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- (54) Title: MAMMALIAN PEROXISOME PROLIFERATOR-ACTIVATED RECEPTORS AND USES THEREOF
- (57) Abstract

Peroxisome proliferator-activated receptor subunits designated PPAR γ and PPAR δ are described. Nucleic acid sequences encoding the receptor subunits, expression vectors containing such sequences and host cells transformed with such vectors are also disclosed, as are heterodimeric PPAR receptors comprising at least one of the invention subunits, and methods for the expression of such novel receptors, and various uses therefor.

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MAMMALIAN PEROXISOME PROLIFERATOR-ACTIVATED RECEPTORS AND USES THEREOF

RELATED APPLICATIONS

This application is a continuation-in-part of United States Application Serial Number 08/270,643, now pending, which is a continuation-in-part of United States Application Serial Number 07/907,908, filed July 2, 1992, now abandoned, which is a continuation-in-part of United States Application Serial Number 07/497,935, filed March 22, 1990, now abandoned.

FIELD OF INVENTION

The present invention relates to novel members of the steroid/thyroid superfamily of receptors, as well as uses therefor.

15 BACKGROUND OF THE INVENTION

Peroxisome proliferators are a structurally diverse group of compounds which, when administered to rodents, elicit dramatic increases in the size and number of hepatic and renal peroxisomes, as well as concomitant 20 increases in the capacity of peroxisomes to metabolize fatty acids via increased expression of the enzymes required for the B-oxidation cycle (Lazarow and Fujiki, Ann. Rev. Cell Biol. 1:489-530 (1985); Vamecq and Draye, Essays Biochem. 24:1115-225 (1989); and Nelali et al., 25 Cancer Res. 48:5316-5324 (1988)). Chemicals included in this group are the fibrate class of hypolipidermic drugs, herbicides, and phthalate plasticizers (Reddy and Lalwani, Rev. Toxicol. 12:1-58 (1983)). Peroxisome proliferation can also be elicited by 30 physiological factors such as a high-fat diet and cold acclimatization.

Insight into the mechanism whereby peroxisome proliferators exert their pleiotropic effects was provided by the identification of a member of the nuclear hormone receptor superfamily activated by these chemicals (Isseman 5 and Green, Nature 347-645-650 (1990)). This receptor, termed peroxisome proliferator activated receptor alpha $(PPAR\alpha)$, was subsequently shown to be activated by a variety of medium and long-chain fatty acids and to stimulate expression of the genes encoding rat acyl-CoA 10 oxidase and hydratase-dehydrogenase (enzymes required for peroxisomal \(\beta \)-oxidation), as well as rabbit cytochrome P450 4A6, a fatty acid ω -hydroxylase (Gottlicher et al., Proc.Natl. Acad. Sci. USA 89:4653-4657 (1992); Tugwood et al., EMBO J. 11:433-439 (1992); Bardot et al., Biochem. Biophys. 15 Res. Comm. 192:37-45 (1993); Muerhoff et al., J. Biol. Chem. 267:19051-19053 (1992); and Marcus et al., Proc. Natl. Acad. Sci. USA 90(12):5723-5727 (1993). foregoing references support a physiological role for PPAR α in the regulation of lipid metabolism. PPARα activates 20 transcription by binding to DNA sequence elements, termed peroxisome proliferator response elements (PPRE), as a heterodimer with the retinoid X receptor. The retinoid X receptor is activated by 9-cis retinoic acid (see Kliewer et al., Nature 358:771-774 (1992), Gearing et al., Proc. 25 Natl. Acad. Sci. USA 90:1440-1444 (1993), Keller et al., Proc. Natl. Acad. Sci. USA 90:2160-2164 (1993), Heyman et al., Cell 68:397-406 (1992), and Levin et al., Nature 355:359-361 (1992)). Since the PPAR α -RXR complex can be activated by peroxisome proliferators and/or 9-cis retinoic 30 acid, the retinoid and fatty acid signaling pathways are seen to converge in modulating lipid metabolism.

BRIEF DESCRIPTION OF THE INVENTION

In accordance with the present invention, there are provided isolated mammalian peroxisome proliferator-

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activated receptor subunit proteins of the γ and δ subtypes, and functional fragments thereof. In addition, there are provided isolated nucleic acids encoding mammalian peroxisome proliferator-activated receptor subunit proteins, as well as fragments thereof. There are also provided vectors containing the above-described nucleic acids, as well as cells containing such nucleic acids and/or vectors.

The present invention also provides methods for the recombinant production of mammalian peroxisome proliferator-activated receptor proteins comprising at least one PPAR subunit protein of the γ and δ subtype, and functional fragments thereof, as well as methods to identify clones encoding the above-described receptor subunit proteins, and functional fragments thereof.

Also provided by the present invention are methods for screening compounds to determine those which bind to mammalian peroxisome proliferator-activated receptor proteins comprising at least one PPAR subunit 20 protein of the γ or δ subtype, or functional fragments thereof, as well as bioassays for evaluating whether test compounds are agonists or antagonists for receptor proteins of the invention, or functional modified forms of said receptor protein(s).

25 <u>BRIEF DESCRIPTION OF THE FIGURES</u>

Figure 1 presents a schematic comparison of the members of the PPAR gene family using mPPAR δ as a reference. Comparisons among the different domains of the proteins are expressed as percent amino acid identity.

Figure 2 demonstrates that PPARy and PPAR δ fail to respond to the peroxisome proliferator Wy 14,643. CV-1 cells were cotransfected with reporter plasmid PPRE₃-TK-LUC

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and either no receptor expression plasmid (-), CMX-PPAR α , CMX-PPAR γ , or CMX-PPAR δ and then incubated in either the absence (-) or presence (+) of 5 μ M Wy 14,643. Luciferase activities are expressed as percentages of the maximal response where 100% is the activity obtained with PPAR α in the presence of 5 μ M Wy 14,643.

Figure 3 illustrates the ability of PPARy and PPARδ to repress PPARα-mediated responsiveness to Wy 14,643. CV-1 cells were cotransfected with reporter plasmid PPRE3-TK-LUC and either no receptor expression plasmid (NONE) or CMX-PPARα (10ng) in either the absence or presence of CMX-PPARγ (100ng) or CMX-PPARδ (100ng). Cells were then incubated in either the absence (-) or presence (+) of 5μM Wy 14,643. Luciferase activities are presented as fold-activation relative to cells which were not transfected with receptor expression plasmid and were not treated with Wy 14,643.

Figure 4 demonstrates that PPAR isoforms are pharmacologically distinct. CV-1 cells were cotransfected with reporter plasmid PPRE₃-TK-LUC and either no receptor expression plasmid (-), CMX-PPARα, CMX-PPARγ, or CMX-PPARδ in either the absence or presence of 5μM Wy 14,643 (WY), 30μM linoleic acid (C18:2), or 30μM LY-171883 (LY). Luciferase activities are presented as the fold activation achieved in compound-treated versus mock-treated cells. Similar results were obtained in triplicate in three independent experiments.

DETAILED DESCRIPTION OF THE INVENTION

Two novel PPAR receptor subunits have been cloned and characterized. These novel γ and δ isoforms (subunits) together with the α subunit display marked differences in their responsiveness to peroxisome proliferators and fatty acids, as well as differences in their temporal and spatial

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patterns of expression. These observations suggest a broad role for the PPAR family during development and in adult physiology.

The existence of multiple PPAR isoforms with distinct expression patterns has been found to correlate with the fact that the three isoforms have different ligand specificities. Indeed, the PPAR isoforms are shown herein to be pharmacologically distinct. Thus, PPARα, PPARγ and PPARδ are most efficiently activated by Wy 14,643, 10 LY-171883, and linoleic acid, respectively. Remarkably, Wy 14,643, which results in approximately 100-fold induction in reporter expression in the presence of PPARα, fails to activate either PPARγ or PPARδ.

With regard to this differential responsiveness 15 to activators of peroxisome proliferation, the relationship among the PPAR isoforms may be analogous to that between the glucocorticoid and mineralocorticoid receptors (GR and MR, respectively). While both receptors can bind to the same response element, and both respond 20 mineralocorticoids and corticosteroids, MR and GR display differential sensitivities to aldosterone and specific glucocorticoids such as dexamethasone, respectively (Arriza et al., Neuron 1:887-900 (1988)). Thus, the ratio of these receptors to their ligands provides a means of determining 25 tissue-specific expression of target genes. Similarly, the existence of multiple PPAR isoforms with overlapping ligand specificities may provide the means for tissue-specific regulation of gene expression by peroxisome proliferators and fatty acids.

In addition to their differential responsiveness to peroxisome proliferators, the three PPAR isoforms also display distinct yet overlapping expression patterns. As previously shown, PPARα mRNA is abundant in liver and kidney (Isseman and Green, supra; Beck et al., Proc. R.

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Soc. Lond. 247:83-87 (1992)), tissues in which peroxisome proliferators result in dramatic increases in the numbers of peroxisomes and concomitant increases in peroxisomal B-oxidation (Nemali et al., supra). In contrast, the 5 levels of PPARy mRNA and PPARδ mRNA, which can act as dominant repressors of PPAR α -mediated responsiveness to Wy 14,643, are low in these tissues. Thus, a pattern emerges in which tissues that are most responsive to peroxisome proliferators such as Wy 14,643 are observed to express high amounts of PPAR α mRNA and relatively low amounts of PPARy mRNA and PPAR& mRNA. These data suggest that the ratio of the PPAR isoforms is likely to play a critical role in establishing the degree of responsiveness of tissues to specific peroxisome proliferators.

15 Widespread expression of PPARS is observed in both the embryo and in adult tissues. This observation suggests that this isoform may play а general "housekeeping" role. In contrast, PPARy is observed to be expressed almost exclusively in the adrenal and spleen. 20 The expression of all three PPAR isoforms in the adrenal is particularly intriguing, since diseases which result in peroxisome dysfunction (e.g. adrenoleukodystrophy and Zellweger syndrome) cause gross morphological changes in adrenal cells and, eventually, adrenal deficiency. 25 observations suggest a critical role for peroxisomes in this tissue (Vamecq and Draye, supra). Interestingly, peroxisomes can be induced to proliferate in hamster adrenals in response to treatment with adrenocorticotropic hormone and corticosteroids (Black and Russo, Amer. J. 30 Anatomy 159:85-120 (1980)), indicating the presence of adrenal-specific signaling pathway(s) for peroxisome proliferation. The differential expression of PPARy in the adrenal suggests that this isoform may respond to an adrenal-enriched ligand.

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Accordingly, in accordance with the present invention, there are provided isolated mammalian peroxisome proliferator-activated receptor subunit proteins of the γ or δ subtype and functional fragments thereof.

5 As employed herein, the phrase "mammalian peroxisome proliferator-activated receptor subunit proteins of the γ or δ subtype" refers to isolated and substantially purified as well as recombinantly produced proteins which members of the steroid/thyroid superfamily 10 receptors, and which mediate the pleiotropic effects of peroxisome proliferators (such as medium and long-chain fatty acids). Such receptors participate in the formation of heterodimeric species with retinoid X receptors (RXRs) and comprise an amino-terminal domain, a DNA binding 15 domain, and a ligand binding domain. Also contemplated within this definition are variants thereof encoded by mRNA generated by alternative splicing of a primary transcript.

Use of the terms "recombinantly produced", "isolated" "substantially pure" or in the 20 specification and claims as a modifier of DNA, polypeptides or proteins means that the modified substances have been produced by the hand of man, and thus are separated from their native in vivo cellular environment. As a result of this human intervention, the recombinant/ isolated/substantially pure DNAs, RNAs, polypeptides and 25 proteins of the invention are useful in ways that the naturally occurring DNAs, RNAs, polypeptides or proteins are not, for example, in assays to identify selective drugs or compounds.

The novel receptors of the present invention also can be included as part of a panel of receptors which are screened to determine the selectivity of interaction of proposed agonists or antagonists of other steroid hormone

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receptors. Thus, a compound which is believed to interact selectively, for example, with the glucocorticoid receptor, should not have any substantial effect on any other receptors, including invention receptors. However, if such 5 a proposed compound does interact with the invention receptors, then the probability of side effects caused by the activation of other receptors in addition to the target receptor, is clearly indicated. For example, the use of many drugs in the treatment of hormone-related disorders is 10 currently restricted by side effects caused by activation of "non-target" receptors. Employment of the invention receptors in a panel of receptors in a screen to determine the selectivity of interaction of potential ligands provides a means to identify receptor-specific ligands that are therapeutically superior than currently used ligands that cause unwanted side effects.

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As used herein, the term splice variant refers to variant encoding nucleic acid(s) produced PPAR differential processing of primary transcript(s) of genomic 20 DNA, resulting in the production of more than one mRNA. derived from differentially processed primary transcript will encode PPAR receptor proteins that have regions of complete amino acid identity and regions having different amino acid sequences. Thus, the same genomic 25 sequence can lead to the production of multiple, related mRNAs and corresponding proteins. Both the resulting mRNAs and proteins are referred to herein as "splice variants".

Accordingly, also contemplated within the scope of the present invention are nucleic acids that encode 30 mammalian PPAR receptor subunit proteins as defined above, but that by virtue a degenerate genetic code do not necessarily hybridize to the nucleic acids set forth in SEQ ID NOs: 1 or 3 under specific hybridization conditions. Nucleic acid fragments encoding invention receptor subunit 35 proteins are capable of forming a functional heterodimer

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with one or more RXR receptor protein isoform(s).

Typically, unless a PPAR receptor protein is encoded by mRNA that arises from alternative splicing (i.e., a splice variant), PPAR receptor encoding DNA and encoded protein share substantial sequence homology with at least one of the PPAR receptor-encoding DNAs and encoded proteins described herein. It is understood that DNA or RNA encoding a splice variant may share less than 90% overall sequence homology with the DNA or RNA provided herein, but include regions of nearly 100% homology to a DNA fragment described herein, and encode an open reading frame that includes start and stop codons and encodes a functional PPAR receptor protein.

Exemplary nucleic acid sequences encoding
15 mammalian peroxisome proliferator-activated receptor subunit proteins of the y subtype are represented by nucleotide sequences which encode substantially the same amino acid sequence as set forth in SEQ ID NO:2. Presently preferred sequences encode the same amino acid sequence as set forth in SEQ ID NO:2.

Exemplary nucleic acid sequences can alternatively be characterized as those nucleotide sequences which encode mammalian peroxisome proliferatoractivated receptor subunit proteins of the y subtype and hybridize under high stringency conditions to SEQ ID NO:1.

Exemplary nucleic acid sequences encoding mammalian peroxisome proliferator-activated receptor subunit proteins of the δ subtype are represented by nucleotides which encode substantially the same amino acid sequence as set forth in SEQ ID NO:4. Presently preferred sequences encode the same amino acid sequence as set forth in SEQ ID NO:4.

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Especially preferred sequences are those which have substantially the same nucleotide sequence as that set forth in SEQ ID NO:1.

Exemplary nucleic acid sequences can alternatively be characterized as those nucleotide sequences which encode mammalian peroxisome proliferatoractivated receptor subunit proteins of the δ subtype and hybridize under high stringency conditions to SEQ ID NO:3.

Especially preferred nucleic acid sequences are those which have substantially the same nucleotide sequence as the coding sequences in SEQ ID NO:3.

The phrase "stringency of hybridization" is used herein to refer to conditions under which polynucleic acid hybrids are stable. As known to those of skill in the art, the stability is reflected in the melting temperature (T_m) of the hybrids. T_m can be approximated by the formula:

$$81.5^{\circ}C - 16.6(\log_{10}[Na^{\dagger}]) + 0.41(\%G+C) - 600/1,$$

where l is the length of the hybrid in number nucleotides. $\rm T_{m}$ decreases approximately 1-1.5°C with every 1% decrease in sequence homology. In general, stability of a hybrid is a function of sodium ion concentration and temperature. Typically, the hybridization reaction initially performed under is conditions of low stringency, followed by washes 25 varying, but higher, stringency. Reference to hybridization stringency relates to such washing conditions. Thus, as used herein:

> (1) HIGH STRINGENCY refers to conditions that permit hybridization of only those nucleic acid sequences that form stable hybrids in 0.018M NaCl at 65°C (i.e., if a hybrid is

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not stable in 0.018M NaCl at 65°C, it will not be stable under high stringency conditions, as contemplated herein). High stringency conditions can be provided, for example, by hybridization in 50% formamide, 5X Denhart's solution, 5X SSPE, 0.2% SDS at 42°C, followed by washing in 0.1X SSPE, and 0.1% SDS at 65°C;

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(2) MODERATE STRINGENCY refers to conditions that permit hybridization in 50% formamide, 5X Denhart's solution, 5X SSPE, 0.2% SDS at 42°C, followed by washing in 0.2X SSPE, 0.2% SDS, at 65°C; and

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(3) LOW STRINGENCY refers to conditions that permit hybridization in 10% formamide, 5X Denhart's solution, 6X SSPE, 0.2% SDS at 42°C, followed by washing in 1X SSPE, 0.2% SDS, at 50°C.

It is understood that these conditions may be varied using a variety of buffers and temperatures well known to skilled artisans.

As used herein, the phrase "substantial sequence homology" refers to nucleotide sequences which share at least about 90% identity, and amino acid sequences which typically share more than 95% amino acid identity. It is recognized, however, that proteins (and DNA or mRNA encoding such proteins) containing less than the above-described level of homology arising as splice variants or that are modified by conservative amino acid substitutions (or substitution of degenerate codons) are contemplated to be within the scope of the present invention.

As used herein, the phrase "substantially the same" refers to nucleotide sequences, ribonucleotide sequences, or amino acid sequences, that have slight and

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non-consequential sequence variations from the actual sequences disclosed herein. Species "substantially the same" are considered to be equivalent to the disclosed sequences, and as such are within the scope 5 of the appended claims. In this regard, "slight and nonconsequential sequence variations" mean that sequences that are substantially the same as invention sequences disclosed and claimed herein, are functionally equivalent to the sequences disclosed and claimed herein. Functionally 10 equivalent sequences will function in substantially the same manner to produce substantially the same results as the nucleic acid and amino acid sequences disclosed and claimed herein. Specifically, functionally equivalent nucleic acids encode proteins that have conservative amino 15 acid variations, such as substitution of a non-polar residue for another non-polar residue or a charged residue for a similarly charged residue. These changes are recognized by those of skill in the art as modifications that do not substantially alter the tertiary structure of 20 the protein.

Fragments of invention nucleic acid sequences are useful as hybridization probes, wherein such fragments comprise at least 14 contiguous nucleotides of the above-described nucleic acids, and wherein the fragment is labeled with a detectable substituent. Suitable detectable substituents can be readily determined by those of skill in the art, and include such species as radiolabeled molecules, fluorescent molecules, enzymes, ligands, and the like.

As used herein, a probe is single- or doublestranded DNA or RNA that has a sequence of nucleotides that includes at least 14 contiguous bases that are the same as (or the complement of) any 14 or more contiguous bases set forth in SEQ ID NOs:1 or 3. Preferred regions for the 35 construction of probes include those regions predicted to

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encode a DNA binding domain. Such regions are preferred because they are most highly conserved among members of the steroid/thyroid superfamily of receptors.

As a particular application of the invention sequences, genetic screening can be carried out using the nucleic acid sequences of the invention as probes. Thus, nucleic acid samples from patients having conditions suspected of involving alteration/modification of any one or more of the PPAR receptor subtypes can be screened with appropriate probes to determine if abnormalities exist with respect to the endogenous PPAR receptor proteins.

In accordance with yet another embodiment of the present invention, there are provided vectors comprising nucleic acid sequences, as well as cells and vectors 15 containing such sequences. Such host cells, including bacterial, yeast and mammalian cells can be used for expressing invention nucleic acids to produce PPAR receptor Incorporation of cloned DNA into a suitable expression vector, transfection of eukaryotic cells with a 20 plasmid vector or a combination of plasmid vectors, each encoding one or more distinct genes, and selection of transfected cells are well known in the art (see, e.g., Sambrook et al. (1989) Molecular Cloning: A Laboratory Manual, Second Edition, Cold Spring Harbor Laboratory 25 Press). Heterologous DNA may be introduced into host cells by any method known to those of skill in the art, such as transfection by CaPO4 precipitation with a vector encoding the heterologous DNA (see, e.g., Wigler et al. (1979) Proc. Natl. Acad. Sci. <u>76</u>:1373-1376), DEAE-dextran, 30 electroporation, microinjection, or lipofectamine (GIBCO BRL #18324-012). Transfected host cells can then be cultured under conditions whereby the receptor subunit protein(s) encoded by the DNA is (are) recombinantly expressed.

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The present invention further provides a mammalian peroxisome proliferator-activated receptor, expressed recombinantly in a host cell. The receptor comprises at least one PPAR subunit, wherein the PPAR subunit is PPARγ or PPARδ, and at least one retinoid X receptor isoform. The invention receptor has the ability to repress PPARα-mediated responses activated by Wy 14,643,

Also provided by the present invention are mammalian peroxisome proliferator-activated subunit 10 proteins, expressed recombinantly in a host cell wherein the receptor subunits have substantially the same amino acid sequence as set forth in SEQ ID NOs: 2 or 4.

In accordance with still another embodiment of the present invention, there is provided a method for the recombinant production of mammalian peroxisome proliferator-activated receptor proteins comprising at least one PPAR subunit of the γ or δ subtype, or functional fragments thereof. Such method comprises expressing the above-described nucleic acid(s) in a suitable host cell.

In accordance with still another embodiment of the present invention, there is provided a method to identify clones encoding mammalian peroxisome proliferator-activated receptor subunit proteins of the γ or δ subtype, or functional fragments thereof. Such method comprises screening a genomic or cDNA library with an invention nucleic acid probe under low stringency hybridization conditions, and identifying those clones which display a substantial degree of hybridization to said fragment.

Nucleic acids encoding mammalian peroxisome proliferator-activated receptor subunit protein of the γ or δ subtype, or functional fragments thereof may be isolated by screening suitable human cDNA or human genomic libraries under suitable hybridization conditions with nucleic acids

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disclosed herein (including nucleotide sequences derived from SEQ ID NOs:1 or 3). Suitable libraries can be prepared from appropriate tissue samples, e.g., brain tissue, heart tissue, intestinal tissue, kidney tissue, liver tissue, spleen tissue, and the like. The library can be screened with nucleic acid including substantially the entire receptor-encoding sequence thereof, or the library may be screened with a suitable probe, as described above.

After screening the library, positive clones are 10 identified by means of a hybridization signal; identified clones are characterized by restriction enzyme mapping and/or DNA sequence analysis, and then examined, by comparison with the sequences set forth herein to ascertain whether they encode a complete PPAR receptor subunit protein (i.e., if they include translation initiation and termination codons). If the selected clones incomplete, they may be used to rescreen the same or a different library to obtain overlapping clones. library is genomic, then the overlapping clones may include exons and introns. If the library is a cDNA library, then 20 the overlapping clones will include an open reading frame. In both instances, complete clones may be identified by comparison with the DNA and encoded proteins provided herein.

The ligand-binding domain (LBD) of nuclear hormone receptors is a complex multifunctional unit containing subdomains for dimerization, transcriptional suppression and hormone-induced transactivation (Forman and Samuels, Mol. Endocrinol. 4:1293-1301 (1990)). The dimerization domain includes a series of heptad repeats flanked by sequences required for ligand binding. Thus, the dimerization domain is embedded within the larger LBD. This structural arrangement raises the possibility that dimerization may serve as an allosteric modulator of ligand binding and transactivation. It has previously been shown

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that the *Drosophila* ecdysone receptor (EcR) acquires ligand binding activity after heterodimerization with USP (*Drosophila* homolog of RXR; see Yao et al., in *Nature* 366:476-479 (1993)). Thus, differential interactions among receptor LBDs can either restrict, redirect or lead to an acquisition of new ligand binding phenotypes.

It has recently been shown that PPARα binds to its cognate response elements as a heterodimer with the RXR (see Kliewer et al., supra, Gearing et al., supra, or Keller et al., supra). The resulting PPARα-RXR complex can respond to both peroxisome proliferators and 9-cis retinoic acid (see Kliewer et al., (1992), supra). It has now been found that PPARγ and PPARδ also cooperate with RXR in the formation of heterodimers, and in binding to DNA as heterodimers. Ultimately, the regulation of peroxisome physiology is likely a consequence of a complex interplay among the multiple PPAR and RXR isoforms and the ligands for these receptors.

In accordance with the present invention, there
20 are provided combinations of receptors comprising at least
two different members of the steroid/thyroid superfamily of
receptors, wherein one receptor is either PPARy or PPARδ,
and wherein said receptors are associated in the form of a
multimer, preferably a heterodimer. A particularly
25 preferred combination of receptors is a heterodimer
comprising either PPARy or PPARδ and a subtype of RXR.

Combinations contemplated by the present invention can broadly be referred to as "multimeric species," which is intended to embrace all of the various oligomeric forms in which members of the steroid/thyroid superfamily of receptors (including fragments thereof comprising the dimerization domains thereof) are capable of associating in combination with either PPARy or PPARS. Thus, reference to "combinations" of steroid receptors or

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"multimeric" forms of steroid receptors homodimeric combinations of a single PPARy or PPARS receptor (including fragments thereof comprising the dimerization domains thereof), heterodimeric combinations 5 of either a PPARy or PPARδ receptor and another different receptor (including fragments thereof comprising the dimerization domains thereof), homotrimeric combinations of a single PPARy or PPARS receptor (including fragments thereof comprising the dimerization domains thereof), 10 heterotrimeric combinations of two or three different receptors including PPAR γ or PPAR δ (including fragments thereof comprising the dimerization domains thereof), homotetrameric combinations of a single PPAR γ or PPAR δ receptor (including fragments thereof comprising the 15 dimerization domains thereof), heterotetrameric combinations of two or more different receptors including PPARy or PPAR& (including fragments thereof comprising the dimerization domains thereof), and the like.

As employed herein, the phrase "members of the steroid/thyroid superfamily of receptors" (also known as "nuclear receptors" or "intracellular receptors") refers to hormone binding proteins that operate as ligand-dependent transcription factors, including identified members of the steroid/thyroid superfamily of receptors for which specific ligands have not yet been identified (referred to hereinafter as "orphan receptors"). These hormone binding proteins have the intrinsic ability to bind to specific DNA sequences. Following binding, the transcriptional activity of target gene (i.e., a gene associated with the specific DNA sequence) is modulated as a function of the ligand bound to the receptor.

The DNA-binding domains of all of these nuclear receptors are related, consisting of 66-68 amino acid residues, and possessing about 20 invariant amino acid residues, including nine cysteines. A member of the

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superfamily can be identified as a protein which contains the above-mentioned invariant amino acid residues, which are part of the DNA-binding domain of such known steroid receptors as the human glucocorticoid receptor (amino acids 421-486), the estrogen receptor (amino acids 185-250), the mineralocorticoid receptor (amino acids 603-668), the human retinoic acid receptor (amino acids 88-153). The highly conserved amino acids of the DNA-binding domain of members of the superfamily are well-known as set forth, for example in PCT WO 94/01558. Thus, the DNA-binding domain is a minimum of 66 amino acids in length, but can contain several additional residues.

Exemplary members of the steroid/thyroid superfamily of receptors contemplated for use in the practice of the present invention (including the various 15 isoforms thereof) include steroid receptors such as mineralocorticoid receptor, progesterone receptor, androgen receptor, vitamin $\mathbf{D_3}$ receptor, and the like; plus retinoid receptors, such as the various isoforms of RAR (e.g., $\mbox{RAR}\alpha,$ 20 RAR β , or RAR γ), the various isoforms of RXR (e.g., RXR α , $\mathtt{RXR}\beta$, or $\mathtt{RXR}\gamma$), and the like; thyroid receptors, such as ${
m TR}lpha$, ${
m TR}eta$, and the like; as well as other gene products which, by their structure and properties, are considered to be members of the superfamily, as defined hereinabove, 25 including the various isoforms thereof. Examples of orphan receptors include HNF4 [see, for example, Sladek et al., in Genes & Development 4: 2353-2365 (1990)], the COUP family of receptors [see, for example, Miyajima et al., in Nucleic Acids Research 16: 11057-11074 (1988), and Wang et al., in 30 Nature 340: 163-166 (1989)], COUP-like receptors and COUP homologs, such as those described by Mlodzik et al., in Cell 60: 211-224 (1990) and Ladias et al., in Science 251: 561-565 (1991), the ultraspiracle receptor [see, example, Oro et al., in Nature 347: 298-301 (1990)], and the like. Presently preferred members of the superfamily

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for use in the practice c. the present invention are the various isoforms of RXR (e.g., RXR α , RXR β , or RXR γ).

species can modulate the ability of the first receptor to trans-activate transcription of genes maintained under expression control in the presence of ligand for said first receptor. The actual effect on activation of transcription (i.e., enhancement or repression of transcription activity) will vary depending on the receptor species which is combined with either a PPARy or PPAR& receptor to form the multimeric species, as well as on the response element with which the multimeric species interacts.

In accordance with the present invention, there are provided multimeric receptor species which belong to the steroid/thyroid superfamily of receptors, comprising at least the dimerization domain of at two different members of the steroid/thyroid superfamily of receptors, wherein one of the members is selected from the invention PPARy or PPARδ.

20 employed herein, the term "dimerization domain" of a member of the steroid/thyroid superfamily of receptors refers to that portion of the receptor which is believed to be involved in the formation of multimeric receptor species. This domain typically comprises the 25 carboxy-terminal portion of the receptor, i.e., that portion of a receptor which is 3' with respect to the DNA-binding domain of the receptor. See, e.g., Evans, in Science 240:889-895 (1988), and Forman and Samuels, Mol. Endocrinol. 4:1293-1301 (1990). Presently preferred 30 members of the superfamily for use in deriving the dimerization domain are the various isoforms of RXR (e.g., $RXR\alpha$, $RXR\beta$, or $RXR\gamma$).

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In accordance with the present invention, there are also provided heterodimer complexes comprising either PPAR γ or PPAR δ and a silent partner therefor.

As employed herein, the term "silent partner" refers to members of the steroid/thyroid superfamily of receptors which are capable of forming heterodimeric species with either PPARy or PPARδ, wherein the silent partner of the heterodimer does not have any ligand bound to the ligand-binding domain (LBD) when the silent partner is complexed with a PPAR subtype (i.e., only the PPAR copartner of the heterodimer binds ligand). Presently preferred silent partners for use in the practice of the present invention are the various isoforms of RXR (e.g., RXRα, RXRβ, or RXRγ).

In accordance with a further embodiment of the present invention, there is provided a method for screening compounds to determine those which bind to mammalian peroxisome proliferator-activated receptor proteins comprising at least one PPAR subunit of the γ or δ subtype, or functional fragments thereof. Such method comprises employing receptor protein(s) of the invention in a binding assay, which comprises, contacting receptor protein(s) of the invention with test compound, and identifying those compounds which bind to invention receptor protein(s).

In accordance with a still further embodiment of the present invention, there is provided a bioassay for evaluating whether test compounds are agonists for receptor proteins of the invention, or functional modified forms of said receptor protein(s). Such bioassay comprises:

(1) contacting suitable host cells expressing said receptor protein with test compound under physiological conditions, wherein said host cells contain DNA encoding a reporter

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protein, wherein said DNA is operatively linked to a PPAR-response element;

(2) monitoring said host cells for expression of reporter gene, wherein expression of reporter protein reflects transcriptional activity of the receptor protein and, therefore, the presence of an activated receptor-ligand complex.

In accordance with yet another embodiment of the present invention, there is provided a bioassay for evaluating whether test compounds are antagonists for receptor proteins of the invention, or functional modified forms of said receptor protein(s). Such bioassay comprises:

contacting suitable host cells with

- (i) increasing concentrations of at least one compound whose ability to inhibit the transcription activation activity of agonists of mammalian peroxisome proliferator-activated receptor proteins of the γ or δ subtype is sought to be determined, and
- (ii) a fixed concentration of at least
 one agonist for said receptor protein(s) or
 functional modified forms thereof,

wherein suitable test cells express mammalian peroxisome proliferator-activated receptor proteins comprising at least one PPAR subunit of the γ or δ subtype and DNA encoding a reporter protein, wherein said DNA is operatively linked to a PPAR-response element; and thereafter

assaying for evidence of transcription of said reporter gene in said cells as a function of the concentration of said compound in said culture medium, thereby indicating the ability of

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said compound to inhibit activation of transcription by agonists of mammalian peroxisome proliferator-activated receptor proteins comprising at least one PPAR subunit of the γ or δ subtype.

In accordance with a still further embodiment of the present invention, there is provided a method for identifying ligands selective for heterodimers comprising either PPARy or PPAR\$\delta\$ and a silent partner therefor. Such method comprises

comparing the level of expression of reporter when cells containing a reporter construct, either PPARy or PPARδ and silent partner therefor are exposed to test compound, relative to the level of expression of reporter when cells containing a reporter construct, either PPARy or PPARδ and a member of the steroid/thyroid superfamily which is not a silent partner therefor are exposed to test compound, and

selecting those compounds which activate only the combination of either PPAR γ or PPAR δ and silent partner therefor.

In accordance with yet another embodiment of the present invention, there are provided antibodies generated against the invention proteins. Such antibodies can be employed for studying receptor tissue localization, subunit composition, structure of functional domains, as well as in diagnostic applications, therapeutic applications, and the like. Preferably, for therapeutic applications, the antibodies employed will be monoclonal antibodies.

The above-described antibodies can be prepared employing standard techniques, as are well known to those of skill in the art, using the invention receptor proteins or portions thereof as antigens for antibody production. Both anti-peptide and anti-fusion protein antibodies can be

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used [see, for example, Bahouth et al. (1991) <u>Trends Pharmacol Sci.</u> vol. <u>12</u>:338-343; <u>Current Protocols in Molecular Biology</u> (Ausubel et al., eds.) John Wiley and Sons, New York (1989)]. Factors to consider in selecting portions of the invention receptor protein subunit sequences for use as immunogen (as, for example, a synthetic peptide or a recombinantly produced bacterial fusion protein) include antigenicity, accessibility (i.e., internal or external domains), uniqueness to the particular protein subunit, and the like.

The availability of sequence-specific antibodies enables use of immunohistochemical techniques to monitor the distribution and expression density of various protein subunits (e.g., in normal versus diseased brain tissue). Such antibodies can also be employed for diagnostic and therapeutic applications.

In accordance with yet another embodiment of the present invention, there are provided methods for modulating processes mediated by mammalian peroxisome 20 proliferator-activated receptor proteins comprising at least one PPAR subunit of the γ or δ subtype. Such methods comprise contacting mammalian peroxisome proliferator-activated receptor proteins of the γ or δ subtype with an effective, modulating amount of agonist, antagonist or antibody according to the present invention.

The antibodies, agonists and/or antagonists of the invention can be administered to a subject employing standard methods, such as, for example, by intraperitoneal, intramuscular, intravenous, or subcutaneous injection, implant or transdermal modes of administration, and the like. One of skill in the art can readily determine dose forms, treatment regiments, etc, depending on the mode of administration employed.

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Processes which are mediated by mammalian peroxisome proliferator-activated receptor proteins of the γ or δ subtype include, for example, macrophage production in the spleen which is believed to be important in atherosclerosis.

The invention will now be described in greater detail with reference to the following non-limiting examples.

EXAMPLE 1

10 <u>Screening of cDNA libraries</u>

PPARy was isolated by screening an adult mouse liver AZAP cDNA library (Stratagene) with a synthetic oligonucleotide (GGNTTYCAYTAYGGNGTNCAYCG; SEQ ID NO:5) under conditions previously described by Blumberg et al., in Proc. Natl. Acad. Sci. USA 89:2321-2325 (1992). This oligonucleotide is a mixture of all possible DNA sequences encoding the amino acid sequence GFHYGVHA (SEQ ID NO:6), a sequence present in the loop of the first zinc finger in the Xenopus PPARα, PPARß and PPARy isoforms.

PPARδ was isolated by screening an E6.5 mouse AZAPII cDNA library (a gift of D.E. Weng and J.D. Gerhart, Johns Hopkins University) under low stringency conditions with a cDNA fragment encoding the human retinoic acid receptor αDNA binding domain (Mangelsdorf et al., Nature 345:224-229 (1990)). In both screens, positive clones were converted to plasmids by the automatic excision process.

EXAMPLE 2 Cotransfection Assay

The mammalian expression vectors pCMX-PPAR α , 30 pCMX-PPAR γ and pCMX-PPAR δ were constructed by inserting the cDNA inserts of PPAR α , PPAR γ , and PPAR δ into pCMX as

previously described by Umesono et al., in Cell 65:1255-1266 (1991)). Construction of the reporter $PPRE_3$ -TK-LUC has also been previously described by Kliewer et al., (1992) supra. Cotransfection assays in CV-1 cells were done in 48 5 well plates using N-[1-(2,3-dioleoyloxy)-propyl[N,N,Ntrimethyl ammonium methyl sulfate (DOTAP) according to the manufacturer's instructions (Boehringer Mannheim). Transfections contained 10ng of receptor expression plasmid vector, 20ng of the reporter $PPRE_3$ -TK-LUC, 60ng of pCMX-BGAL 10 (ß-galactosidase) as an internal control, and 210ng of carrier plasmid pGEM. Cells were incubated in the presence of DOTAP for 8 hours, washed, and incubated in the presence of peroxisome proliferators or fatty acids for 36 hours. Cell extracts were prepared and assayed for luciferase and 15 ß-galactosidase activity as previously described (Umesono, supra). All experimental points were done in triplicate.

EXAMPLE 3 Northern Analysis

Preparation of poly(A) * RNA from rat tissues and Northern analysis were performed as previously described (Mangelsdorf et al., supra). Thus, Northern blot analysis of PPAR mRNA was carried out employing adult and embryonic tissue. Adult male rat tissues and mouse embryos from gestation day 10.5 to 18.5 were employed. The exposure time for each of the blots was 48 hours. The sizes of the transcripts, based on RNA size markers, were 8.5kb (PPARa), 1.9kb (PPARa), and 3.5kb (PPARa).

EXAMPLE 4 DNA Binding Assays

Gel mobility shift assays were performed as previously described by Kliewer et al. (1992) supra. PPAR α , PPAR γ , PPAR δ , RXR α , RXR β and RXR γ were synthesized in vitro using the TNT coupled transcription/translation

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system (Promega) according to the manufacturer's instructions.

EXAMPLE 5

Isolation of three murine PPAR isoforms

The function of peroxisome proliferators has been most extensively studied in rodents, where treatment with these compounds results in marked increases in peroxisome size and number and concomitant increases in the expression of the genes encoding the enzymes of the peroxisomal β-oxidation pathway. To gain insight into the function of PPAR isoforms, mouse embryonic and adult liver libraries were screened for PPARα-related gene products. In addition to PPARα, two types of PPARα-related clones were isolated.

The first clone encodes a 475-amino acid protein that is 56% identical to mouse (m) PPARα and 76% identical to Xenopus (x) PPARγ. Since this clone is 97% and 84% identical to the DNA binding and ligand binding domains of xPPARγ, respectively, it is designated mPPARγ (see SEQ ID NOs:1 and 2).

The second clone encodes a 440-amino acid protein that is closely related to NUC-1 (see SEQ ID NOs:3 and 4, and Figure 1), a PPARα-related receptor recently isolated from a human osteosarcoma library (see Schmidt et al., in Mol. Endo. 6:1634-1641 (1992)). Since this second clone is not highly homologous to any of the previously identified PPAR isoforms (i.e., mPPARα, xPPARα, xPPARβ or xPPARγ; see Figure 1), it appears to represent a novel receptor, and is, therefore, designated mPPARδ. Of the approximately 50 positives characterized during the course of screening, no mouse homolog of xPPARβ was identified.

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EXAMPLE 6

PPARα, PPARγ, and PPARδ are differentially expressed in the adult and embryo

The expression patterns of the murine PPAR isoforms were examined in the embryo and adult. Northern analysis of poly(A) * RNA isolated from adult male rat tissues revealed differential yet overlapping patterns of expression of the three isoforms. Both PPARα and PPARδ are widely expressed, with PPARα message levels highest in the liver, kidney, heart, and adrenal, and PPARδ message highest in the heart, adrenal, and intestine. In contrast, PPARγ displays a more restricted distribution pattern, with abundant expression in only the adrenal and spleen, although message is also detectable in the heart, kidney, and intestine.

The developmental expression of the PPAR isoforms was also examined through Northern analysis of whole mouse embryo RNA. PPARα and PPARγ displayed similar expression patterns during mouse embryogenesis, with message first appearing at day 13.5 postconception and increasing until birth. In contrast, PPARδ message was abundant at all the embryonic time points tested, suggesting a broad role for this isoform during development. Thus, the PPAR isoforms are seen to be differentially expressed in both the embryo and the adult.

EXAMPLE 7

Evidence for pharmacological differences between $\frac{\text{PPAR}\alpha,\ \text{PPAR}\gamma\ \text{and}\ \text{PPAR}\delta}{\text{PPAR}}$

The relatively high degree of conservation within 30 the ligand binding domains of PPAR α , PPAR γ and PPAR δ suggested that these PPAR isoforms might respond to the same activators. Accordingly, each of the PPAR isoforms

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was first tested for responsiveness to Wy 14,643, a peroxisome proliferator and potent activator of PPARα (Reddy and Lalwani, Crit. Rev. Toxicol. 12:1-58 (1983)). Cotransfection of PPARα expression plasmid resulted in a dramatic (>100-fold) increase in activation of a reporter construct containing three copies of the acyl-CoA oxidase PPRE (AOX-PPRE) upstream of the thymidine kinase promoter driving luciferase expression (PPRE3-TK-LUC) in response to Wy 14,643 (Figure 2).

10 In contrast, no activation of reporter expression was seen in the presence of Wy 14,643 upon cotransfection of PPAR γ or PPAR δ expression plasmids (Figure 2). lack of activation is unlikely to reflect differences in binding site specificity, as each of the PPAR isoforms 15 bound efficiently to the AOX-PPRE as a heterodimer with RXR (as determined by gel mobility shift assays done using in vitro synthesized PPAR α , PPAR γ , and PPAR δ , and/or RXR γ , and 32P-labeled AOX-PPRE oligonucleotide). Additional experiments revealed that overexpression of PPAR γ and PPAR δ interfered with the ability of PPAR α to activate through the AOX-PPRE (Figure 3). Thus, both PPAR γ and PPAR δ are expressed and can function as dominant repressors of $\mathtt{PPAR}\alpha extsf{-}$ mediated responsiveness to Wy 14,643.

Since no activation of PPAR γ and PPAR δ was 25 detected with Wy 14,643, other potential activators were tested, including a broad spectrum of peroxisome proliferators and fatty acids. As shown in Figure 4, significant activation of PPARy was obtained upon treatment with LY-171883, a leukotriene antagonist and peroxisome 30 proliferator which lacks the carboxyl group typically found in this class of compounds (Foxworthy and Eacho, Biochem. Pharmacology 42:1487-1491 (1991)). Conversely, activation of PPARy was seen in the presence of linoleic acid (Figure 4).

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In contrast to the results obtained with PPARy, PPARδ was activated in the presence of lincleic acid, but was not activated upon treatment with LY-171883. Both LY-171883 and linoleic acid are strong activators of PPARα (Figure 4). Interestingly, each of the three PPAR isoforms was activated with a distinct rank order of efficacy by these compounds:

PPAR α :

Wy 14,643 > LY-171883 > linoleic acid;

10 PPARy:

LY 171883 > linoleic acid > Wy 14,643;

PPAR δ :

linoleic acid > LY-171883 and Wy 14,643.

See Figure 4. These data provide evidence that PPARy and 15 PPARδ can function as regulated activators of gene expression and that the three PPAR isoforms are pharmacologically distinct.

While the invention has been described in detail with reference to certain preferred embodiments thereof, it will be understood that modifications and variations are within the spirit and scope of that which is described and claimed.

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SEQUENCE LISTING

SEQ ID NO:1 Mouse PPARy

- 1	ATCHATCCCGCCCCAGGCGCTCTGAGTGCGACGGGCCCGCCTGGCCGGCC	
61	GGAGGACGCGGAAGAAGACCTGGGGCGCTGCCTGGGGTATTGGGTCGCGCGCAGTGAG	
121	GGGACCGAGTGTGACGACAAGGTGACCGGGCTGAGGGACGGGCTGAGGAGAAGTCACAC	
181	TCTGACAGGAGCCTGTGAGACCAACAGCCTGACGGGGTCTCGGTTGAGGGGACGCGGGCT	
241	GAGAAGTCACGTTCTGACAGGACTGTGTGACAGACAAGATTTGAAAGAAGCGGTGAACCA	
301	CTGATATTCAGGACATTTTTAAAAACAAGACTACCCTTTACTGAAATTACCATGGTTGAC	
	M V D	3
361	ACAGAGATGCCATTCTGGCCCACCAACTTCGGAATCAGCTCTGTGGACCTCTCCGTGATG	3
301	~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~	
401		23
421	GAAGACCACTCGCATTCCTTTGACATCAAGCCCTTTACCACAGTTGATTTCTCCAGCATT	
	EDHSHSFDIKPFTTVDFSSI	43
481	TCTGCTCCACACTATGAAGACATTCCATTCACAAGAGCTGACCCAATGGTTGCTGATTAC	
	SAPHYEDIPFTRADPMVADY	63
541	AAATATGACCTGAAGCTCCAAGAATACCAAAGTGCGATCAAAGTAGAACCTGCATCTCCA	
	K Y D L K L Q E Y Q S A I K V E P A S P	83
601	CCTTATTATTCTGAAAAGACCCAGCTCTACAACAGGCCTCATGAAGAACCTTCTAACTCC	63
001	P Y Y S E K T O L Y N R P H E E P S N S	
661		103
661	CTCATGGCCATTGAGTGCCGAGTCTGTGGGGATAAAGCATCAGGCTTCCACTATGGAGTT	
	L M A I E C R V C G D K A S G F H Y. G V	123
721	CATGCTTGTGAAGGATGCAAGGGTTTTTTCCGAAGAACCATCCGATTGAAGCTTATTTAT	
	HACEGCKGFFRRTIRLKLIY	143
781	GATAGGTGTGATCTTAACTGCCGGATCCACAAAAAAAGTAGAAATAAAT	
	D R C D L N C R I H K K S R N K C Q Y C	163
841	CGGTTTCAGAAGTGCCTTGCTGTGGGGGATGTCTCACAATGCCATCAGGTTTGGGCGGATG	103
		100
901	R F Q K C L A V G M S H N A I R F G R M	183
901	CCACAGGCCGAGAAGGAGAAGCTGTTGGCGGAGATCTCCAGTGATATCGACCAGCTGAAC	
061	PQAEKEKLLAEISSDIDQLN	203
961	CCAGAGTCTGCTGATCTGCGAGCCCTGGCAAAGCATTTGTATGACTCATACATA	
	PESADLRALAKHLYDSYIKS	223
1021	TTCCCGCTGACCAAAGCCAAGGCGAGGGGGGGTCTTGACAGGAAAGACAACGGACAAATCA	
	F P L T K A K A R A I L T G K T T D K S	243
1081	CCATTTGTCATCTACGACATGAATTCCTTAATGATGGGAGAAGATAAAATCAAGTTCAAA	240
	D D U T V D V N A * 14 14	262
1141	CATATCACCCCCTGCAGGAGCAGAGCAAGAGGTGGCCATCCGAATTTTTCAAGGGTGC	263
	H I T P L O E O S K E V A I R I F O G C	
1201		283
1201	CAGTTTCGATCCGTAGAAGCCGTGCAAGAGATCACAGAGTATGCCAAAAATATCCCTGGT	
	Q F R S V E A V Q E I T E Y A K N I P G	303
1261	TTCATTAACCTTGATTTGAATGACCAAGTGACTCTGCTCAAGTATGGTGTCCATGAGATC	
	FINLDLNDOVTLLKYGVHET	323
1321	ATCTACACGATGCTGGCCTCCCTGATGAATAAAGATGGAGTCCTCATCTCAGAGGGCCAA	
	I Y T M L A S L M N K D G V L I S E G O	343
1381	GGATTCATGACCAGGGAGTTCCTCAAAAGCCTGCGGAAGCCCTTTGGTGACTTTATGGAG	343
		262
1441	CCTAAGTTTGAGTTTGCTGTGAAGTTCAATGCACTGGAATTAGATGACAGTGACTTGGCT	363
	D V D D D X 17 V D 31 3	
1501		383
1501	ATATTTATAGCTGTCATTATTCTCAGTGGAGACCGCCCAGGCTTGCTGAACGTGAAGCCC	
	I F I A V I I L S G D R P G L L N V K P	403
1561	ATCGAGGACATCCAAGACAACCTGCTGCAGGCCCTGGAACTGCAGCTCAAGCTGAATCAC	
	I E D I Q D N L L Q A L E L O L K T. N H	423
1621	CCAGAGTCCTCTCAGCTGTTCGCCAAGGTGCTCCAGAAGATGACAGACCTCAGGCAGATC	
	PESSQLFAKVLQKMTDLRQI	442
1681	GTCACAGAGCACGTGCAGCTACTGCATGTGATCAAGAAGACAGAC	443
1001	V T E H V O L L H V I K K T F T D M C I	
7747		463
1741	CACCCCTGCTCCAGGAGATCTACAAGGACTTGTATTAGCAGGAAAGTCCCACCCGCTGA	
	HPLLQEIYKDLY*	475
1801	CAACGTGTTCCTTCTATTGATTGCACTATTATTTTGAGGGAAAAAATCTGACACCTAAG	
1861	AAATTTACTGTGAAAAAGCATTTAAAAACAAAAGTTTTAGAACATGATCTATTTTATGC	
1921	ATATTGTTTATAAAGATACATTTACAATTTACTTTTAAATATTAAAAATTACCACATTATA	
1981	AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	

SEQ ID NO:2

MVDTEMPFWPTNFGISSVDLSVMEDHSHSFDIKPFTT V D F S S I S A P H Y E D I P F T R A D P M V A D Y K Y D L K L Q E Y Q S AIKVEPASPPYYSEKTQLYNRPHEEPSNSLMAIECRV C G D K A S G F H Y G V H A C E G C K G F F R R T I R L K L I Y D R C D L N C R I H K K S R N K C Q Y C R F Q K C L A V G M S H N A I R F G R M P Q A E K E K L L A E I S S D I D Q L N P E S A D L R A L A K H L Y D S Y I K 225 240 255 SFPLTKAKARAILTGKTTDKSPFVIYDMNSLMMGEDK IKFKHITPLQEQSKEVAIRIFQGCQFRSVEAVQEITE YAKNIPGFINLDLNDQVTLLKYGVHEIIYTMLASLMN KDGVLISEGQGFMTREFLKSLRKPFGDFMEPKFEFAV KFNALELDDSDLAIFIAVIILSGDRPGLLNVKPIEDI Q D N L L Q A L E L Q L K L N H P E S S Q L F A K V L Q K M T D L R Q I V TEHVQLLHVIKKTETDMSLHPLLQEIYKDLY

SEQ ID NO:3

Mouse PPAR δ

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	GCG	GGA	Cgg	Jugo	CAC	JCGG	GCG	CGA	GAG	GCG	GCC	GGg	ACA	GTG	CTG:	rgca	GCG	GTG	TGG	GTA	
121	TGC	GCA	TGG	GAC	CTC	CT	CAG	AGG	CTC	CTG	CTC	ACT	'GAC	'AGA'	TGA	AGAC	'AAA	CCC	ACG	GTA	
181	AAG	GCA	GTC	CAT	CTG	CGC	CTC	AGA(CCC	AGA	TGG	TGG	CAG	AGC:	TAT	ACC	AGG	CCT	GCA	GCG	
241	CCA	CGc	CAA	GTC	GGG	GT(CAG'	rca'	TGG:	AAC	AGC	CAC	AGG	AGG	AGA	CcCC	TGA	GGC	CCG	GGA	
								M						E		P	E	Α	R	E	13
301	AGA	GGA	GAA	AAG	AGG?	AAG'	TGG	CCA	TGG	GTG	ACG	GAG	CCC	CGG	AGC'	CA	TGG	GGG	ACC	AGA	
	E	E	K	E	E	V				D						N	G	G	P	E	33
361	ACA	CAC	GCI	TCC	CTTC	CCA	GCA	GCT	GTG	CAG	ACC	TCT	ccc	AGA	aTT(CTC		
	Ħ	T	L	P	s				A					N Q		S	P	s	S	T.	53
421	GCT	GGA	CCA			AGA'				ATG	GGG	CCT	יראני זמטי	GCG	בראו מראו					እጥር ፲	
	T.						G				A			G			N.	M	E.	WIG.	7.3
481	_	_												GGG						CEC	13
201	R				D	K	A			F	H	Y	G							GIG	93
541		•	_	_	_			_	_					v AGT							93
241																				GAT	
	K	_	F	F	R		_	I				L				K	С	D	R	I	113
601														ACT				GAA	GTG	CCT	
	_	K												. C				K		L	133
661	GGCACTCGGCATGTCGCACAACGCTATCCGCTTTGGACGGATGCCGGACGGCGAGAAGAG																				
	A	L	G	M	s	Н	• • •	Α	_							D	G	E	K	R	153
721	GAA	GCT	GGI	GGC	CGGC	GC'	TGA	CTG	CCA	GCG	AGG	GGT	GCC	AGC	ACA	ACCC	CCA	GCT	GGC	CGA	
	K	_	V	Α		L		Α				С		H		P	Q	L	Α	D	173
781	CCT	GAA	GGC	CTI	CTC	CTA	AGC	ACA'	TCT.	ACA	ACG	CCT	ACC	TGA	AAA	ACTI	CĀA	CAT	GAC	CAA	
	L	K	Α		S	K						Y				F	N	М	Т	K	193
841	AAA	GAA	GGC	CCC	GAC	CA!	TCC'	rca (CCG	GCA	AGT	CCA	GCC	ACA	ACG	CACC	CTT	TGT	CAT	CCA	
	K	K	A	R	s	I	L	Т	G	K						P	F	v	T	Н	213
901	CGA	CAT	CGA	GAC	'AC'	ርርጥ(GGC	AGG	_		_	-		TGT							213
	D	T	E	Т		W					G		. 1				L	V.			233
	U	_	ند	_		44	7	u	£,	1	G	· +	, ,	, w	V.	Q	L	V	N	G	233

961	GCTGCCGCCCTACAACGAGATCAGTGTGCACGTGTTCTACCGCTGCCAGTCCACCACAGT											
	L P P Y N E I S V H V F Y R C O S T T V	253										
1021	GGAGACAGTCCGAGAGCTCACCGAGTTCGCCAAGAACATCCCCAACTTCAGCAGCCTCTT											
	ETVRELTEFAKNIPNFSSLF	273										
1081	CCTCAATGACCAGGTGACCCTCCTCAAGTATGGCGTGCACGAGGCCATCTTTGCCATGCT											
	LNDQVTLLKYGVHEAIFAML	293										
1141	GGCCTCCATCGTCAACAAGACGGGCTGCTGGTGGCCAACGGCAGTGGCTTCGTCACCCA	•										
	ASIVNKDGLLVANGSGFVTH	313										
1201	CGAGTTCTTGCGAAGTCTCCGCAAGCCCTTCAGTGACATCATTGAGCCCAAGTTCGAGTT											
	EFLRSLRKPFSDIIEPKFEF	333										
1261	TGCTGTCAAGTTCAATGCGCTGGAGCTCGATGACAGTGACCTGGCGCTCTTCATCGCGGC	-										
	AVKFNALELDDSDLALFIAA	353										
1321	CATCATTCTGTGTGGAGACCGGCCAGGCCTCATGAATGTGCCCCAGGTAGAAGCCATCCA	000										
	I I L C G D R P G L M N V P Q V E A I O	373										
1381	GGACACCATTCTGCGGGCTCTAGAATTCCATCTGCAGGTCAACCACCCTGACAGCCAGTA											
	D T I L R A L E F H L Q V N H P D S Q Y	393										
1441	CCTCTTCCCCAAGCTGCTGCAGAAGATGGCAGACCTGCGGCAGCTGGTCACTGAGCATGC	• • • • • • • • • • • • • • • • • • • •										
	LFPKLLQKMADLRQLVTEHA	413										
1501	CCAGATGATGCAGTGGCTAAAGAAGACGGAGAGTGAGACCTTGCTGCACCCCCTGCTCCA											
	Q M M Q W L K K T E S E T L L H P L L O	433										
1561	GGAAATCTACAAGGACATGTACTAAGGCCGCagCCCaggCCtCCCCtCaggCtCtgCtgg											
	E I Y K D M Y 440											
1621	gCCCagCCaCggaCtGTTCAGAGGACCAGCCACAGGCACTGGCAGTCAAGCAGCTAGAGC											
1681	CTACTCACAACACTCCAGACACGTGGCCCAGACTCTTCCCCCAACACCCCCCACCCCCACC											
1741	AACCCCCCATTCCCCCAACCCCCTCCCCCACCCGCTCTCCCCATGGCCCGTTTCCTG											
1801	TTTCTCCTCAGCACCTCCTGTTCTTGCTGTCTCCCTAGCGCCCCTTGCTCCCCCCCC											
1861	CCTTCCTTCTCTAGCATCCCCCTCCTCCCAGTCCTCACATTTGTCTGATTCACAGCAGAC											
1921	AGCCCGTTGGTACGCTCACCAGCAGCCTAAAAGCAGTGGGCCTGTGCTGGCCCAGTCCTG											
1981	CCTCTCCTCTATCCCCTTCAAAGGGAATTC 2012											

SEQ ID NO:4

MEQPQEETPEAREEEKEEVAMGDGAPELNGGPEHTLP S S S C A D L S Q N S S P S S L L D Q L Q M G C D G A S G G S L N M E C R V C G D K A S G F H Y G V H A C E G C K G F F R R T I R M K L E Y E K C D RICKIQKKNRNKCQYCRFQKCLALGMSHNAIRFGRMP D G E K R K L V A G L T A S E G C Q H N P Q L A D L K A F S K H I Y N A Y LKNFNMTKKKARSILTGKSSHNAPFVIHDIETLWQAE KGLVWKQLVNGLPPYNEISVHVFYRCQSTTVETVREL T E F A K N I P N F S S L F L N D Q V T L L K Y G V H E A I F A M L A S I V N K D G L L V A N G S G F V T H E F L R S L R K P F S D I I E P K F E F AVKFNALELDDSDLALFIAAIILCGDRPGLMNVPQVE AIQDTILRALEFHLQVNHPDSQYLFPKLLQKMADLRQ LVTEHAQMMQWLKKTESETLLHPLLQEIYKDMY

SEQ ID NO:5

GGNTTYCAYT AYGGNGTNCA YCG 23

SEQ ID NO:6

GFHYGVHA

That which is claimed is:

- 1. An isolated mammalian peroxisome proliferator-activated receptor subunit protein of the γ or δ subtype, or functional fragments thereof.
- 2. A protein according to claim 1 wherein said protein is of the γ subtype, or functional fragments thereof.
- 3. A protein according to claim 2 having an amino acid sequence substantially the same as set forth in SEQ ID No:2.
- 4. A protein according to claim 1 wherein said protein is of the δ subtype, or functional fragments thereof.
- 5. A protein according to claim 4 having an amino acid sequence substantially the same as set forth in SEQ ID NO:4.
- 6. A receptor recombinantly expressed in a cell containing a vector comprising isolated nucleic acid encoding a protein according to claim 1.
- 7. A receptor according to claim 6 comprising at least one PPARy subunit or at least one PPAR δ subunit and at least one retinoid X receptor isoform.
- 8. A receptor according to claim 6, wherein said receptor is a heteromultimer.
- 9. A receptor according to claim 7, wherein the PPAR subunit is PPARy.

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10. A receptor according to claim 7, wherein the PPAR subunit is PPAR δ .

- 11. A mammalian peroxisome proliferator-activated receptor, expressed recombinantly in a host cell, said receptor having the ability to repress PPARα-mediated responses activated by Wy 14,643, comprising at least one PPAR subunit, wherein the PPAR subunit is PPARγ or PPARδ, and at least one retinoid X receptor isoform.
 - 12. A mammalian peroxisome proliferator-activated subunit protein, expressed recombinantly in a host cell comprised of an amino acid sequence shown in SEQ ID Nos: 2 or 4.
- 13. A method for screening compounds to determine those which bind to mammalian peroxisome proliferator-activated receptor proteins comprising at least one PPAR subunit of the γ or δ subtype, or functional fragments thereof, said method comprising:

contacting receptor protein according to claim 11 with a test compound, and

identifying those compounds which bind to invention receptor proteins.

14. A bioassay for evaluating whether test compounds are agonists for receptors according to claim 11, or functional modified forms of said receptor(s), said bioassay comprising:

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- (1) contacting suitable host cells expressing said receptor with test compound under physiological conditions, wherein said host cells contain DNA encoding a reporter protein, wherein said DNA is operatively linked to a PPAR-response element;
- (2) monitoring said host cells for expression of reporter protein, wherein expression of

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reporter protein reflects transcriptional activity of said receptor protein and the presence of an activated receptor-ligand complex.

15. A bioassay for evaluating whether test compounds are antagonists for receptors according to claim 11, or functional modified forms of said receptor(s), said bioassay comprising:

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contacting suitable host cells with

- (i) increasing concentrations of at least one compound whose ability to inhibit the transcription activation activity of agonists of mammalian peroxisome proliferator-activated receptors of the γ or δ subtype is sought to be determined, and
- (ii) a fixed concentration of at least
 one agonist for said receptor protein(s) or
 functional modified forms thereof,

wherein suitable host cells express mammalian peroxisome proliferator-activated receptors comprising at least one PPAR subunit of the γ or δ subtype, and wherein said host cells contain DNA encoding a reporter protein, wherein said DNA is operatively linked to a PPAR-response element; and thereafter

assaying for evidence of transcription of said reporter gene in said cells as a function of the concentration of said compound in said culture medium, thereby indicating the ability of said compound to inhibit activation of transcription by agonists of mammalian peroxisome proliferator-activated receptors of the γ or δ subtype.

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- 16. An antibody generated against the receptor of claim 11.
- 17. An antibody according to claim 16, wherein said antibody is a monoclonal antibody.
- 18. A heterodimer complex comprising PPAR γ or PPAR δ and an isoform of RXR.
- 19. A heterodimer complex according to claim 18 wherein said isoform of RXR is selected from RXR α , RXR β or RXR γ .
- 20. A combination of receptors comprising at least two different members of the steroid/thyroid superfamily of receptors, wherein said receptors are associated in the form of a multimer, and wherein one of said members is selected from PPARy or PPARS.
- 21. A combination of receptors according to claim 20 wherein said combination is in the form of a heterodimer.
- 22. A multimer comprising at least the dimerization domain of at least two different members of the steroid/thyroid superfamily of receptors, wherein one of said members is selected from PPARy or PPARs.
- 23. An isolated nucleic acid encoding a mammalian peroxisome proliferator-activated receptor subunit protein of the γ or δ subtype, or functional fragments thereof.
- 24. An isolated nucleic acid according to claim 23, wherein said protein is of the y subtype, or functional fragments thereof.

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- $25.\,$ An isolated nucleic acid according to claim 23, wherein said protein is of the δ subtype, or functional fragments thereof.
- 26. An isolated nucleic acid according to claim 23 having a contiguous nucleotide sequence substantially the same as set forth in SEQ ID NO:1 [PPAR-y], or SEQ ID NO:3 [PPAR-δ], or variations thereof which encode the same 5 amino acid sequence, but employ different codons for some of the amino acids, or splice variant nucleotide sequences thereof.
 - 27. An isolated nucleic acid according to claim 23 having a contiguous nucleotide sequence substantially the same as set forth in SEQ ID NOs:1 or 3.
 - 28. Isolated and purified nucleic acid, or functional fragments thereof encoding the protein of claim 1, selected from:
 - (a) nucleic acid encoding the amino acid sequence set forth in SEQ ID NO:2 or SEQ ID NO:4, or
 - (b) nucleic acid that hybridizes to the nucleic acid of (a) under moderately stringent conditions, wherein said nucleic acid encodes biologically active PPAR- γ or PPAR- δ , or
 - (c) nucleic acid degenerate with respect to either (a) or (b) above, wherein said nucleic acid encodes biologically active PPAR- γ or PPAR- δ .
 - 29. A vector comprising a nucleic acid according to claim 23.
 - 30. A cell containing a nucleic acid according to claim 23.

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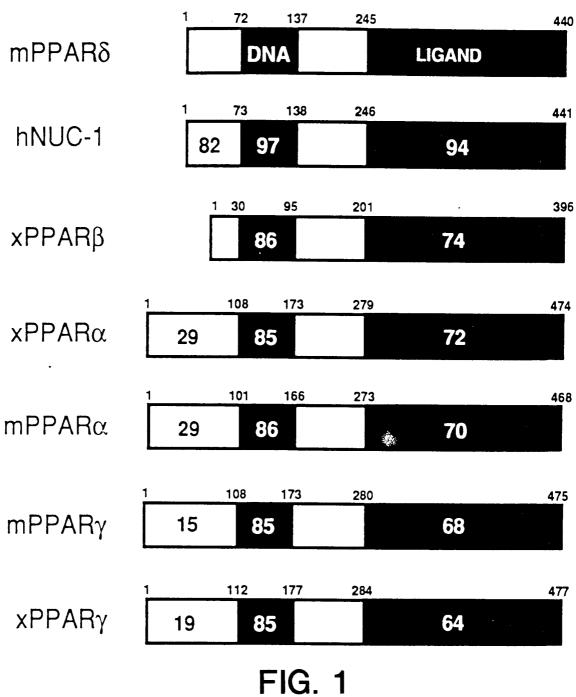
- 31. A cell according to claim 30, wherein said cell further comprises nucleic acid encoding at least one retinoid X receptor isoform.
- 32. A cell containing a vector according to claim 29.
- 33. A cell according to claim 32, wherein said cell further comprises nucleic acid encoding at least one retinoid X receptor isoform.
- 34. A method for the recombinant production of a mammalian peroxisome proliferator-activated receptor comprising at least one PPAR subunit of the γ or δ subtype, or functional fragments thereof, said method comprising expressing the nucleic acid of claim 23 in a suitable host cell.
 - 35. A method according to claim 34, wherein the suitable host cell contains nucleic acid encoding at least one retinoid X receptor.
 - 36. A host cell according to claim 35, wherein said host cell is a CV-1 cell.
 - 37. An isolated nucleic acid fragment useful as a hybridization probe, wherein said fragment comprises at least 14 contiguous nucleotides of the nucleic acid according to claim 23, and wherein said fragment is labeled with a detectable substituent.
 - 38. A method to identify clones encoding mammalian peroxisome proliferator-activated receptor subunit proteins of the γ or δ subtype, or functional fragments thereof, said method comprising:

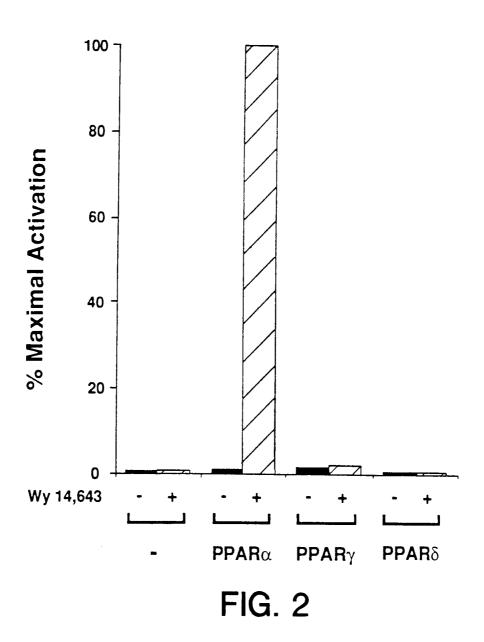
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screening a genomic or cDNA library with a nucleic acid probe according to claim 37 under low stringency hybridization conditions, and

identifying those clones which display a substantial degree of hybridization to said fragment.





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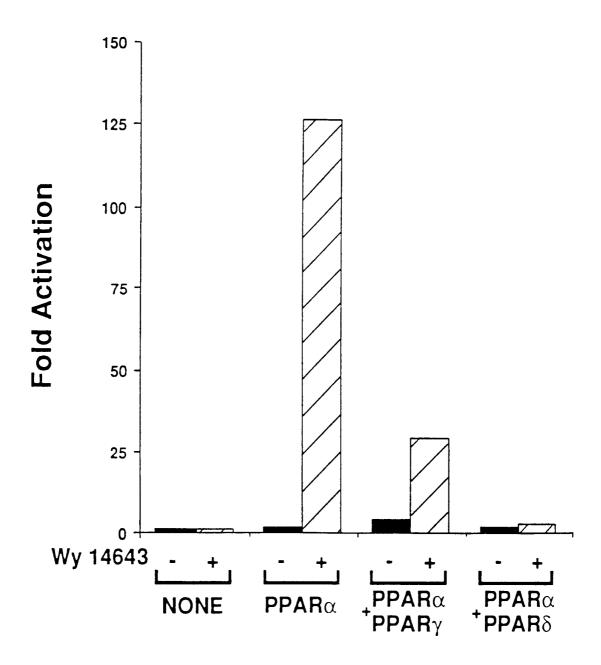


FIG. 3

