Nuclear Receptor Resource Research Article The Retinoid-Related Orphan Receptors P.P. Albrecht, J.P. Vanden Heuvel, INDIGO Biosciences Inc., State College PA

Retinoid-Related Orphan Receptors (RORs)

Retinoid-related orphan receptors (RORs) are nuclear receptors (NR) which belong to the family of Thyroid hormone receptors. RORs play a role in various biological processes such as circadian rhythm, immune response, regulation of metabolic pathways, energy homeostasis, and thymopoiesis. They are also known to play a role in several pathologies such as cancer, osteoporosis, hypoxia, autoimmune diseases, asthma etc (Jetten A.M. 2009).

ROR genes encode proteins of 459 to 556 amino acids. Similar to other nuclear receptors, they have four major functional domains (Figure 1). Domain A/B, also referred to as AF-1 domain is the N-terminal domain. It is followed by DNA-binding domain (DBD) which is highly conserved in different RORs. Other two domains include a hinge domain and a ligand-binding domain which is also the C-terminal domain. (Jetten A.M. 2009).

There are three subtypes of RORs namely ROR α (NR1F1, RORA or RZR α), ROR β (NR1F2, RORB or RZR β), and ROR γ (NR1F3, RORC, TOR). ROR α is present in several organs such as brain, heart, liver, testis, and skin. ROR β is found in sensory processing areas of central nervous system such as pineal gland and retina. ROR γ is highly expressed in thymus and other tissues such as liver (hepatocytes), kidney, heart, lung etc (Jetten A.M. & Ueda E. 2002, Kallen J.A. et al 2002).

ROR gene generates several isoforms of each subtype depending upon alternative promoter usage and exon splicing. All these isoforms differ in mainly their amino terminus. Four isoforms have been identified for ROR α (α 1-4). ROR β gene is expressed in only one isoform in humans whereas for ROR γ - two isoforms have been identified (γ 1 and γ 2 also known as γ t). Most isoforms present a distinct tissue-specific expression which results in involvement of regulation of different physiological processes and target genes. For example, human ROR α 3 is only found in human testis (Steinmayr M. et al 1998). All three RORs play important roles, more specifically: **ROR\alpha**

• Plays a critical role in the development of the cerebellum



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Figure 1. Basic Structure of the retinoid-related orphan receptors (RORs). Shown is the crystal structure of RARa bound to DNA or its natural ligand cholesterol sulfate

- Required for maturation of photoreceptors in the retina
- May play a role in thymopoiesis and lymphocyte development
- Plays a role in Th17 cell differentiation and IL-17 expression

RORβ

- Required for maturation of photoreceptors in the retina
- May play a role in control of circadian rhythm

RORy

- Is essential for the development of secondary lymphoid tissues including lymph nodes
- Plays a critical role in thymopoiesis
- Plays a role in Th17 cell differentiation and IL-17 expression

2. Basic Mechanism of Action

RORs bind to ROR-specific DNA response elements (ROREs) as monomers. They do not form heterodimers with retinoid-X-receptors (RXRs) like RARs. The ROREs consist of the consensus RGGTCA core motif preceded by a 6-bp A/T-rich sequence, in the regulatory region of target genes (Jetten A.M. 2009). Different subtypes of RORs recognize and show different affinities to different ROREs as a result of the role of A/B domain and due to different promoters.

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Figure 2. Regulation of gene expression by RORs.

It is also known that REV-ERB nuclear receptors are able to inhibit ROR-mediated transcriptional activation by competing for binding with ROREs.

Many co-regulators are found among protein complexes which affect the gene transcription of RORs. For example, coactivators such as NCOA1 (SRC1), NCOA2 (TIF2 or GRIP1), PGC-1 α , p300, and CBP or corepressors such as NCOR1, NCOR2, RIP140, NIX1, and FOXP3.

3. RORs as a Possible Therapeutic Target

Spinocerebellar ataxia type 1 (SCA1) - is an autosomal dominant inherited neurodegenerative disorder characterized by progressive loss of motor coordination, speech impairement, and problems swallowing. It was concluded that decreased levels of ROR α protein in SCA1 Purkinje cells affects development and function of Purkinje cells which results in neurodegeneration (Serra H.G. et al 2006, Steinmayr M. et al 1998).

Autoimmune and inflammatory diseases such as lupus, arthritis, multiple sclerosis, atopic dermatitis, psoriasis, asthma, inflammatory bowel disease – Th17 cells have been implicated in all of the above pathologies. Additionally IL-17 induces secretion of various proinflammatory cytokines. It was shown in studies with mice with loss of ROR α and γ function greatly reduces susceptibility to autoimmune diseases or inflammatory



conditions such as lung inflammation induced by ovalbumin.

ROR α is known to impede NF- \varkappa B activity resulting in reduced secretion of interleukins and cox-2 (Duez H. & Staels B. 2008). Additionally ROR α has been shown to play a role in ischemia-induced angiogenesis and to regulate expression of endothelial NO synthase (Duez H. & Staels B. 2008).

Lipid and steroid metabolism – Many recent studies suggest that RORs play an important role in regulation of lipid and steroid metabolism. ROR α activates transcription of apolipoproteins. Additionally ROR α in skeletal muscle may regulate muscle lipid metabolism (Duez H. & Staels B. 2008).

Cancer – Many studies indicate that RORs may play a role in cancer. Mice deficient in the expression of ROR γ exhibit high incidence of thymic lymphomas that metastasize (Jetten & Ueda, 2002). Gastric tumors and ovarian cancers show an increase in the population of Th17 cells (Miyahara Y. et al 2008, Zhang et al 2008). Studies also indicate that ROR α gene is down-regulated in several tumor-types such as breast and lung cancer (Lu et al 2007)

4. Natural ROR Modulators

In vitro studies indicate that cholesterol, 7dehydrocholesterol and cholesterol sulfate can act as ROR α agonists (Kallen JA et al 2002). In vitro studies also indicate ATRA can bind to ROR β and ROR γ to inhibit the receptor-mediated transcription (Stehlin-Gaon et al 2003) thus acting as antagonists. It is still not known whether in vivo ROR activity is regulated by endogenous ligands.

5. References

- Duez, H. and Staels, B. (2008). The Nuclear Receptors Rev-erbs And RORs Integrate Circadian Rhythms and Metabolism. Diab Vasc Dis Res 5: 82-8.

- Jetten, A.M. (2009). Retinoid-related Orphan Receptors (RORs): Critical Roles In Development, Immunity, Circadian Rhythm, and Cellular Metabolism. Nucl Recept Signal 7: e003.

- Jetten, A.M. & Ueda, E. (2002). Retinoid-Related Orphan Receptors (RORs): Roles in Cell survival, Differentiation And Disease. Cell Death Differ 9: 1167-71.

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Kallen, J.A. et al (2002). X-ray Structure of the hRORalpha LBD at 1.63A: Structural And Functional Data That Cholesterol Or A Cholesterol Derivative Is The Natural Ligand of RORalpha. Structure 10: 1697-707.
Lu, Y. et al (2007). Common Human Cancer Genes Discovered by Integrated Gene-Expression Analysis PLoS ONE 2: e1149.

- Miyahara, Y. et al (2008). Generation And Regulation of Human CD4+ IL-17-producing T Cells in Ovarian Cancer. Proc Natl Acad Sci USA 105: 15505-10.

- Serra, H.G. et al (2006). RORα-mediated Purkinje Cell Development Determines Disease Severity In Adult SCA1 Mice. Cell 127: 697-708.

- Stehlin-Gaon, C. et al (2003). All-trans Retinoic Acid Is A Ligand For The Orphan Nuclear Receptor ROR β . Nat Struct Biol 10: 820-5.

- Steinmayr, M. et al (1998). staggerer Phenotype In Retinoid-Related Orphan Receptor α -deficient Mice. Proc Natl Acad Sci USA 95: 3960-5.

- Zhang, F. et al (2008). Interactions Among The Transcription Factors RunX1, RORgamma and Foxp3 Regulate The Differentiation of Interleukin 17-Producing T Cells. Nat Immunol 9: 1297-306.



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