect of greater physiological relevance in the homeostatic control of hyperglycemia and cardiac dysfunction associated with diabetes. Accordingly, an emerging trend in PPAR research is formulation of pan-PPAR ligands capable of simultaneously activating two or all three PPAR isoforms (7, 24, 25, 71, 172-175), with the hope of identifying more potent pharmacological response. Accordingly, several dual PPAR alpha/gamma agonists including aleglitazar, muraglitazar and tesagliptazar are currently under development at different phases of clinical trials (7, 25, 174, 175). The results of these clinical trials have yielded mixed results because of serious adverse effects such as bladder cancer and hyperplasia observed with ragaglitazar, MK-0767, and navagliptazar (176, 177),

posing further challenges for the formulation of agonist with better safety profile. The search for more-specific PPAR gamma and PPAR alpha agonists continues and candidate products such as INT131, MK0533 and ATx008-001/ FK614 for PPAR gamma (18, 178, 179), as well as other candidate compounds like derivatives of bis-oximinoalkanoic acid for PPAR alpha (180) are presently under development. However, many of such agonists are also noted for excessive side effects because the cofactors that modulate PPAR activity also have the intrinsic ability to regulate a wide spectrum of metabolic processes (71, 181, 182). Besides the development of dual PPAR alpha/gamma agonists, ambitious studies have been designed to formulate agonists capable of concomitantly stimulating all three PPAR isoforms (PPAR alpha, PPAR gamma and PPAR delta) (172, 173). Accordingly, the PPAR alpha/gamma/delta agonist, sipoglitazar is at phase II clinical trials for possible use against type-2 diabetes (172, 173). Whether these generations of new agonist would yield the desired pharmacological effects remains the subject of more intense investigations.

Although the therapeutic potential of PPAR agonists remains enormous, many challenges have to be overcome. Besides collateral effects, recent findings indicate that the different PPAR agonists may have opposing effects. For example, pioglitazone a PPAR gamma agonist, has been shown to improve cardiovascular function, while another PPAR gamma agonist, rosiglitazone, reportedly aggravated the risk of myocardial infarction in clinical trials (25). Similarly, pan-PPAR alpha/gamma agonists such as muraglitazar and tesagliptazar have also encountered setbacks in clinics because of excessive adverse effects (25), whereas another pan-PPAR alpha/gamma agonist such as aleglitazar showed promising results in phase-II clinical trials with benefits on glucose and lipid metabolism (25). These conflicting clinical observations from different PPAR gamma agonists constitute an important drawback that needs to be thoroughly investigated. To improve the benefit-to-risk ratio, it is imperative to formulate novel PPAR agonists with greater selectivity, enhanced efficacy, but less collateral effects. The quest for PPAR agonists with such rigorous prerequisites have led to the inception of several novel candidate compounds including the partial PPAR gamma agonists INT131, MK0533 and ATx008-001/ FK614 (18, 178, 179). Although these studies are still preliminary, the initial results are quite promising.

Given that diverse role of PPARs in regulating physiological functions and the existence of different isoforms (PPAR alpha, PPAR gamma and PPAR delta), the formulation of specific agonists and blockers of these isoforms are needed to assess the specific input and relative contribution of each isoform in regulating physiological functions. Whether the different tissue distribution of PPARs in different tissues (39-42) (43-47) would affect the relative effect in specific tissue needs further clarification because tissue-specific responses is a common phenomenon in physiological milieu (183, 184). Nevertheless, the formulation of specific triad PPAR gamma/alpha/delta agonists may be more effective because

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the synergism and/or additive effects of the single agonists for PPAR gamma, PPAR alpha, and PPAR delta. Since PPAR gamma agonists like thiazolidinediones improve insulin resistance and glucose metabolism, while PPAR alpha agonists such as fibrates improve dyslipidemia, and emerging evidence indicate that PPAR delta modulates energy balance, weight loss, skeletal muscle endurance, lipid metabolism and insulin sensitivity (12-25, 45, 46, 64-66, 171, 185), the formulation of selective triad PPAR gamma/alpha/delta agonists may confer greater benefits in patients co-morbid with insulin resistant diabetes, dyslipidemia, obesity and related cardiometabolic complications.

To further elucidate the role of the different PPAR in the regulation of physiological functions, each of the different isoforms, PPAR gamma, PPAR alpha, and PPAR delta will be elaborated in turn.

3.1. Peroxisome proliferator-activated receptor gamma (PPAR gamma)

Thiazolidinediones is the most common class of PPAR gamma agonists, with congenic such as troglitazone, rosiglitazone and pioglitazone (7, 25, 174, 175). Thiazolidinediones are well known for their role in enhancing insulin sensitivity and glucose metabolism by increasing insulin-stimulated glucose uptake in adipose tissue, skeletal muscle cells and hepatocytes, while decreasing hepatic gluconeogenesis (12-28). An interesting pharmacological characteristic of PPAR gamma agonists is their ability to evoke lasting glucose-lowering effect in type-2 diabetic patients without causing hypoglycaemia and gastrointestinal problems that commonly occur with the administration of other anti-diabetic agents like metformin and sulphonylureas (186). In spite of these positive attributes of PPAR gamma agonists, adverse effects associated with PPAR gamma such as bone fractures, macular oedema, heart failure, peripheral oedema and weight gain are amongst the setbacks that must be addressed (187-190). Thus the formulation of alternative PPAR gamma agonists with reduced collateral effects is imperative. A recent development in this line include the formulation of glitazars, an investigational class of dual-PPAR alpha/gamma agonists in phase-III clinical trials and have been shown to improve insulin sensitivity and glucose metabolism (7, 25, 174, 175). However, further research to fully characterize glitazars and more-importantly for the inception of more selective PPAR gamma agonist besides thiazolidinediones is needed.

Besides their ability to improve glucose metabolism, the activation of PPAR gamma is also beneficial in the cardiovascular system (28, 191-193). PPAR gamma agonists improve cardiovascular risk factors such as blood pressure, lipid metabolism, enhance adiponectin and mitigate inflammation peripherally (28, 191-197), as well as inflammation associated with chronic and acute neurological insults (29). Other relevant physiological effects of PPAR gamma include: (i) the suppression of fibrosis, (ii) reduction of deposition of extracellular matrix/profibrotic proteins such as fibronectin, collagen and transforming growth factor-beta in tissues,

(iii) the attenuation of apoptosis, (iv) the suppression of necrosis, and (v) the enhancement of vasorelaxation and thus the regulation of blood pressure (34-36, 147-150, 154-158). Thus, PPAR gamma is implicated in a wide range of physiological functions relevant to energy metabolism, tissue defense, the maintenance of intact structural morphology and blood pressure homeostasis. Although PPAR gamma abates hypertrophic growth mediated by extracellular matrix/profibrotic (34-36), emerging evidence indicate that PPAR gamma regulates the differentiation of trophoblasts, adipocytes and mesenchymal cells (48, 49, 198-200). In a related-study, PPAR gamma was shown to play an integral role in sustaining and optimizing trophoblast differentiation (48). In a similar way, cyclin G2, an unconventional cyclin that is generally upregulated during apoptosis or growth inhibition was shown to activate PPAR gamma during adipocyte differentiation (49). These studies suggest that PPAR gamma may be important for tissue turnover by selectively abating hypertrophic growth while promoting tissue regeneration and neogenesis.

The molecular mechanisms underlying the effects of PPAR gamma are not completely elucidated. Although the major effects of PPAR are largely mediated via the regulation of gene expression, emerging evidence indicate that PPAR gamma undergoes posttranslational modifications via SUMOylation to abate inflammation (201-204). Small Ubiquitin like-MODifier (SUMOylation) is a process of posttranslational modification for the regulation of proteins and cellular functions (201-203). However, further exploration of posttranslational modification by PPAR gamma is needed. Besides its involvement in inflammation novel studies are needed to clarify whether the mechanism of posttranslational modification is more diffused and implicated in other pathophysiological events modulated by PPAR gamma.

3.2. Peroxisome proliferator-activated receptor alpha (PPAR alpha)

PPAR alpha is the target receptor of the lipid-lowering class of drugs known as fibrates and related compounds including fenofibrate, gemfibrozil, bezafibrate, ciprofibrate, and clofibrate which were originally used as an adjunct therapy against hypercholesterolemia (24, 171). PPAR alpha regulates the transcription of many genes involved in lipid metabolism (51), and does so at intracellular and extracellular levels (51). At the intracellular level, PPAR alpha modulate hepatic fatty acid metabolism by enhancing fatty acid oxidation with increased catabolism and reduced synthesis of triglycerides (50, 51). In addition PPAR alpha enhances several important enzymes implicated in lipid metabolism namely: (i) acyl-CoA oxidase, a fundamental enzyme of peroxisomal β -oxidation; (ii) medium-chain acyl-CoA dehydrogenase, a key factor of mitochondrial β -oxidation; and (iii) cytochrome P450, a substance implicated in microsomal ω -hydroxylation of fatty acids (51-55). At extracellular level, PPAR alpha modulate lipid metabolism by increasing lipolysis of triglycerides (51, 205), an event that would eventually lead to the formation of HDL like apolipoproteins A-I and A-II (51, 56, 57).

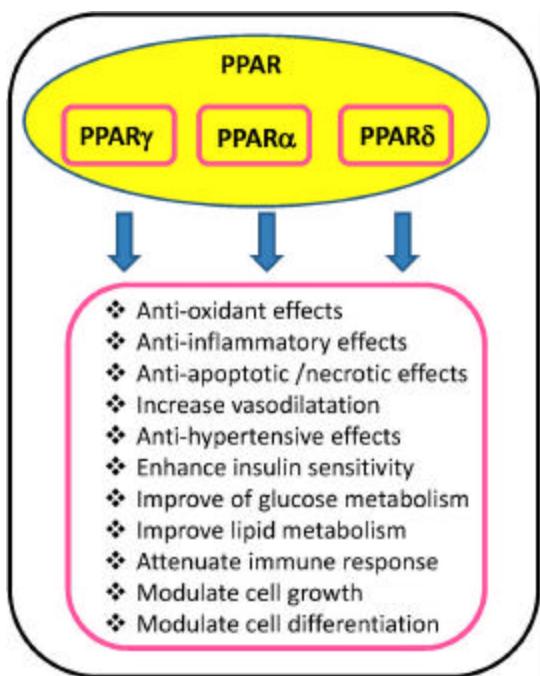


Figure 2. The Heme oxygenase (HO) system and its role in physiological functions. HO catalyzes heme breakdown to yield carbon monoxide, biliverdin, bilirubin and iron, while the iron formed enhances the synthesis of the antioxidant, ferritin. Carbon monoxide, bilirubin, biliverdin and ferritin are implicated in the potentiation of insulin sensitivity, improvement of glucose metabolism, enhancement of lipid metabolism, suppression of inflammation/oxidative stress, attenuation of immune response, reduction of apoptosis and necrosis, and modulation of cell growth and differentiation.

Besides its effects on lipid metabolism, agonists of PPAR alpha are known to suppress inflammation (58-62), modulate the immune response (37, 38), improve glucose metabolism (62, 152, 206), mitigate apoptosis/necrosis and relax vascular tissue (143-153), ameliorate nicotine-induced seizures (207-211) and neuroprotective in Parkinson's disease (212). Similarly, the PPAR alpha agonist, fenofibrate is cardioprotective and has been shown to inhibit left-ventricular hypertrophy, attenuate abnormalities in left-ventricular relaxation and improve systolic dysfunction in Dahl salt-sensitive hypertensive rats (213). Thus, the functions of PPAR alpha agonist not only to optimize lipid and glucose metabolism, but to attenuate inflammatory/immune responses, improve vasodilatation and regulate blood pressure, as well as conferring neuroprotection against seizures and Parkinson's disease. However PPAR alpha agonist have also been shown to induce apoptosis and cancer (214), suggesting the need for more in-depth studies to fully characterize the effects of PPAR alpha agonist in different tissues, especially when administered at different doses.

3.3. Peroxisome proliferator-activated receptor delta (PPAR delta)

PPAR delta (also known as PPAR beta) is widely distributed in tissues and modulate a wide spectrum of

physiological functions (43-47, 64-66). These include regulation of cell differentiation, vasodilation, energy balance, weight loss, skeletal muscle endurance and insulin sensitivity (43-47, 63-66, 159-161). Moreover, agonist of PPAR delta such as tesaglitazar, muraglitazar, ragaglitazar, imiglitazar, aleglitazar have been shown to reduce elevated blood glucose levels by altering the body's energy substrate preference from glucose to lipids (24). In addition, PPAR delta suppress inflammation, abates apoptosis, reduce remodeling of vascular tissue and is protective against lipotoxicity (44, 46, 47, 67, 68, 162, 163). These wide spectrum of effects of PPAR delta suggest a in both cellular physiology and pathophysiology effects.

In contrast to agonists of PPAR alpha and PPAR gamma that are used clinically, agonists of PPAR delta are still undergoing development and are at different phases of clinical trials (215-217). Although cytoprotection by PPAR delta (44, 46, 47, 67, 68, 162, 163) and anti-tumorigenic effects of PPAR delta have been reported (218), some reports in literature are not in accordance with this notion. These suggest a proapoptotic role of PPAR delta and its involvement in carcinogenesis (219, 220). These conflicting observations can only be clarified by more intense investigations in this area.

4. THE HEME OXYGENASE (HO) SYSTEM

In the human body, carbon monoxide is produced at a rate of 16.4 micromole per hour and daily production can reach 500 mM (89, 90, 221). About 86% comes from HO-catalyzed degradation of heme while 14% from lipid peroxidation, xenobiotics and other sources (221, 222). HO is a microsomal enzyme that degrades the pro-oxidant heme, generating cytoprotective products including biliverdin, bilirubin and carbon monoxide (Figure 2) (89, 137-139).

Although HO has three isoforms HO-1 (inducible), HO-2 and HO-3 (constitutive), the enzymatic activity is mainly derived from HO-1 and HO-2 because HO-3, has no functional genes in rat and is considered a pseudo-transcript of HO-2 (140-142). In physiological systems, the basal HO activity is maintained by HO-2 (90, 223-226), while HO-1 is activated by physical, chemical and pathophysiological stimuli (90, 226-230).

As such, HO-1 may be considered a sensitive index that is triggered during the onset of pathophysiological changes in cells as an attempt to counteract adverse conditions. However, the pathophysiological activation of HO-1 evokes only a transient or sub-threshold value of HO-activity that is incapable of activating the downstream signaling components of the HO system like the cyclic guanosine monophosphate (cGMP) (90, 99, 100, 231-234), suggesting the need for a robust enhancement of HO-1 by pharmacological agents like hemin, heme arginate, stannous mesoporphyrin, copper protoporphyrin and cobalt protoporphyrin (99, 100, 231-234). Therefore the transient up-regulation of HO-1 that accompanies many pathophysiological conditions may represent the first line of defense mounted by the HO system against tissue insults.

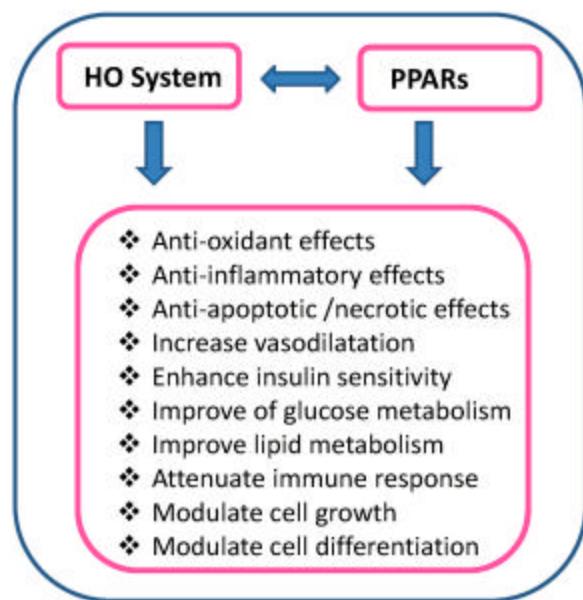


Figure 3. Crosstalk between the HO system and PPARs in regulating physiological functions. There is an intimate reciprocal relationship between PPARs and the HO system. Both PPARs and the HO system implicated in a wide range of physiological functions including enhancement of insulin sensitivity, improvement of glucose and lipid metabolism, suppression of inflammatory/oxidative insults, modulation of the immune response and the regulation of cell growth and differentiation.

The HO system and related products such as carbon monoxide, bilirubin, biliverdin and ferritin are of great physiological relevance (89-136). Accordingly, the HO system have been shown to normalize blood pressure in hypertensive animals, enhance insulin sensitivity, improve glucose/lipid metabolism, suppress inflammation/oxidative stress, abate immune response, attenuate apoptosis, and modulate cell growth and differentiation (Figure 2) (89-139, 235-257).

5. CROSSTALK BETWEEN THE HO SYSTEM AND PPARs

Mounting evidence indicates that there is an intimate, reciprocal, stimulatory and synergistic relationship between the HO system and PPARs-signaling (Figure 3) (4, 5, 76-88, 149, 258-274).

Accordingly, PPAR alpha, PPAR gamma and PPAR delta have been shown to upregulate the HO system (4, 5, 76, 78, 82, 84, 149, 268, 270-274). Conversely, the HO system potentiates PPAR alpha and PPAR gamma (81-86, 269, 275). In related studies, PPAR delta and PPAR gamma reportedly induced HO-1 to suppress oxidative stress (4, 78), while PPAR gamma and PPAR delta enhanced HO-1 to attenuate myocardial inflammation and fibrosis with corresponding reduction of myocardial infarction (149, 272). Similarly, PPAR-mediated repression of inflammation and nephrotoxicity was reportedly mitigated via induction of HO-1 (79-81). Furthermore, PPAR delta

has been shown to attenuate oxidative stress-induced premature cellular senescence by activating the HO system (271). Interestingly, a product generated by the HO system such as carbon monoxide has also been shown to modulate the expression of PPAR (259, 276). Accordingly, carbon monoxide mitigated inflammation by activating PPAR delta (258). Similarly, carbon monoxide reportedly preserved adequate PPAR gamma levels and suppressed inflammatory insults via PPAR gamma-mediated transcriptional repression of pro-inflammatory genes and proteins (259-265, 276). Furthermore, PPAR gamma and the HO system were shown to evoke vascular relaxation by a common mechanism involving potassium channels (155, 277-280). In related studies PPAR gamma and the HO system triggered vasorelaxation by activating ATP-potassium channels and calcium-activated potassium channels (155, 277-280). Collectively, these studies underscore the multifaceted interaction between the HO system and PPARs in combating tissue insults, potentiating insulin signaling, enhancing lipid/glucose metabolism, modulating vascular tone and relaxation, and thus the maintenance of optimal cellular function and homeostasis.

Although there is compelling evidence supporting the reciprocal, stimulatory and synergistic relationship between the HO system and PPARs-signaling (Figure 3) (4, 5, 76-88, 149, 258-274), further studies are still needed to fully characterize the close interaction between the HO system and PPARs. The interaction may be multifaceted and complex. However, emerging evidence indicates that the HO-1 gene promoter contains motifs for PPAR response element (82, 84, 85). Accordingly, PPAR alpha and PPAR gamma have been shown to transcriptionally induce HO-1 by binding to PPAR responsive elements in HO-1 gene (76, 77). Given that dual-PPAR alpha/gamma agonists (7, 25, 174, 175) and PPAR delta agonist (24) are all known to improve glucose metabolism, and HO-1 gene promoter has motifs for PPAR response element (82, 84, 85), it is possible that the mechanism through the HO system improves glucose/lipid metabolism (89, 91-96, 120, 237, 240, 245, 281) is via the modulation of PPAR agonists. Alternatively, the PPAR response element in HO-1 gene (82, 84, 85) could activate the glucocorticoid-responsive-element in HO-1 gene-promoter (282) to enhance insulin signaling and improve glucose metabolism given that glucocorticoids regulate glucose metabolism and insulin resistance (283). Nevertheless, the interaction between the HO system and PPARs is complex and multifaceted, and the presence of PPAR responsive elements in HO-1 gene (76, 77) may just be one of the facets to a more complex puzzle. Thus, these initial findings may simply represent the tip of the iceberg that should be explored further given the convergence of a wide spectrum physiological functions including the enhancement of insulin sensitivity, improvement of glucose and lipid metabolism, suppression of inflammatory/oxidative insults, attenuation of immune response and the modulation of cell growth and differentiation by PPARs (1-5, 11, 29-38) and the HO system (89-136) (Figure 3).

6. CONCLUSION

The activation of PPARs triggers specific systematic responses through gene regulation. Interestingly,

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PPARs can induce or repress the expression of target genes to evoke responses which are channeled to enhance and improve insulin sensitivity, improve glucose/lipid metabolism, but suppress inflammation/oxidative stress, abate the immune response and regulate cell-growth and differentiation (1-5, 29-38). Incidentally, PPAR-signaling is implicated in many effects of the HO system (76-88), and PPAR is potentiated by the HO system (81, 82, 84, 85). Thus, a possible cross-talk between PPARs and the HO system can be envisaged. The existence of an intimate reciprocal relationship between PPAR and the HO system has been further sustained by the presence of motifs of PPAR in HO-1 gene promoter (82, 84, 85). Interestingly, many of the reported physiological effects of PPARs are similar to those evoked by the HO system. Accordingly, the HO system and related products including carbon monoxide, bilirubin, biliverdin and ferritin enhance insulin sensitivity, improve glucose/lipid metabolism, abate apoptosis, inflammation/oxidative stress and immune response and modulate cell growth and differentiation (89-136). Thus, the interaction between the HO system and PPARs may constitute the basis of many beneficial effects against insulin resistant diabetes, hypertension, obesity and related cardiometabolic complications.

Therefore, the coordinated activity between HO system and PPARs is important for fine tuning physiological functions, the maintenance of cellular homeostasis, and more-importantly may constitute the basis for future drug design.

7. ACKNOWLEDGEMENTS

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