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## Nuclear receptors in osteoclasts

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### Abstract

Osteoclasts are bone-resorbing cells that play an essential role in the remodeling of bone under physiological conditions and numerous pathological conditions, such as osteoporosis, bone metastasis, and inflammatory bone erosion. Nuclear receptors are crucial to various physiological processes, including metabolism, development and inflammation, and function as transcription factors to activate target genes. Synthetic ligands of nuclear receptors are also available for the treatment of metabolic and inflammatory diseases. However, dysregulated bone phenotypes have been documented in patients who take synthetic nuclear receptor ligands as a therapy. Therefore, the effect of nuclear receptors on bone cells has become an important area of exploration; additionally, the molecular mechanisms underlying the action of nuclear receptors in osteoclasts have not been completely understood. Here, we cover the recent progress in our understanding of the roles of nuclear receptors in osteoclasts.

### Introduction

Nuclear receptors are a protein family of transcription factors, activated by ligand binding [1–3]. The body expresses more than 48 different nuclear receptors (NRs), officially divided into six groups based on phylogeny [4,5]. Group 1 includes the receptors TRs, RARs, VDR, and PPARs, RORs, Rev-erbs, CAR, PXR, LXRs, and others. Group 2 contains RXRs, COUP-TF, and HNF-4. Group 3 includes the steroid receptors (ERs, GRs, PRs, and ARs) as well as the ERRs. Group 4 contains the nerve growth factor-induced clone B group of orphan receptors including NGFI-B, NURR1, and NOR1. Group 5 includes the steroidogenic factor 1 and the receptors related to the *Drosophila* FTZ-F1. Group 6 contains

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only the GCNF1 receptor, which does not fit well into any other subfamilies. NRs signal in both genomic and non-genomic ways (Figure 1). Ligand binding to NRs facilitates conformational changes promoting DNA binding to NR/ligand complexes and allowing for interactions with transcription factors. Ultimately, NRs regulate target gene expression through a consensus DNA binding site. However, recent findings suggest that NRs bind to other genomic sequences by interacting with additional transcription factors [6]. NRs also function via non-genomic effects, including regulation of protein activity, modulation of ion channels and intracellular calcium levels, and production of secondary messengers [7].

Nuclear receptors show notable structural similarity [1–3]. Most members of the nuclear receptor group share a general structure (Figure 2). A-B is the N-terminal domain containing activation function-1 (AF-1) and is variable in size and sequence. C has the highly conserved DNA binding domain (DBD), and D is the hinge domain. E has the Ligand binding domain (LBD) that contains AF-2, which is important for the binding of coactivators. Ligand binding to LBD causes the conformational change of NRs. F is the highly variable C-terminal domain. The function of AF-1 is independent of ligand binding and AF-1 alone leads to a weak transcriptional activation. In contrast, the action of AF-2 is dependent on ligands and co-activation of AF-1 with AF-2 promotes a strong transcriptional activation of target genes. In the absence of ligand, nuclear receptor corepressors such as SMRT and NcoR bind to NRs to repress the basal transcription of the target genes. Although corepressors bind to corepressor NR boxes in the LBD, their binding is independent of AF-2. Ligand binding facilitates the conformational changes of AF-2 to trigger the release of corepressor and to induce the subsequent interaction with nuclear receptor coactivators such as SRC1 and TIF2. The deletion of AF-2 helix enhances the binding of corepressors to some NRs such as VDR. Moreover, AF-2 dependent recruitment of coactivators enhances the binding of transcriptional machinery to NRs and facilitates transcriptional activation of the target genes.

Osteoclasts are multinucleated, bone-resorbing cells derived from myeloid lineage cells [8–10]. Osteoclastogenesis progresses in three different phases; commitment, fusion/maturation, and resorption [8] (Figure 3). Macrophage-colony stimulatory factor (M-CSF) and receptor activator of nuclear factor  $\kappa$ B ligand (RANKL) are essential cytokines for osteoclast differentiation. M-CSF is a crucial factor for function, survival, and differentiation of monocytes/macrophages and osteoclasts [11,12]. Engagement of RANKL to its receptor, RANK, activates multiple signaling pathways, including p38, ERK, JNK, AKT, and NF- $\kappa$ B [13], and this induces expression of osteoclast transcription factors, such as c-FOS and MYC [8]. With ITAM-mediated signals, RANKL induces and activates NFATc1, initiating osteoclast differentiation and promoting cell-cell fusion to generate multinuclear osteoclasts by induction of the membrane fusion promoter, DC-STAMP, and other fusion genes. During osteoclastogenesis, osteoclasts also undergo cytoskeletal remodeling to migrate and attach to the bone surface via the PI3K dependent mechanism [14]. In early phase of osteoclastogenesis, columnar actin puncta, known as podosomes, starts to form and is organized as actin ring as osteoclastogenesis progresses [15]. Fusion of mononuclear preosteoclasts occurs in a site-specific manner [16]. Mature osteoclasts are able to resorb bone by secreting bone resorbing factors. Osteoclastogenesis is regulated by RANKL and various factors including inflammatory mediators, PTH, and ligands of NRs such as

hormones and vitamin D. Ostoprotegrin (OPG) is a decoy receptor for RANKL and is secreted from stromal cells and osteoblasts [17]. OPG prevents RANKL and RANK interactions and the ratio of RANKL-to OPG correlates with *in vivo* osteoclastogenesis and fine-tune osteoclast differentiation and bone remodeling.

Multiple NRs are expressed in osteoclasts [18,19] (Table1). It is established that NRs play a critical role in bone homeostasis and remodeling [18,19]. NRs affect skeleton size and shape during development and maintain skeletal homeostasis through both direct and indirect regulation of osteoclast differentiation and activity. Activation of NRs directly regulates the expression of target genes in osteoclasts. NRs indirectly influence osteoclastogenesis by modulating the RANKL/OPG ratio. Ligands for many NRs have not been discovered to date, and these NRs are often named orphan NRs based on their structural domain properties. Adopted NRs emerge when their endogenous ligands are characterized[20]. Many synthetic ligands for NRs have been developed, and pharmacological activation of NR with specific agonists has been used for the treatment of metabolic and inflammatory diseases. However, they are often associated with abnormal skeletal phenotypes and increased risk of fracture [21], and thus, it is important to understand the underlying mechanism of how NRs regulate osteoclast formation and function. In this review, we provide an update on six selected NRs with profound effects on osteoclasts and whose synthetic ligands are used in clinical treatments; additionally, the effects of other NRs on osteoclasts are reviewed in [18,19,22].

## Estrogen receptor (ER)

Estrogen plays an important role in reproductive organ development, energy metabolism and bone homeostasis [23,24]. In post-menopausal women, estrogen is diminished, and estrogen treatment prevents bone loss [25]. SERM (selective estrogen receptor modulators) are FDA approved synthetic estrogen analogs that are also used to prevent bone loss [26]. ERs are major mediators for estrogen actions and have two isoforms; ER $\alpha$  and ER $\beta$ . ER $\alpha$  regulates gene expression in a tissue and cell typespecific manner through the coordinated actions of transcription factors and co-factors. Following estrogen binding, activated nuclear ERs bind to genomic DNA, containing an estrogen response element (ERE), and co-regulators are recruited to modulate ER-mediated gene transcription.

Estrogen-deficiency in postmenopausal women, ovariectomized women, and ER $\alpha$ -deficient men all have diminished cortical and trabecular bone mineral density and enhanced osteoclast-mediated bone resorption [27,28]. In contrast, female ER $\alpha$ -deficient mice show increased bone mass with low bone turnover, and ER $\beta$ -deficient mice exhibit a milder bone phenotype compared to ER $\alpha$ -deficient mice [29,30]. In addition, ovariectomized female ER $\alpha$ -deficient mice show decreased bone mass. The discrepancy in bone phenotype for female global ER $\alpha$ -deficient mice can be explained through elevated circulating testosterone and androgen signaling. However, ER $\alpha^{f/f}$ -LysozymeM cre mice, carrying a conditional deletion of ER $\alpha$  in osteoclast precursor cells, show increased osteoclast number and decreased trabecular bone mass [31]. ER $\alpha^{f/f}$ -CathepsinK cre mice, resulting in the deletion of ER $\alpha$  in mature osteoclasts, exhibit decreased trabecular bone mass only in females [32].

These results suggest that activation of ERs in osteoclasts negatively regulates osteoclast differentiation and bone resorption.

Estrogen promotes osteoclast apoptosis and regulates osteoclast lifespan by inducing expression of FAS ligands (FASL)[32]. Activation of ER $\alpha$  also induces the destabilization of HIF1 $\alpha$  (hypoxia inducible factor) in osteoclasts [33]. The enhanced osteoclastogenesis in cells from ER $\alpha^{f/f}$ -CathepsinK cre mice is reversed by HIF1 $\alpha$ - deficiency. HIF1 $\alpha$  is stabilized and accumulated during osteoclastogenesis, positively regulating osteoclast differentiation[34]. Together, estrogen and SERM suppress the accumulation of HIF1 $\alpha$  in osteoclasts [35]. Estrogen indirectly regulates osteoclastogenesis by inducing FASL in osteoblasts and regulating the RANKL/OPG ratio [36]. ER $\alpha$  regulates gene expression in a ligand-independent manner. ER $\alpha$  regulates CREB phosphorylation by activating other kinases, such as mitogen activated protein kinase (MAPK), and controlling CREB-mediated gene regulation[37]. ER $\alpha$  also undergoes posttranslational modification through signaling pathways independent of ligands [38]. The action of ER $\alpha$  is regulated by growth factors and cytokines [39] and by competing with ERE-BP (estrogen response element-binding protein) for occupancy of ERRs. ERE-BP overexpression causes estrogen resistance [40] and enhances osteoclast differentiation[41]. Finally, myeloid-specific deletion of Sirt6 decreases ER $\alpha$  protein levels and apoptosis in pre-osteoclasts. Sirt6 deacetylates ER $\alpha$  protein, preventing further proteasome degradation [42]. The effect of ER $\beta$  on osteoclasts has not been characterized. Taken together, estrogen therapy would be beneficial for treating bone loss by suppressing osteoclasts and promoting osteoblast apoptosis. However, other cells also express ERs and estrogen therapy shows the adverse effect on other organs such as uterine and breast [43,44]. Importantly, SERMs such as roxifene is used to prevent or treat bone loss in women after menopause and functions as an estrogen blocker in uterine and breast to diminish the side effects of estrogen therapy.

## Estrogen related receptor (ERR)

ERRs are important in energy metabolism and cancer development and express in tissues with high-energy demands [45]. ERRs were discovered from their sequence similarity to DNA binding domains of ER $\alpha$  [46]. However, ERRs do not bind to estrogen. They contain three members; ERR $\alpha$ , ERR $\beta$ , and ERR $\gamma$  [46]. Estrogen-related receptors (ERRs) bind to the ERR response element (ERRE) and the ERE [46]. The ligands that interact with ERRs are unclear, so they are orphan NRs [47]. However, a recent study identified cholesterol as a potential endogenous ligand for ERR $\alpha$  [45].

ERR $\alpha$  regulates many key functions in osteoclasts. It is a positive regulator of osteoclast differentiation by mediating metabolic reprogramming [48,49]. ERR $\alpha$  also regulates metabolic genes controlling mitochondrial respiration in osteoclasts [49]. During osteoclastogenesis, MYC induces ERR $\alpha$  to regulate oxidative phosphorylation [49]. Several cofactors interact with ERRs, including peroxisome proliferator-activated receptor gamma coactivator 1-alpha (PGC1- $\alpha$ ) and PGC1- $\beta$  [50]. In osteoclasts, ERR $\alpha$  complexes with PGC1 $\beta$  upon RANKL stimulation to regulate target genes such as Ndg2, Aco2, and IDH3 [48]. Beyond regulating osteoclast functions, ERR $\alpha$  also affects pathological bone loss. Deletion or pharmacological inhibition of ERR $\alpha$  attenuates ovariectomy-induced bone loss

in mouse models of osteoporosis [49,51]. In addition, cholesterol, a natural  $ERR\alpha$  ligand, promotes osteoclastogenesis, and its effect on osteoclast differentiation was diminished in  $ERR\alpha$ -deficient mice [45].  $ERR\alpha$  is a potential therapeutic target in pathologic bone loss. In addition to  $ERR\alpha$  agonists and antagonists, cholesterol level modulation can regulate  $ERR\alpha$  activity. The physiological role of  $ERR\alpha$  in osteoclasts is not conclusive due to contradicting reports using global  $ERR\alpha$ -deficient mice. While an increase in osteoblast number, and a decrease in osteoclast number are reported in  $ERR\alpha$ -deficient mice by Wei et al[48], other studies only detect an increase in osteoblast number with no change in osteoclast number[51,52]. Bisphosphonates are an anti-resorptive drug and are widely used for suppressing osteoclast-mediated bone resorption[53]. Bisphosphonates inhibit two enzymes in the cholesterol biosynthesis pathway. Although the linking of the action of bisphosphonates to  $ERR\alpha$  is not well established, pharmacological inhibition of  $ERR\alpha$  can be useful treatment for pathological bone loss due to the dual effect of  $ERR\alpha$  on bone cells and on cholesterol-mediated actions in bone.

Compared to  $ERR\alpha$  less is known about  $ERR\beta$  and  $ERR\gamma$  in osteoclasts. To date, no study addresses  $ERR\beta$ 's role in osteoclasts.  $ERR\gamma$  is high in osteoclast precursor cells and decreased in osteoclasts. Overexpression of  $ERR\gamma$  suppresses osteoclast differentiation by inhibiting RANKL-induced NF- $\kappa$ B pathway [54], suggesting that  $ERR\gamma$  is a negative regulator of osteoclastogenesis. Administration of  $ERR\gamma$ -agonists also protects mice from pathological bone loss[54]. However,  $ERR\gamma$ -deficiency increases bone mass with no changes in osteoclasts [55], suggesting that the effect of  $ERR\gamma$  on bone remodeling is attributed to regulating osteoblast function under physiological conditions.

## Glucocorticoid receptor (GR)

Endogenous glucocorticoids (GCs) are essential for physiological and developmental processes [56]. Moreover, synthetic GCs are an effective therapy for autoimmune and inflammatory diseases. Both endogenous and synthetic GCs bind to GRs to regulate transcription of targets [57]. Human GRs are endocrine NRs, containing  $GR\alpha$  and  $GR\beta$ . Human  $GR\alpha$  is present in the cytoplasm of almost every cell. It is a ligand-dependent transcription factor, mediating many known actions of glucocorticoids. In contrast to  $GR\alpha$ , human  $GR\beta$  is in the nucleus and does not bind to GCs or activate GC-response genes [58].  $GR\beta$  is a dominant negative factor for  $GR\alpha$  and confers glucocorticoid resistance in inflammatory diseases [58]. The cellular response to GCs varies in sensitivity and specificity among individuals and within tissues of the same individual [59]. GR binds to the GRE (GC-responsive element) of targets to regulate gene expression by interacting with transcription factors [60].

Therapeutic use of GCs come with serious side effects, such as glucocorticoid-induced osteoporosis (GIO) [61,62]. GIO has been studied, focusing on the effects of GCs on osteogenesis. GCs block osteoblast differentiation and accelerate apoptosis of osteoblasts and osteocytes [63]. Several different mechanisms underlie the impaired osteogenesis by GCs have been suggested. GR-mediated suppression on osteogenic genes via regulation of chromatin remodeling complexes, BRG1 and BRM [64], suppressing IL-11 [65], and inducing WNT antagonists such as DKK1, sFRP1, and WIF [66–68]. In addition, GCs

reduce OPG and increase RANKL in osteocytes/osteoblasts, and an altered RANKL to OPG ratio by GCs promote osteoclast-mediated cortical bone resorption [69]. Although the indirect role of GCs in osteoclasts are well established, GCs' direct effect on osteoclasts is inconclusive [70]. GCs enhance osteoclast activity and differentiation *in vitro*, especially, when precursor cells were seeded in bone. These promoting effects of GCs decrease in dimerization-defective GR<sup>dim</sup> osteoclasts [71], showing that GC-mediated enhancement of osteoclastogenesis is GR-dependent. Other studies show that the administration of GCs suppresses osteoclast precursors, but osteoclast numbers remain constant due to their prolonged life span [72]. However, treatment with dexamethasone (DEX) suppresses osteoclast-spreading and bone resorption. DEX-mediated suppression of osteoclastogenesis is undetectable in GR $\alpha$ <sup>f/f</sup>-LysozymeM cre mice, a conditional deletion of GR $\alpha$  in osteoclast precursor cells [73]. Although one study showed that GCs regulate osteoclast activity by PKCII $\beta$ -mediated TRPV1 and CB2 regulation [74], the mechanisms underlying GR's activity under physiological and pathological conditions are not fully characterized. Glucocorticoid-induced bone loss is biphasic with a rapid early phase of bone loss (5~15% per year) followed by a constant low rate of bone loss (2% per year) [75]. Despite the controversial effects of GCs on osteoclasts, the timing and dosage of GCs generate the differential effect on osteoclastogenesis (Kaneko et al., unpublished observations), and the combinational effects of GCs that increase apoptosis of osteoblasts and osteocytes, increase adipocyte hypertrophy, decrease bone vasculature, and decrease osteoclast apoptosis might explain the clinical effects of GCs.

### Peroxisome Proliferator-Activator receptor (PPAR $\gamma$ )

The peroxisome proliferator-activated receptor (PPAR) family of nuclear receptors regulates lipid-related genes and functions as a metabolic sensor [76]. PPARs are activated by natural ligands, such as fatty acids [77]. They contain three members, PPAR $\alpha$ , PPAR $\beta/\delta$ , and PPAR $\gamma$ . PPAR $\delta$  and PPAR $\gamma$  express in human and murine osteoclasts, but PPAR $\alpha$  expresses only in human osteoclasts. PPARs form heterodimeric complexes with RXR (retinoid X receptor), and PPAR/RXR complexes bind to peroxisome proliferator response elements (PPRE) in the genome with different cofactors. PPAR $\gamma$  has been involved in regulation of the bone metabolism [78–80], but the role of other PPARs in osteoclast differentiation and function is not well characterized. PPAR $\gamma$  is originally identified as the master regulator of adipogenic differentiation [81]. Ligands of PPAR $\gamma$  are naturally occurring lipophilic ligands, such as polyunsaturated fatty acids and oxidized prostaglandin J<sub>2</sub>, as well as the synthetic agonist, TZDs [81,82]. PPAR $\alpha$ -deficient mice have no bone phenotype, but treatment with fenofibrate, a synthetic ligand for PPAR $\alpha$ , decreases osteoclasts, increases osteoblast differentiation [83], and it protects from OVX-induced bone loss in rats [84]. PPAR $\delta$  is a key regulator of fatty acid metabolism in muscle [85] and has been implicated as a regulator of the crosstalk between osteoblasts and osteoclasts [86]. The treatment of GW501516, an agonist of PPAR $\beta/\delta$ , promotes Wnt-dependent and  $\beta$ -catenin-dependent signaling in osteoblasts and the differentiation of osteoblasts [86]. Activation of PPAR $\beta/\delta$  by GW501516 suppresses osteoclastogenesis in an osteoblast dependent manner and also protects mice from OVX-induced bone loss by altering the OPG/RANKL ratio [86]. Conversely, PPAR $\delta$ -Sox2 cre mice exhibit osteopenic phenotype with diminished

osteoclast number *in vivo*. Although bone formation *in vivo* is not altered in PPAR $\delta$ -Sox2 cre mice, osteoblasts increase the expression of OPG to attenuate osteoclastogenesis. Therefore, PPAR $\delta$  regulates osteoclasts by controlling the crosstalk between osteoclasts and osteoblasts.

The thiazolidinedione (TZD) family is a synthetic agonist for PPAR $\gamma$  used to treat insulin-resistant type II diabetes, and both pioglitazone and rosiglitazone are the currently approved for clinical use. However, it has been shown that long-term usage of TZD increases fracture risk and bone loss [87,88] with higher incidence in women than men [89]. TZDs regulate bone metabolisms by supporting osteoclastogenesis [48,80] and suppressing osteoblastogenesis [78]. Rosiglitazone-fed mice show significant decrease in BMD, lower cortical thickness and trabecular bone volume [90,91]. TZDs change marrow structure and function showing a decrease in osteoblast number, an increase in marrow fat cells and osteoclast number [92]. Although TZDs decrease osteoblasts and induce osteocyte apoptosis, in aged animal, TZD-mediated bone loss is dominantly associated with an increased osteoclastogenesis by increased RANKL expression [92]. PGC1 $\beta$  is required for TZD-enhanced osteoclastogenesis and is induced by  $\beta$ -catenin in cells treated with rosiglitazone [48]. PGC1 $\beta$  is a co-activator for PPAR $\gamma$  activation of osteoclastogenesis and regulates osteoclastogenesis [93]. Both PGC1 $\beta$  and PPAR $\gamma$  bind to ERR $\alpha$ 's promoter to induce ERR $\alpha$  [48], suggesting several NRs cooperatively regulate osteoclast function. However, the physiological role of PPAR $\gamma$  in osteoclasts is controversial because the opposite bone phenotype of PPAR $\gamma$ - deficient mice has been observed from two different laboratories. Wan *et al* show that osteoclastogenesis and bone mass of PPAR $\gamma$ -Tie2cre mice decreased compared to control mice, and PPAR $\gamma$  promoted osteoclastogenesis by enhancing c-FOS expression [80]. In contrast, the Teitelbaum group demonstrated that various osteoclast-lineage cell specific deletion of PPAR $\gamma$  using Tie2cre, LysozymeM cre, Vav1 cre, and RANK cre mice (PPAR $\gamma$  cKO mice) all exhibits no changes in *in vitro* osteoclastogenesis and unaltered bone mass [94], and PPAR $\gamma$  cKO mice express similar levels of osteoclast-specific proteins including c-FOS, cathepsin K, and integrin  $\beta$ 3 compared to wild type mice. Furthermore, PPAR $\gamma$  deficiency cannot protect mice from OVX-induced bone loss [94], suggesting that PPAR $\gamma$  does not regulate physiological and pathological osteoclastogenesis. In other studies, ASXL2-deficient mice exhibit osteopenic phenotype with diminished osteoclast activity [95]. As ASXL2 is a coactivator of PPAR $\gamma$ , this data may also suggest the physiological role of PPAR $\gamma$  in osteoclastogenesis. Although these contradicting results argue that basal activation of PPAR $\gamma$  in early stages of osteoclast differentiation, it is clear that drug-induced activation of PPAR $\gamma$  promotes osteoclastogenesis, and thus, osteoclasts likely contribute to the increased fracture risk by TZDs. Therefore, it is important to understand the mechanisms how TZDs regulate osteoclast differentiation and activity.

## Liver X receptors (LXRs)

LXR $\alpha$  and LXR $\beta$  are important regulators of cholesterol, glucose metabolism, and inflammatory response [96]. While LXR $\beta$  is ubiquitously expressed, LXR $\alpha$  expression is limited to few tissues including liver, adipose tissue, intestines, lungs, and macrophages. LXR $\alpha$  and LXR $\beta$  proteins have considerable sequence homology and respond to the same

ligands. The endogenous ligands for LXR are sterol metabolites, including oxysterols [97,98]. LXRs form heterodimers with Retinoid X Receptor (RXR) and bind to LXR response elements (LXREs). Synthetic agonists including GW3965 and T0901317, inverse agonists such as SR9238, and antagonists such as GSK2033 for LXRs have been developed. Although bone mass is not altered in LXR $\alpha$ -, LXR $\beta$ -, or LXR $\alpha\beta$ -deficient mice, some bone parameters are changed in LXR-deficient mice-LXR $\alpha$ -deficient female mice that significantly increase total BMD and femur length at 16 weeks, and LXR $\beta$ -deficient male mice significantly decrease trabecular BMD at 16 weeks [99]. In addition, bone mass is not altered by the treatment with the synthetic LXR ligands GW3965 or T0901317 in mice [100]. However, activation of LXRs by GW3965 regulates both osteoclasts and osteoblasts, and it is well documented that LXR agonists are important players for osteoclastogenesis. Activation of LXRs inhibits osteoclast differentiation and activity by regulating MITF (microphthalmia-associated transcription factor)-NFATc1 expression via an LXR $\beta$ -dependent mechanism [101]. The treatment of GW3965 or T0901317 protects mice from OVX-induced bone loss and PTH-induced bone loss by decreasing the RANKL-to OPG ratio [102]. GW3965 treatment block osteoblast-induced osteoclast formation in osteoclast/osteoblast co-culture by regulating the RANKL to OPG ratio [102]. GW3965 also suppresses LPS-induced osteoclastogenesis by regulating Akt pathway via an LXR $\beta$ -dependent mechanism [103]. LXR agonists inhibit RANKL-induced c-Fos and NFATc1 expression and accelerate apoptosis in mature osteoclasts by the induction of caspase-3 and -9 activity and Bim expression [104]. LXR agonists have a minimal effect on physiological bone homeostasis, but activation of LXRs is useful for estrogen deficiency-induced bone loss or inflammatory bone loss. These results suggest that activation of LXRs would be beneficial for the treatment of pathological bone loss. However, in animal models, LXR agonists also increase fatty acid and triglyceride (TG) synthesis, which also affect osteoclast activity and function. Thus, both beneficial and adverse effects of LXR agonists in other cells need to be considered for the use of LXR agonists for bone diseases.

## Retinoic X receptor (RXR)

Retinoid X receptor (RXR) controls a wide variety of cellular processes such as lipid and glucose metabolism. RXRs consist of three subtypes- RXR $\alpha$ , RXR $\beta$ , and RXR $\gamma$ . RXR $\alpha$  and RXR $\beta$  express in osteoclasts and myeloid cells [105]. RXRs form either homodimers or heterodimers with other nuclear receptors, including LXRs, VDR, or PPARs [106,107]. RXR $\alpha\beta$ -Mx1cre mice (RXR KO), which have deletion of RXR in any IFN-responsible cells including osteoclasts and osteoblasts, exhibit increased cortical and trabecular bone mass, but the number of osteoclasts remains the same [108]. RXR-deficient osteoclasts form bigger osteoclasts with low bone resorption activity than wild type osteoclasts and express low MAFB, resulting in altered proliferation of osteoclasts. Paradoxically, both RXR-deficiency as well as pharmacological activation of RXR by bexarotene protect mice from OVX-induced bone loss. This protective role of RXR activation in OVX-induced bone loss can be explained by altered MAFB expression. While RXR homodimers can directly bind to the promoter of *Mafb* to induce MAFB expression, pharmacological activation of RXR leads to the formation of RXR/LXR heterodimers that indirectly increase MAFB by inducing SREBP1c expression. Bexarotene is the only approved RXR agonist for the clinical use and



has been used as a treatment for cutaneous T cell lymphoma[109]. Bexarotene has no effect on physiological bone remodeling in mice [108]. Although RXR activation protects mice from OVX-induced bone loss [108], the bone mass is significantly but marginally reversed by bexarotene. Therefore, it further needs to be evaluated the effect of RXR activation on physiological and pathological bone remodeling.

## Conclusions

Despite the importance of NRs, many aspects of NR's skeletal actions are not fully characterized. Studies have reported the alteration of osteoclast differentiation and activity by synthetic ligands of NRs (Figure 4); however, knowledge of the mechanisms of NR's action in osteoclasts is limited. Several factors should be considered. First, genome-wide screening of NRs has identified distinct cell-type specific NR binding sites, their consensus DNA binding sites, and cofactors. For example, genomic mapping of ER $\alpha$  identifies co-regulators of ER $\alpha$ , including FOXA1 in breast cancer cells [110]. However, there are no genomic mapping studies of NRs in osteoclasts. It has been shown that nuclear receptor binding profiles are tissue-specific and species-specific. In addition, in macrophages the co-binding of PU.1, a key factor for myeloid and B cell development, with PPAR $\gamma$  has been found[111], suggesting the co-binding between NRs and key transcription factors facilitate the function of cells. Thus, identifying the osteoclast-specific binding sites of NRs and other co-factors, and mapping the interaction of NRs with the osteoclast-specific transcription factors such as NFATc1 and IRF8 needs to be determined [112]. Secondly, as NRs are regulated by other signals, cross-regulation of NRs with important signals in osteoclasts needs to be determined. Third, multiple NRs co-exist in osteoclasts, and the interplay between NRs reprograms chromatin to open sites in addition to NR's designated binding sites. The crosstalk between NRs needs to be characterized during osteoclastogenesis. Fourth, alternative translation and post-translation modification of NRs generate multiple functional isoforms, and the exact isoform relevant to osteoclast activity needs to be characterized. Fifth, NRs have profound effects on osteoblasts, bone-forming cells. Since the effect comes from global knock mice, it is difficult to differentiate contributions to osteoclasts and osteoblasts. Therefore, osteoclast-specific NR-deficient mice are needed to understand the contributions to osteoclasts. Since NRs play an important role in skeletal biology, further studies to uncover transcriptional networks driven by NRs, binding sites in osteoclasts and evidence that they will affect bone under physiological and pathological conditions will provide a better method of targeting NRs without creating skeletal abnormalities.

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Papers of particular interest have been highlighted as:

\*of special interest

\*\*of outstanding interest

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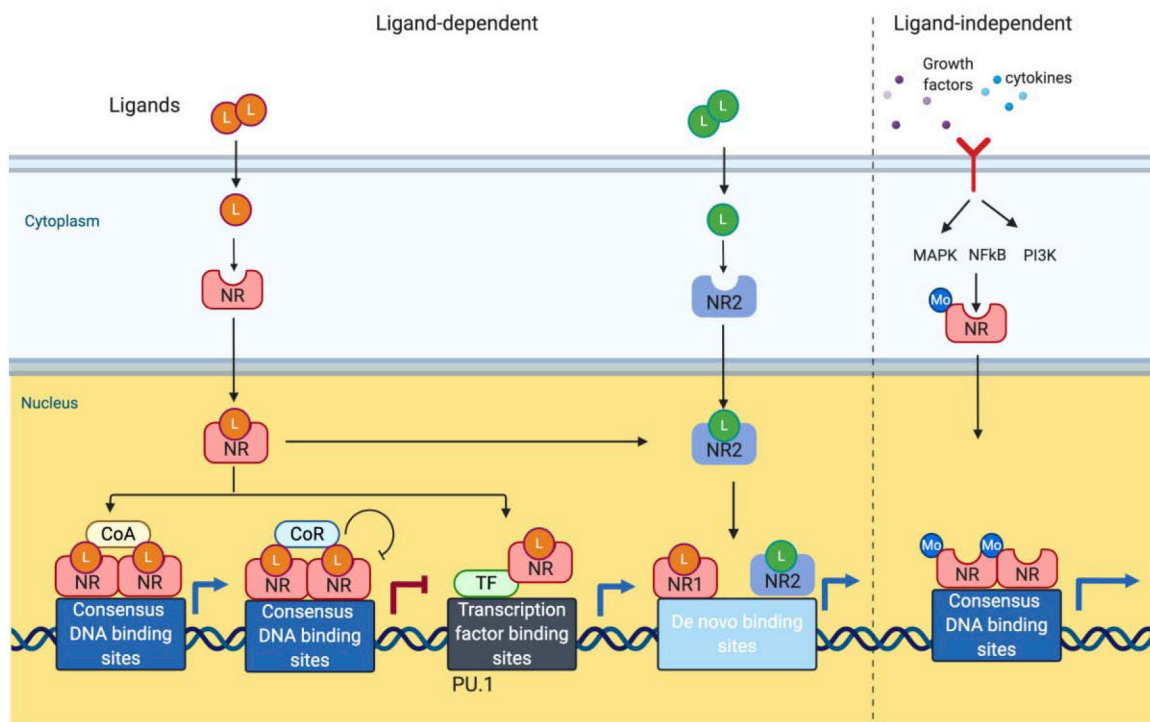
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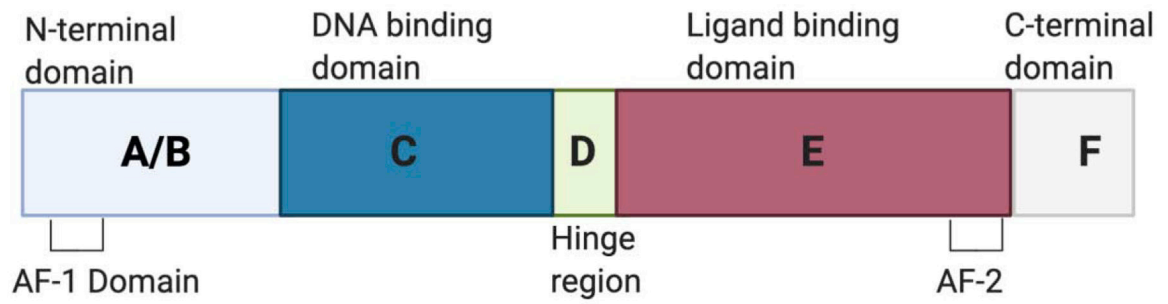
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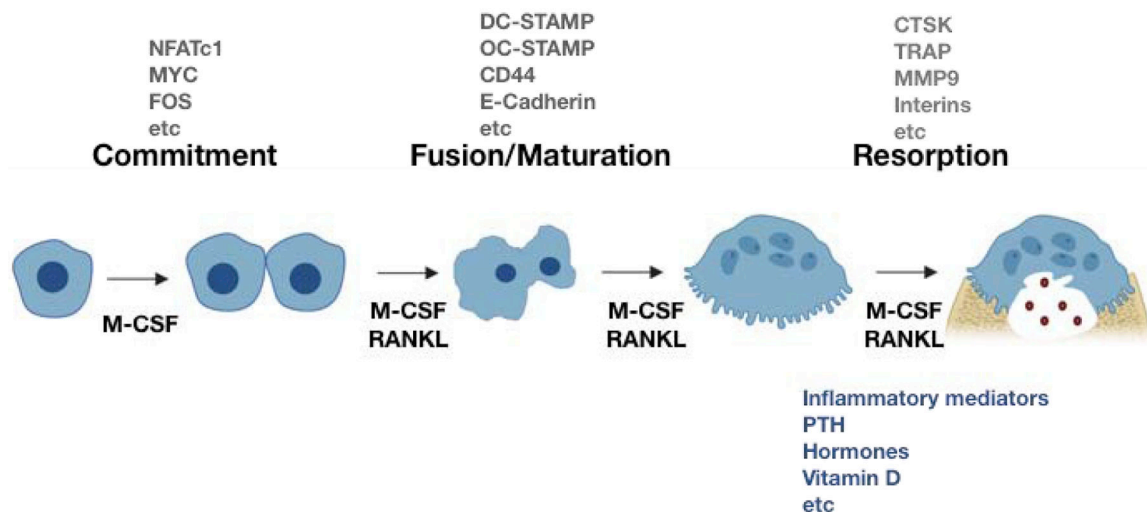
**Figure 1. Nuclear receptor (NR) signaling.**

Upon binding on ligands (agonist or antagonist), cytoplasmic NRs undergo the conformational changes and translocate into the nucleus, where they can signal in genomic ways. Activated NRs also exert their action in non-genomic ways (not shown). In the nucleus, ligand-bound NRs directly bind to consensus DNA binding sites of NRs and activate or suppress target gene expression by interacting with co-activator (CoA) or co-repressor (CoR), respectively. Activated NRs also activate target by binding to other transcription factors (TF). Co-binding of NRs with pioneer factors such as PU.1 and osteoclast-specific TFs including NFATc1 need to be determined. Interplay between two different NRs open up de novo binding sites in chromosome and can regulate gene expression. Some NRs can be activated by different modification (Mo) including phosphorylation through other stimulus such as growth factors or cytokines.



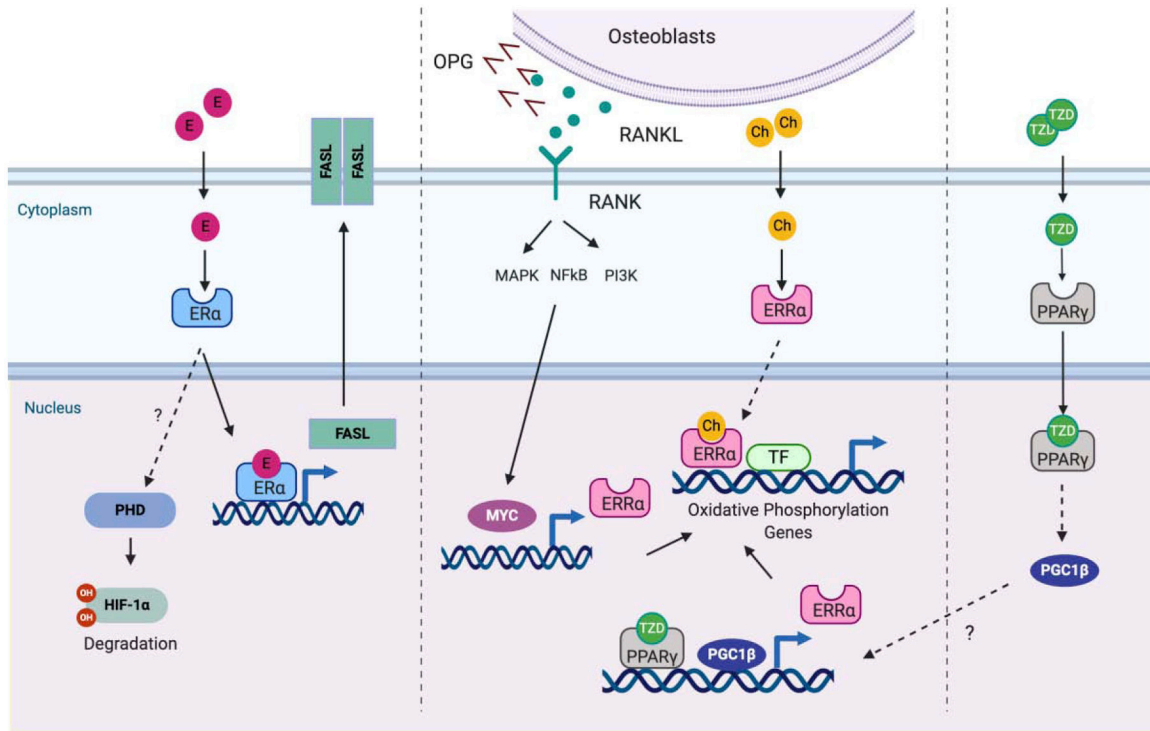
**Figure 2. The structure of nuclear receptors.**

Nuclear receptors share structural homology. A-B is the N-terminal domain containing ligand-independent activation function -1 (AF-1) domain. C has the highly conserved DNA binding domain (DBD), and D is the hinge domain linking DBD to LBD. E has the Ligand binding domain (LBD) that contains AF-2, which is important for the binding of coactivators. Ligand binding to LBD causes the conformational change of nuclear receptors. F is the less conserved C-terminal domain.



**Figure 3. Osteoclastogenesis.**

Osteoclastogenesis progresses in three different phases; commitment, fusion/maturation, and resorption. Osteoclasts differentiate from multipotential, myeloid lineage precursor cells under the influence of a variety of cytokines and local factors. M-CSF and RANKL are essential factors for osteoclastogenesis and induce various transcription osteoclasts. Cell-cell adhesion is initiated by E-cadherin, integrins, and cytoskeletal rearrangement, resulting in the podosome formation. Subsequently, cell-cell fusion is mediated by induction of fusion related genes such a DC-STAMP and OC-STAMP. Osteoclasts tightly attach to bone via the sealing zone and form the resorption pit. To break bone, osteoclasts resorb bone by releasing several enzymes, including CTSK, TRAP, and MMP9 and ions such as  $H^+$  and  $Cl^-$ . The microenvironment surrounding osteoclasts such as inflammatory mediators, PTH, hormones, vitamin D greatly affects osteoclast differentiation and function.



**Figure 4. Nuclear receptor (NR) signaling in osteoclasts.**

On binding of RANKL to RANK, MYC is induced. MYC binds to the promoter of *ERRα* to induce the expression of *ERRα*. *ERRα* subsequently activates genes related to oxidative phosphorylation which is a key metabolic mechanism of osteoclast differentiation. The MYC-*ERRα* axis plays an important role in osteoclast differentiation. The thiazolidinedione (TZD) family binds to and activates *PPARγ*. *PPARγ* indirectly induces *PGC1β*. Both *PPARγ* and *PGC1β* bind to the promoter of *ERRα*. Estrogen negatively regulates osteoclastogenesis. Estrogen binds to and activates *ERα*. Activated *ERα* induces *HIF1α* degradation and *FASL* expression to enhance apoptosis of osteoclasts. Osteoblasts/osteocytes secrete OPG and RANKL. The activation of NRs in osteoblasts regulates the ratio of RANKL to OPG to modify RANKL-induced osteoclastogenesis. factors to initiate the osteoclastogenenic program. Then committed cells (osteoclast precursor cells) fuse to each other and generate multinucleated

Table 1.

The function of NRs in osteoclasts

Type	NRs	Natural ligands	Synthetic ligands	Effect on osteoclasts	Mechanisms in osteoclasts
	RXR $\alpha$	Retinoic acid, Retinol	Bexarotene	Regulate the proliferation of osteoclasts	<ul style="list-style-type: none"> <li>RXRs directly bind to the promoter of MAFB and regulate the expression of MAFB, a negative regulator of osteoclastogenesis</li> <li>Bexarotene induces SREBP1c to regulate MAFB expression</li> </ul>
	RXR $\beta$		BRF110		
			IRX4204		
	RXR $\gamma$			No expression in osteoclasts	N/A
	PPAR $\alpha$	Polysaturated fatty acid	Fenofibrate, statin	Ligand-induced suppression of osteoclastogenesis	<ul style="list-style-type: none"> <li>Mechanism is unknown</li> </ul>
	PPAR $\beta/\delta$	Unsaturated/saturated fatty acid	GW0742, L-165041	Ligand-induced suppression of osteoclastogenesis	<ul style="list-style-type: none"> <li>Mechanism is unknown</li> </ul>
	PPAR $\gamma$	15-deoxy-12, 14 prostaglandin J2 (15d-PGJ2)	Thiazolidinedione (rosiglitazone, ciglitazone, pioglitazone)	Ligand-induced stimulation of osteoclastogenesis and apoptosis in <i>in vivo</i> ; contradicting results	<ul style="list-style-type: none"> <li>Induce PGC1<math>\beta</math> expression</li> </ul>
	LXR $\alpha$	Oxysterols (25OHC, 27OH, 24OHC, etc)	GW3965, T0901317	Activation of LXRs by agonists suppresses osteoclastogenesis and apoptosis of enhances mature osteoclasts	<ul style="list-style-type: none"> <li>Activation of LXRs by agonists suppresses the MTF/p38/ NFATc1 pathway via LXR<math>\beta</math></li> <li>Suppress RANKL-induced c-FOS and NFATc1</li> </ul>
	LXR $\beta$				
	ER $\alpha$	Oestrogen (oestrone (E1), oestradiol (E2), and oestriol (E3))	SERMs Synthetic estrogen (ethinyl estradiol (EE2), diethylstilbestrol (DES))	Negative regulation of osteoclastogenesis Induction of apoptosis in mature osteoclasts	<ul style="list-style-type: none"> <li>Induce HIF1<math>\alpha</math> degradation</li> <li>Induce FASL expression</li> </ul>
	ER $\beta$			N/A	<ul style="list-style-type: none"> <li>Mechanism is unknown</li> </ul>
	GR	Glucocorticoid	Synthetic glucocorticoid (Dexamethasone, Prednisolone RU38486, A348441)	Suppress or increase osteoclast activity <i>in vitro</i> Prolonged life span of osteoclasts	<ul style="list-style-type: none"> <li>Mechanism is unknown</li> </ul>
	ERR $\beta$	N/A	4-methylensterols DY131 (GSK9089) GSK4716	N/A	<ul style="list-style-type: none"> <li>Mechanism is unknown</li> </ul>
	ERR $\gamma$	N/A	Bisphenol A	Overexpression of ERR $\gamma$ suppresses osteoclastogenesis	<ul style="list-style-type: none"> <li>Mechanism is unknown</li> </ul>
	ERR $\alpha$	Cholesterol	inverse agonist, XCT790, Isoflavones, flavone		<ul style="list-style-type: none"> <li>Regulate oxidative phosphorylation</li> </ul>