The Function of Retinoid X Receptor α in Cancer Cells

Gui-Li Huang[1], Qing-Xi Chen[1] and Dong-Yan Shen[2*]

[1]State Key Laboratory of Cellular Stress Biology, School of Life Sciences, Xiamen University, China

[2]Biobank, The First Affiliated Hospital of Xiamen University, China

*Corresponding author: Dong-Yan Shen, No. 55, Road of Zhenhai, Siming Zone, Xiamen 361003, China, Tel: +86-592-2137507; Fax: +86-592-2137509; E-mail: shendongyan@163.com

Abstract

The retinoid X receptor (RXR) is a member of the steroid/thyroid hormone superfamily of nuclear receptors (NRs) which are transcription factors that are essential in embryonic development, maintenance of differentiated phenotypes, metabolism and cell death. This review is to provide an overview of the mechanism of RXRα and RXRα signaling pathways involving RXR/TR3, RAR/RXR, PPAR/RXR, VDR/RXR, LXR/RXR, FXR/RXR in cancer cells and other diseases, which will enhance our ability to design rational therapeutic drugs for cancer. Recent studies have shown that an N-terminally truncated RXRα (tRXRα) exists in several cancer cell lines and primary tumors, which is considered as a kind of oncoprotein, demonstrating the new suitability of targeting tRXRα for cancer therapy.

Keywords: Retinoid X receptor α; Cancer; Cancer therapy

Introduction

Nuclear receptors are a class of transcription factors that can directly bind to DNA and modulate target gene transcription. Nuclear receptors play key roles in reproduction, development, and homeostasis of organisms [1-3]. On the basis of their ligands, nuclear receptors are classified into three families. The first is the classic and most extensively characterized group, steroid- and thyroid-hormone receptors [1], including retinoid nuclear receptors. The second class is the orphan nuclear receptors, which are structurally related to nuclear hormone receptors but for which no ligand has yet been discovered. The third class of nuclear receptors is adopted orphan nuclear receptors, whose regulation has been shown to range from true ligand-independence to highly promiscuous ligand-dependence. The retinoid receptor subfamily contains two classes, namely, the retinoic acid receptors (RARs) and retinoid X receptors (RXRs). Each class consists of three subtypes (α, β and γ) [4].

The structure of nuclear receptors is similar despite wide variation in ligand sensitivity. With few exceptions, they contain an NH2-terminal region (also known as the A/B region) that harbors a transactivation domain (AF-1); a core DNA-binding domain (the C region), also contains a dimerization interface that determines target gene specificity [5-7], containing two highly conserved zinc finger motifs that are common to the entire family except for dosage-specific sex reversal-adrenal hypoplasia congenita critical region on the X chromosome-1 (DAX1) and short heterodimeric partner (SHP) [8]; a hinge region (also named D region) that permits protein flexibility to allow for simultaneous receptor dimerization and DNA binding; and the E region, ligand-binding domain (LBD), contains a dimerization interface, and a ligand-dependent activation function (AF-2),The rest part is a variable F region whose entire function has not been known so far (Figure 1).
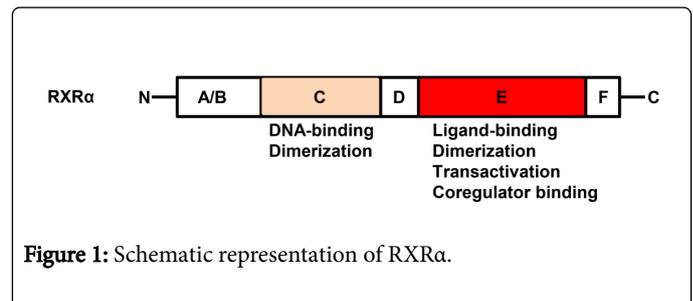

Figure 1: Schematic representation of RXRα.

Nuclear receptors are major targets for drug discovery and have key roles in development and homeostasis, as well as in many diseases such as cancer [8]. Retinoid X receptor α (RXRα) are becoming increasingly appreciated not simply as silent heterodimerization partners of other NRs, but also as therapeutic targets for cancer therapy and prevention by interacting with its ligands and several related signaling pathways. RXRα plays a role in many physiological processes including carcinogenesis [9]. Several RXRα ligands containing 9-cis-Retinoic acid (9-cis-RA), Targretin and the NSAID Etodolac and Sulindac could bind to RXRα to regulate different biological functions. One case of Retinoids' cancer therapy is that the effects of retinoid-based "differentiation therapy" have been impressively shown in the case of Acute Promyelocytic Leukaemia (APL) [10]. Genetic data also indicate that RXRα are involved in the chemopreventive activity of RA in experimental skin carcinogenesis [10]. It has been reported that targretin, a synthetic RXRα ligand, the major side effect of which is the induction of hypertriglyceridaemia, is recently used for treating persistent or refractory cutaneous T cell lymphoma [9,11,12], indicating the possibility of targeting RXRα for cancer therapy (Table 1).

Cell type	Available/Possible Ligands)	Treatments(RXRα
Acute Promyelocytic Leukaemia	9-cis-Retinoic acid	

kin carcinogenesis	9-cis-Retinoic acid
cutaneous T cell lymphoma	targretin

Table 1: Specific cancers and available/possible treatments targeting RXRα.

RXRα plays a central role in the regulation of many intracellular receptor signaling pathways and can mediate ligand-dependent transcription. RXRα enhances human cholangiocarcinoma growth via simultaneous activation of Wnt/β-catenin and NF-κB pathways [13]. The expression of the dominant-negative RXR affected the expression levels of a number of genes, some of which have been implicated in transcription, signal transduction, protein synthesis and protein trafficking [14]. The features of hepatocyte RXRα deletion in mice are that genes related to angiogenesis (Nos3, Kdr) were down-regulated, which leads to inhibition of angiogenesis, whereas genes connected with adipogenesis (Cebpb, Srebf1), pro-inflammatory pathway (NF-κB, TNFα) and apoptosis (Gzmb, Bcl-2) were up-regulated [15]. RXRα is known to heterodimerise with a number of nuclear receptors including TR, RAR, PPAR, VDR, LXR, FXR and several orphan receptors [2,16,17]. So, RXRα ligands (agonists and antagonists) have the potential to affect the signaling of numerous other pathways. Many recent reviews have described the mechanisms of RXRα as heterodimerization partners in various circumstances including development, metabolic diseases, and cancer [18-25]. Here we will discuss the latest insights into these various mechanisms how RXRα interacts with other nuclear receptors in cancer and other diseases in order to develop improved target-based drugs for cancer and other disease therapy.

RXRα Regulates TR3-Dependent Apoptosis by Modulating Its Nuclear Export and Mitochondrial Targeting

TR3 (also known as Nur77 or NGFI-B), an orphan member of the nuclear receptor superfamily [26-28], is an immediate-early response gene whose expression is rapidly induced in response to a variety of extracellular stimuli [29], including growth factors, the phorbol ester 12-O-tetradecanoyl-13-phorbol acetate (TPA) and cyclic-AMP-dependent pathways. The expression of TR3 is rapidly induced during apoptosis of immature thymocytes, T-cell hybridomas and various cancer cell types [28,30-34]. The apoptosis-associated translocation of TR3 from the nucleus to the cytoplasm has been observed in a lot of cancer cells such as lung cancer, ovarian cancer, colon cancer, gastric carcinoma and breast cancer [35-46]. RXRα is essential for nuclear export and mitochondrial targeting of TR3 through their dimerization interfaces located in their DNA-binding domain [43]. A nuclear export sequence (NES) present in RXRα's carboxyl-terminal region is required for the efficient nuclear export of RXRα/TR3 heterodimers [43]. RXRα has two dimerization interfaces, which are located in the DBD and the LBD [5,29]. The formation of the RXRα/TR3 heterodimer is mediated by dimerization interfaces in their DBDs, suggesting that the RXRα NES situating in the LBD is in its active conformation, resulting in the RXRα/TR3 heterodimer nuclear export [43]. Here RXRα, acting as a helper factor, facilitates the translocation of TR3 from the nucleus to the mitochondria, inducing cytochrome c release and cell apoptosis (Figure 2).
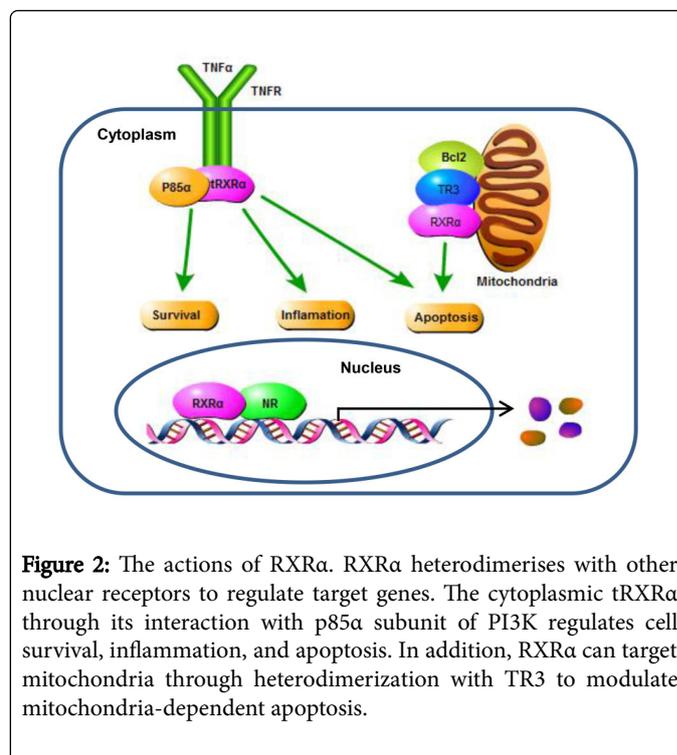

Figure 2: The actions of RXRα. RXRα heterodimerises with other nuclear receptors to regulate target genes. The cytoplasmic tRXRα through its interaction with p85α subunit of PI3K regulates cell survival, inflammation, and apoptosis. In addition, RXRα can target mitochondria through heterodimerization with TR3 to modulate mitochondria-dependent apoptosis.

RXRα ligands suppress apoptosis by preventing of TR3 and RXRα mitochondrial targeting. RXRα ligands 9-cis-RA and SR11237, for instance, effectively inhibited the release of cytochrome c induced by TPA or SR11453 in LNCaP cells [43]. Ligand binding can favor RXRα/Nur77 interaction to DNA binding and transactivation. The inhibition of RXRα/Nur77 DBD-mediated dimerization is related with the induction of heterodimer DNA binding and transactivation by 9-cis-RA. Ligand binding allows RXRα to interact with Nur77 through their LBD dimerization interfaces, silencing the RXRα NES. The nuclear export of the RXRα/TR3 heterodimer may be suppressed by 9-cis-RA through its induction RXRα homodimerization or modulation of RXRα/TR3 heterodimerization interfaces [43]. Accumulating evidence that RXRα ligands regulate apoptosis by modulating RXRα/TR3 heterodimer nuclear export in response to different apoptosis stimuli represents a novel approach for developing RXRα-based apoptosis regulators.

Therapeutic Applications of RAR/RXR Heterodimer Modulators in Cancer

Retinoic acid receptors (RARs) are ligand-inducible transcription factors that function as heterodimers with retinoid X receptors (RXRs) to regulate cell growth, differentiation, survival and death [47]. RXRs heterodimerizing with RARs in various tissues come into play mainly, if not exclusively [48]. These heterodimers have two distinct functions: First, they modulate the frequency of transcription initiation of target genes after binding to retinoic acid receptor response elements (RAREs) in their promoters; and second, they affect the efficiency of other signaling pathways ('crosstalk') [10]. RARs and RXRs form heterodimers, which are "non-permissive", that is unresponsive to RXR ligands on their own, but these agonists super activate transcription by synergizing with RAR ligands [21]. It is assumed that in these processes RAR heterodimerizes with RXR responding to RXR-selective

ligands, which are inactive alone, strengthening by RAR-selective ligands [49].

Having a better understanding of the biological role of RARs and RXRs is beneficial in the design of selective receptor modulators that might overcome the limitations of current drugs. The panRAR–RXR agonist, 9-cis-retinoic acid (9cRA), an active metabolite of vitamin A, has a higher affinity to RXRα [50]. PanRXR-agonists can induce higher-order RXRα/RARs fusion hetero-oligomeric oncogenic complexes aberrantly recruit transcriptional co-repressors to downstream targets which are essential for transformation in acute promyelocytic leukemia [51], suggesting the pathological significance of their potential value as a therapeutic target. RXRα/RAR heterodimers play a role in the retinoid-stimulated increase in steroid sulfatase activity which was blocked by pharmacological inhibition of the RAF-1 and ERK MAP kinases [52]. Accumulating evidence that ligand-induced promoter activity of RXRα/RAR heterodimer is significantly suppressed by high glucose (HG) which promoted protein destabilization and serine-phosphorylation of RAR and RXR is mediated through oxidative stress/JNK signaling [53]. The impaired RXRα/RAR signaling and oxidative stress/JNK pathway forms a vicious circle, which significantly contributes to cardiomyocyte apoptosis induced by hyperglycemia [53]. The RXRα/RAR signaling pathway plays critical roles in hippocampal synaptic plasticity and greatly contributes to memory performance, and long-term potentiation (LTP) in the hippocampus in the adult brain [54]. RXRα/RAR signaling pathway also improves axonal regeneration and modulate reactions of glia cells in physiological reactions after spinal cord injury [55]. So the modulators of RXRα/RAR promise to be a useful target after spinal cord or brain lesions. A large amount of RAR- and RXR-selective ligands have been designed and the corresponding structural and functional analyses have provided deep insight into the molecular basis of ligand action, which is useful for drug discovery.

Delineation of the molecular mechanisms that regulate RXR specification and function should be important for understanding a number of diseases. The post-maturation apoptosis of HL60 leukaemia cells requires both retinoids and rexinoids via RXRα/RAR signaling pathway [8]. Retinoids inhibit the progression stage during chemical skin carcinogenesis. Structural overlap of a retinoic acid response element with these retinoid X response elements led to a high affinity of RXRα/RAR heterodimer to the retinoic acid response element in the p21 promoter, resulting in the prevention of RXR ligand-mediated p21 transactivation whose up-regulation facilitated G(1) arrest [56]. Recent studies revealed that LG1506, a selective RXR modulator, had a distinct mechanism of action in that it facilitated the co-repressors recruit to the RXRα/RAR heterodimer complex at target gene promoters, inhibiting the differentiation of hematopoietic stem cells (HSCs) in culture [57]. So, studies on the molecular basis and selectivity of the RXRα/RAR complexes that modulate various events during tumorigenesis, and their effect on differentiation and apoptogenic pathways, might provide ideas about promising avenues for efficacious cancer therapies. Further investigations will clarify the RXRα/RAR-dependent antitumor activity or the receptor-independent anticancer action.

Modulation of Permissive PPAR/RXR Heterodimers

Peroxisome proliferator-activated receptors (PPARs) are ligand-dependent transcription factors, which can regulate gene expression by binding to peroxisome proliferator-responsive element (PPRE) located in the promoter region of their target genes as heterodimers with the RXRs after ligand binding. PPARs that are involved in the regulation of energy homeostasis have recently drawn much attention as therapeutic targets. PPARs are comprised of three closely related isotypes (α, β/δ and γ), which are encoded by different genes. Recent studies have shown that PPARγ agonists can regulate differentiation and induce growth arrest and apoptosis in a variety of cancer types [58,59], which require RXRs as an obligate heterodimeric partner [60].

RXRα forms a permissive heterodimeric complex with PPARγ which activates PPAR regulated gene expression [61-63]. Thiazolidinediones (TZDs), which is one of the most important PPARγ agonists, inhibited cell proliferation of human bladder carcinoma cell lines by increasing cyclin-dependent kinase inhibitor expression and induced cell death [64], which also needs RXRα co-expressed. It has been reported that a number of combinations of the RXRα agonists with the PPARγ agonists are useful for treating various cancers. The combined treatment with the PPARγ ligand Rosiglitazone (BRL) and the RXR ligand 9-cis retinoic acid (9cRA) induces human breast cancer cells apoptosis [65]. The combination of the PPARγ ligand ciglitazone and the RXRα ligand 9-cis-retinoic acid (9cRA) by activation of the RXRα/PPARγ heterodimer is useful on inhibition of cell growth of human colon cancer cells [66]. The combination of rexinoids (synthetic retinoids specific for RXR) with PPAR ligands may enhance the antiproliferative effects [67], arguing for the evaluation of combination therapies. The combination of RXR agonists rexinoids and PPARγ agonists thiazolidinedione (TZD) represents novel therapeutic targets in melanoma [68]. The combination of the RXRα agonist, bexarotene, with the PPARγ agonist, rosiglitazone, has greater efficacy in growth inhibition than either single agent in colon cancer [69], suggesting a potential role for utilizing a combination regimen of an RXRα and PPARγ agonist for colon cancer. Combination therapies' advantage is that one drug may reverse a cancer-selective block of an antiproliferative signaling pathway, thus allowing the second to become active in cells that are otherwise resistant [70,71]. Therefore, the combined use of RXRα and PPARγ ligands may offer therapeutic strategies in the treatment of cancer.

Beyond the treatment of a variety of cancers, targeting RXRα/PPARγ heterodimer opens the way to novel therapeutic opportunities of other diseases. PPAR and RXRα play crucial role in transcription regulation of inflammation response. The expression of PPARγ and RXRα which have been recognized as crucial players in the pathogenesis of atherosclerosis was associated with a more pronounced disease progression in patients with advanced carotid atherosclerotic lesions [72]. More targeted modulation of function through ligands design has been proposed as a strategy to possibly overcome observed rexinoids side effects that have limited the use of these compounds in the treatment of metabolic diseases [21,73]. It is exciting options that the therapy of metabolic diseases originating from the synthesis of heterodimer-selective rexinoids for PPARγ/RXRα combining with PPARγ agonists. It also deserves to be mentioned that co-administration of a rexinoid with a TZD produces enhance anti-diabetic activity without the increase in triglycerides associated with rexinoid administration [74-76]. Anti-inflammatory properties of targeting PPARγ/RXRα are well documented in the periphery [55].

Further studies are required to investigate the molecular mechanisms by which PPARγ/RXRα heterodimer ligands inhibit cell growth and induce differentiation. Extension of these results into the clinic may provide an opportunity to find a potentiated treatment effect of cancer, as well as other diseases, produced by the combination of PPARγ ligands with RXRα ligands.

RXR Dominates the Nuclear Import and Export of the Unliganded Vitamin D Receptor

Liganded and unliganded vitamin D receptors (VDRs) carry out distinct functions, both of which require heterodimerization with retinoid X receptors (RXRs) [77]. Ligand-dependent functions of VDRs heterodimerizing with RXR regulate calcium homeostasis, immune functions, endocrine functions, vitamin D metabolism, and cellular differentiation [77]. Ligand binding induces conformational changes in the VDR, which promote heterodimerization with retinoid X receptor (RXR) and recruitment of a number of nuclear receptor co-activator proteins [78]. Ligand-independent activation of VDR/RXR heterodimers activates a reporter driven by the prolactin promoter, which is in the presence of Ets-1 that induces a conformational change in the receptor, which creates an active interaction surface with co-activators even in the AF2-defective mutants [79].

Many nuclear proteins shuttle between the cytoplasm and the nucleus. The steady-state nuclear localization of RXR and liganded VDR is mediated by means of the molecular mechanisms that the receptors shuttle between cytoplasm and nucleus, but the residence time in the nucleus is longer than in the cytoplasm [77]. After synthesis in the cytoplasm the import process of RXRs which can be promoted by liganded VDR is dependent on a nuclear localization sequence (NLS) in the DNA-binding domain (DBD) of RXR. RXRα increases the ligand-independent nuclear import of VDR. Both RXRα and VDR export require proteins of the export machinery. Unliganded VDR distributed evenly between the cytoplasm and the nucleus. VDR/RXR heterodimers through dimerization interfaces in the DNA-binding domain (DBD) of RXRα are formed in the cytoplasm and translocate together to the nucleus upon calcitriol (VDR ligand) binding [77].

In view of the above molecular mechanism that RXR dominates the nuclear import and export of the unliganded vitamin D receptor, it's worth noticing that the VDR/RXR heterodimer also plays a role in cancer. The effect that VDR ligand 1,25-dihydroxyvitamin D3 (1,25(OH)2D3) and its non-hypercalcemic analog, EB1089, decrease parathyroid hormone-related protein (PTHrP) mRNA and cellular protein levels which increase the growth and osteolytic potential of prostate cancer cells, is mediated via a negative Vitamin D response element (nVDRE) within the human PTHrP gene and involves an interaction between nVDREhPTHrP and the Vitamin D receptor (VDR) [80]. RXRα which is a frequent heterodimeric partner of the VDR forms part of the nuclear protein complex that interacts with nVDREhPTHrP along with the VDR in prostate cancer cells [80].

The Interaction of RXR and Other Nuclear Receptor

As the promiscuous partner of heterodimeric associations, RXRs play a key role within the Nuclear Receptor (NR) superfamily. The RXR contributes to the regulation of diverse biological pathways via its role as a heterodimeric partner of several nuclear receptors. Steroid and xenobiotic receptor (SXR) dimerizes with retinoid X receptor (RXR) and regulates the transcription of genes encoding xenobiotic-metabolizing enzymes such as CYP3A4, which is activated by retinoids [81]. The human pregnane X receptor (hPXR) is an orphan nuclear receptor that binds to its response elements present in steroid-inducible cytochrome P-450 gene promoters, which requires the participation of RXRα [82]. It has been reported that a nuclear location of both hPXR and RXR in infiltrative breast cancer which is associated with an increased risk of recurrent disease [82]. Rexinoid bexarotene, a clinically used antitumoral agent, modulates triglycerides metabolism

in plasma whose increase, the most frequent side-effect, is an independent risk factor of cardiovascular disease, but not cholesterol metabolism via a selective permissivity on target genes of the RXRα/LXR heterodimer in the liver [83]. The antitumoral agent bexarotene (Targretin, Bexarotene) regulates target genes by binding to the nuclear RXRα. The evidence has proven a favorable pharmacological effect of bexarotene on atherosclerosis despite the induction of hypertriglyceridemia, likely via a beneficial action on intestinal absorption and macrophage efflux [84]. RXRα/LXR heterodimer might contribute to the beneficial effects of rexinoids on atherosclerosis and warrant further evaluation of RXRα/LXR agonists in prevention and treatment of atherosclerosis. RXRα function as heterodimers with liver X receptors (LXRs), which are involved in glucose/lipid metabolism. All these findings indicate that RXRα is central to the regulation of many important physiological functions in the organism. This expands the number of possible pharmaceutical targets for intervention with RXRα agonists or antagonists.

Regulation of tRXRα Production and Its Function

RXRα regulates diverse biological functions. Except its well-known action in the nucleus, RXRα also have extranuclear actions. RXRα exists in the cytoplasm at different stages during development in certain cell types [85]. In response to differentiation [86], inflammation [87,88] and apoptosis [43], RXRα transfers from the nucleus to the cytoplasm. The truncated RXRα (tRXRα) proteins, only existing in the cytoplasm, are produced through limited proteolytic cleavage of RXRα in cancer cells [89,90]. The cytoplasmic fraction but not in the nucleus [89] was shown to contain proteases like cathepsin L-type protease and m-calpain [89-93] that cleaves RXRα at its amino terminus. The truncated RXRα is produced in tumor tissues but not in normal tissues [94] is in line with other findings that RXRα is cleaved in tumor but not in premalignant or normal tissues from patients with malignant human prostatic tumor [90] or thyroid cancer [93]. The 54 kDa full-length RXRα (fl-RXRα) protein level is often reduced in cancer cells and tumor tissues [93,95], which is in part due to limited proteolytic processing of RXRα. Proteolytic processing of RXRα is an important mechanism in the regulation of the phosphatidylinositol-3-OH kinase (PI3K)/Akt signaling pathway and provides its potential value as a therapeutic target.

The N-terminally truncated 44 kDa RXRα protein and the 54 kDa fl-RXRα protein play significant roles in various tissues with different effects. The extensive cytoplasmic tRXRα interacts with the p85α subunit of phosphatidylinositol-3-OH kinase (PI3K) to activate the PI3K/AKT survival pathway and induce anchorage-independent cell growth in vitro and tumor growth in animals (Figure 2) [94], conforming that tRXRα provide a therapeutic advantage in cancer treatment. Due to deletion of the N-terminal sequences, RXRα in several cancer cell lines and primary tumors confers its ability to interact with p85α. The p85α-binding motifs in RXRα are probably masked by the N-terminal end sequences. The region, amino acids from 80 to 100 in RXRα critical for tRXRα binding to p85α [94], is enriched with proline resides, can presumably form several polyproline helices (PPII helix) known to bind to the SH3 domain [96] present in p85α. Cleavage of RXRα may represent a mechanism that triggers tRXRα signaling by removing the inhibitory N-terminal domain, allowing tRXRα to expose its p85α-binding motif. The tRXRα detected in the cytoplasm of cancer cells to modulate carcinogenesis acts nongenomically to activate the PI3K/AKT pathway to promote cancer cell growth and survival [94]. Hence, agents targeting tRXRα-mediated

pathway can be effective. Nonsteroidal anti-inflammatory drugs (NSAIDs) sulindac could inhibit the tRXRα-dependent PI3K/AKT activation [94], suggesting that Sulindac stands for a type of anticancer drugs targeting this pathway. The tRXRα was critical for AKT activation by TNFα that could also activate PI3K/AKT signaling [97,98]. Transfection of RXRα siRNA, which inhibited both the expression of the fl-RXRα and the 44 kDa tRXRα, significantly impaired the ability of TNFα to activate AKT [94]. Sulindac can lower tRXRα-mediated activities that tRXRα contributes to the growth and survival of cancer cells by activating AKT, suggesting that tRXRα serves as an intracellular target mediating the apoptotic effect of Sulindac. The fact that sulindac and TNFα synergistically reduce tRXRα-mediated AKT activation [94] provides new understanding of the crosstalk between retinoid receptor and TNFα signaling pathways, implying the further tRXRα-based development for cancer therapy.

Conclusion

RXRs are obligatory DNA-binding partners for a number of nuclear receptors, broadening the spectrum of their biological activity to the corresponding nuclear receptor-signaling pathways. Unliganded RXRα self-associates into tetramers and that each dimer within these tetramers can separately bind to an RXRα response element which may bring about distant genomic effect. Ligand binding induces the dissociation of RXRα tetramers into dimers, which can alter gene expression by modulating the DNA structure. Many other nuclear receptors require RXRα as heterodimerization partner for their function. This places RXRα in the crossroad of multiple distinct biological pathways. The emerging roles of RXRα in the RXRα signaling network and possible implications are helpful for our understanding of nuclear receptor biology and pharmacology. Thus the multiple roles that RXRα plays in all kinds of cells have turned RXRα into an attractive drug target. The priority is to change a tumor-promoting microenvironment to a tumor-inhibiting state and to understand the signaling mechanisms involved, finding a potential target for cancer prevention.

The unique property of RXRα dimerization interfaces allows cross talk among RXRα heterodimerization partners with respect to their subcellular localization and function. The RAR/RXR and PPAR/RXR signaling pathways have recently been implicated in the progression of neurodegenerative and psychiatric diseases, suggesting that the activation of PPAR/RXR and RAR/RXR transcription factors has been proposed as a therapeutic strategy in disorders of the central nervous system [99]. Overexpression of dominant negative RAR mutants may lead to repression of genes that are not normally targeted by co-repressor-associated unliganded RAR/RXR heterodimers [100] and/or interfere with functions of other RXR heterodimeric partners through sequestration of RXRs. RXR-selective ligands (rexinoids) are valuable in the treatment of atherosclerosis, other cardiovascular indications and inflammatory diseases via pathways including the PPARs, the liver X receptors and the farnesoid X receptors [101-103]. RXRα ligands are attractive candidates for clinical application because of their activity against tamoxifen-resistant breast cancer, taxol-resistant lung cancer, metabolic syndrome, and allergy. This then led to investigation of the mechanism in which these compounds inhibited growth and the effects of combination treatment on essential cell growth and differentiation parameters in cancer. A detailed understanding of the multiple physiological effects elicited by various ligands through nuclear receptors is obviously good for drug discovery.

RXRα has been implicated in several neoplastic diseases. Ligands that activate the nuclear RXRα display potent anti-carcinogenic activities through the mechanisms by which these ligands inhibit carcinoma cell growth and promote apoptosis. The case that RXRα ligands inhibit mammary carcinoma cell growth stems from the ability of these ligands to regulate the state of RXRα and is independent of the direct intrinsic transcriptional activity of the receptor [104]. Some combined compounds that target RAR/RXR, RXR/TR3, PPAR/RXR, VDR/RXR, RXR/LXR, RXR/FXR, etc. heterodimers are powerful anticancer drugs. An improved understanding of the mechanism of these heterodimers pathway should enable the rational design of more selective modulators in general. Combinatorial treatments might lead to synergistic effects on growth control or induction of apoptosis, thereby allowing the use of lower concentrations as well as maintaining efficacy and reducing side effects. It has been reported that RXRα ligands such as 9-cis-RA, Targretin, Etodola and Sulindac play a significant role in a lot of cancer therapy, whether alone or combing with its partners' ligands. RXRα and its partner present main targets for pharmacologic interventions, allowing development of therapies targeting different receptors with high efficiency. The combined use of several compounds that act on different signalling pathways also represents an interesting approach in curing cancer. The combination of retinoids and T-cell-based immunotherapy has efficacy in neuroblastoma. The reason why addition of the HDAC inhibitor sodium phenyl butyrate in the treatment of a patient with multiple relapsed RAresistant APL resulted in complete remission [105], is that HDAC inhibitor sensitized the RAinsensitive cells to the differentiative action of RA by restoring retinoid signaling [70].

RXRα plays a central role in controlling multiple hormonal pathways through heterodimerization. Despite their promiscuity in heterodimer formation and activation of multiple pathways, RXRα is a target for drug discovery. Recent studies have shed light on the molecular mechanisms underlying tRXRα action, which has made it possible to design appropriate modulators for cancer therapy. Further studies are required to understand the regulation of tRXRα in all kinds of tissues in order to have far insight in the pathological function of tRXRα. It has been shown that the presence of tRXRα in breast and liver cancer tissues but not in tumor surrounding tissues or distant normal tissues from the same patients. The RXRα-selective Sulindac derivative K-80003 could effectively inhibit the tRXRα pathway and the growth of cancer cells in vitro and in animals, providing an important new treatment for cancer patients.

The accumulated knowledge of the mechanistic, molecular and pharmacological actions of RXRα ligands is the basis for the development of efficient anticancer therapies. Indeed, it has been shown that retinoid agonists can autonomously induce rapid apoptosis under certain conditions [106]. RXRα is an attractive molecular target for drug development. The rational drug design can develop new RXRα selective ligands rexinoids with improved biological properties, warranting further development for cancer therapy. This can be achieved only by interdisciplinary efforts that combine in vivo analysis using genetically engineered animals with in vitro cell and molecular biological analyses to elucidate the detailed mechanisms that drugs promote cancer cells apoptosis by RXRα signaling pathways. This unique class of RXR ligands will provide a means to control distinct target genes at the level of transcription and allow the development of retinoids with a new pharmacological action.

Additional mechanisms to regulate RXRα might also exist. It will be interesting to examine the significance of RXRα signaling pathway. The

complete understanding of RXRα-dependent mechanism and drugs design targeting RXRα signaling for various diseases is therefore a major challenge for future research. There is no doubt that RXRα is important to cancer biology and treatment in the 21st century. It is going to continue investigating the possible involvement of various RXRα regulators in cancer cells and the molecular mechanisms that are involved. Such studies will provide useful information for the design of therapeutic drugs. Obviously, studies on the regulation of RXRα signaling will remain both challenging and exciting in years to come.

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