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## ABSTRACT

Osteoclasts are derived from mononuclear hematopoietic myeloid lineage cells, which are formed in the bone marrow and are attracted to the bloodstream by factors, including sphingosine-1 phosphate. These circulating precursors are attracted to bone surfaces undergoing resorption by chemokines and other factors expressed at these sites, where they fuse to form multinucleated bone-resorbing cells. All aspects of osteoclast formation and functions are regulated by macrophage-colony-stimulating factor (M-CSF) and receptor activator of NF- $\kappa$ B ligand (RANKL), cytokines essential for osteoclast formation and expressed by a variety of cell types, including osteoblast lineage cells. Since the discovery of RANKL in the mid-1990s, mouse genetic and molecular studies have revealed numerous signaling pathways activated by RANKL and M-CSF. More recent studies indicate that osteoclasts and their precursors regulate immune responses and osteoblast formation and functions by means of direct cell-cell contact through ligands and receptors, such as ephrins and Ephs, and semaphorins and plexins, and through expression of chemokines. There is also growing recognition that osteoclasts are immune cells with roles in immune responses beyond mediating the bone destruction that can accompany them. This article reviews recent advances in the understanding of the molecular mechanisms regulating osteoclast formation and functions and their interactions with other cells in normal and pathologic states.

**KEY WORDS:** apoptosis, bone biology, osteoblast(s), bone remodeling/regeneration, cytokine(s), chemokine(s).

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# Advances in the Regulation of Osteoclasts and Osteoclast Functions

## INTRODUCTION

Osteoclasts (OCs) are multinucleated cells that arise by fusion of myeloid hematopoietic precursors formed in the bone marrow. Osteoclast precursors (OCPs) are attracted from the bone marrow to the bloodstream by chemokines and circulate there until they are attracted back into bones by a variety of factors released at sites undergoing resorption, called bone remodeling units (BRUs), and there they differentiate into OCs (Boyce *et al.*, 2012). These factors include chemokines and cytokines, such as macrophage-colony-stimulating factor (M-CSF) and receptor activator of NF- $\kappa$ B ligand (RANKL), which are expressed by cells in and around BRUs and are required for OCP formation and differentiation in normal and pathologic bone remodeling in which bone resorption is increased generally or locally (Boyce *et al.*, 2012).

Until recently, OCs were considered to be bone-resorbing cells with few other functions. However, studies showing that they secrete cytokines (chemokines) and growth factors, and genetic studies in mice have revealed a number of unanticipated roles for OCs and OCPs (Boyce *et al.*, 2009). These include positive and negative regulation of osteoblast (OB) formation, and immune responses. OCs are also found in a number of pathologic conditions outside of bone – for example, within some cancers. Several major signaling pathways regulate the functions of normal and neoplastic cells, including NF- $\kappa$ B, Wnt- $\beta$ -catenin, and Notch, but they also play important roles in OC and OB formation and functions. This paper reviews current knowledge of the molecular mechanisms regulating OC formation and functions.

## RECRUITMENT OF OSTEOCLAST PRECURSORS TO BONE SURFACES

OCPs are held in the bone marrow by chemokines, such as stroma-derived factor-1 (SDF-1). This effect can be modulated by tumor necrosis factor (TNF), which inhibits SDF-1 production by marrow cells, allowing OCPs to mobilize to the blood in inflammatory conditions (Boyce *et al.*, 2012). OCPs are also attracted to the blood by the bioactive sphingolipid, sphingosine-1 phosphate (S1P), which is secreted from red blood cells and platelets. OCPs express S1P receptors (S1PRs) 1 and 2, which have chemo-attractant and chemo-repellent functions, respectively, the latter sending them back to the marrow at BRUs (Kikuta *et al.*, 2011). OCs also express S1P, which could attract OCPs to them for fusion. Interestingly, S1P expression is negatively regulated by cathepsin K, the major matrix-degrading collagenase expressed by OCs (Lotinun *et al.*, 2013), suggesting that cathepsin K inhibitors could stimulate bone formation because S1P expressed by OCs enhances OB precursor differentiation.

S1P levels are increased in the synovial fluid of patients with rheumatoid arthritis (RA), suggesting that it could attract OCPs to affected joints. Mice with inflammatory arthritis treated with FTY720, an S1PR1 agonist, had significantly reduced joint destruction and inflammation, consistent with the

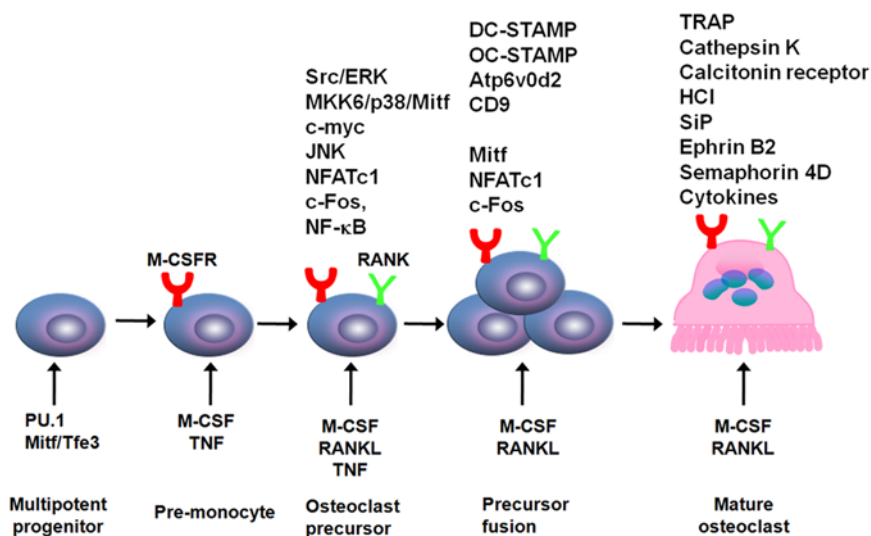
agonist working to retain OCPs and presumably other immune cells in the bloodstream (Kikuta *et al.*, 2011). S1P has positive and negative regulatory effects on inflammation and cytokine expression and functions; thus, agonists may have unanticipated effects in some conditions. S1P levels have not been reported in crevicular fluids around teeth in periodontal disease, but S1P has been shown to increase the production of inflammatory cytokines in periodontitis (Aarthi *et al.*, 2011). Circulating OCPs are also attracted to actively resorbing primary, but not permanent, teeth, presumably by RANKL and M-CSF, which are expressed by pulp and periodontal ligament cells (Wang and McCauley, 2011).

## REGULATION OF OSTEOCLAST FORMATION

Myeloid progenitors differentiate into OCPs in response to signaling induced by several transcription factors, including PU.1, and a heterodimeric complex of microphthalmia-associated transcription factor (MITF) and Tfe3 (Fig. 1). PU.1 and MITF activate expression of the M-CSF receptor (M-CSFR) (Mellis *et al.*, 2011), and mice deficient in the genes encoding any of these proteins have osteopetrosis, which develops during endochondral ossification from failed resorption of bone trabeculae in the medullary cavities. Osteopetrosis occurs because of defective OC formation or activity and is also characterized by failure of incisor eruption in rodents and in abnormal tooth eruption and formation in humans because osteoclasts do not resorb normal channels for them in jaw bones (Del Fattore *et al.*, 2008).

One of the earliest effects of M-CSF is to promote expression of RANK by myeloid progenitors (Fig. 1), a property shared by TNF, which primes the cells to respond to RANKL (Boyce *et al.*, 2012). Binding of M-CSF to M-CSFR attracts a signaling complex comprised of phosphorylated DNAX-activating protein 12 (DAP12) and the non-receptor tyrosine kinase, Syk, and this activates ERK/growth-factor-receptor-bound protein 2 (Grb-2) and Akt/PI3K signaling to regulate several other aspects of OCP and OC activities, including proliferation, differentiation, and survival (Ross and Teitelbaum, 2005). M-CSF also promotes OC survival (Tanaka *et al.*, 2006, 2010) and, like RANKL, mediates all aspects of OC functions.

RANKL is expressed by numerous cell types, including osteoblastic, chondroblastic, and T and B cells (Pacifci, 2012). Hypertrophic chondrocytes appear to be the major source of RANKL for removal of trabeculae during endochondral ossification, while osteocytes within trabeculae appear to be the major source during bone remodeling in adult mice and in response to

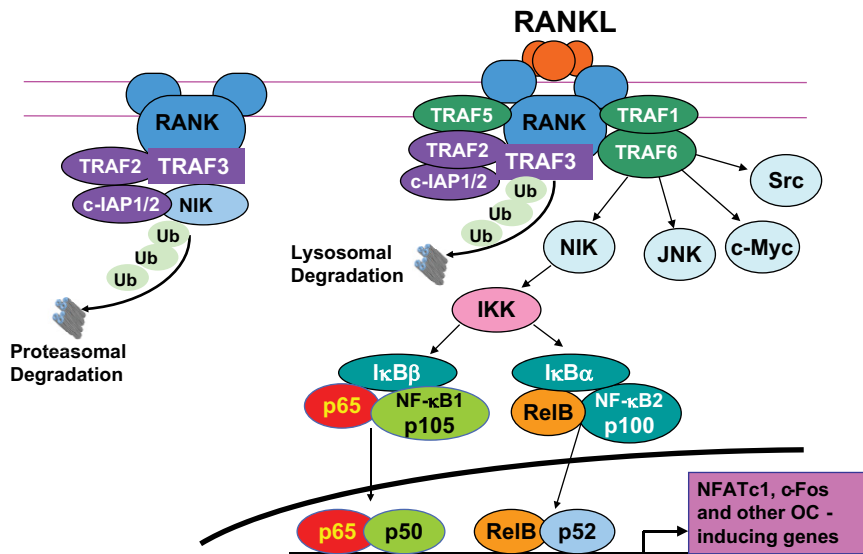


**Figure 1.** Regulation of osteoclast formation and differentiation. An early requirement for myeloid progenitor differentiation into osteoclast precursors is expression of PU.1 and Mitf, which induce expression of M-CSFR, the receptor for M-CSF. M-CSF induces expression of RANK on these cells, priming them for further differentiation in response to RANKL. TNF also induces expression of RANK, which can enhance osteoclast formation in inflammatory bone disease. In response to RANKL and TNF, expression of a number of transcription factors that regulate further differentiation of RANK-expressing cells is increased. These include NF- $\kappa$ B, c-Fos, and NFATc1. They induce expression of several gene-encoding proteins involved in osteoclast activation, including tartrate-resistant acid (TRAP), cathepsin-K, and the calcitonin receptor, and mediate production of H<sup>+</sup> and Cl<sup>-</sup>, which form HCl under the ruffled borders of osteoclasts. RANKL and TNF also induce activation of c-myc, which promotes further proliferation of these cells, as well as map kinase/kinase 6 (MKK6), p38, and Mitf, along with Src and Erk, which have multiple effects to activate osteoclasts and promote their survival.

mechanical stress (Xiong and O'Brien, 2012). However, the major cellular sources of RANKL in common bone diseases remain to be identified.

Like other TNF receptor family members, RANK lacks intrinsic kinase activity to phosphorylate and activate downstream signaling molecules and recruits TNF receptor-activating factors (TRAFs), particularly TRAFs 1, 2, 3, 5, and 6, which are adapter proteins that recruit protein kinases (Boyce *et al.*, 2012). In unstimulated OCPs, TRAFs2 and 3 and cIAP1/2 form a complex that degrades NF- $\kappa$ B-inducing kinase (NIK), and this limits OC formation. RANKL induces degradation of TRAF3, thus allowing for the accumulation of NIK and the induction of OC formation (Fig. 2). TRAF2<sup>-/-</sup> mice die *in utero*, and TRAF3<sup>-/-</sup> mice die around 1 wk after birth, making definitive study of their roles in OC formation challenging. TRAF 1<sup>-/-</sup> and 5<sup>-/-</sup> mice have no bone phenotype, while TRAF6<sup>-/-</sup> mice generated separately by 2 groups have either no OCs or dysfunctional OCs (Wada *et al.*, 2006; Boyce, 2013). This apparent dual role for TRAF6 may explain why RANK/TRAF6 signaling activates NF- $\kappa$ B, c-Jun N-terminal kinase (JNK)/activator protein-1 (AP-1, including c-Fos), c-myc, and calcineurin/nuclear factor of activated T cell c1 (NFATc1) signaling to induce OC formation, as well as Src and MKK6/p38/MITF to mediate osteoclast resorption, and Src and ERK to mediate OC survival (Fig. 1) (Boyce, 2013).

NF- $\kappa$ B signaling is required for osteoclastogenesis, based on NF- $\kappa$ B p50/p52 double-knockout (dKO) mice having marked



**Figure 2.** Functions of TRAFs in osteoclast formation and differentiation. Under basal conditions, TRAF2, c-IAP1/2, and TRAF3 form a complex on the intracellular portion of RANK. This complex polyubiquitinates NIK, which is transported to the proteasome for degradation, resulting in very low levels of NIK in unstimulated OCPs. RANKL binding to RANK on OCPs leads to recruitment of TRAFs 1, 5, and 6 and polyubiquitination of TRAF3 by TRAF2/c-IAP1/2 with subsequent lysosomal degradation of TRAF3 (our unpublished observations). This leads to the release of NIK, and signaling through it and TRAF6 results in the activation of NF-κB and subsequent increased expression of NFATc1 and c-Fos to induce further OCP differentiation. TRAF6 also mediates the activation of signaling through JNK, c-Myc, and Src to promote OCP differentiation and activation.

osteopetrosis due to the failure of OC formation (Boyce, 2013). These dKO, RANKL<sup>-/-</sup>, and RANK<sup>-/-</sup> mice also have failure of B-cell and lymph node development and defective T-cell maturation, suggesting that the inhibition of RANK signaling – for example, with Denosumab (a monoclonal antibody to RANKL) – could have adverse effects on immune responses. However, thus far, studies with Denosumab have not reported such adverse effects (Miller, 2011). Interestingly, despite Denosumab and bisphosphonates having different mechanisms of action, osteonecrosis of jawbones has now been reported in a small percentage of patients treated with Denosumab for metastatic bone disease, suggesting that both drugs may be interfering with the functions of similar cells, which could be osteoclasts and/or other immune cells (Kalyan *et al.*, 2013).

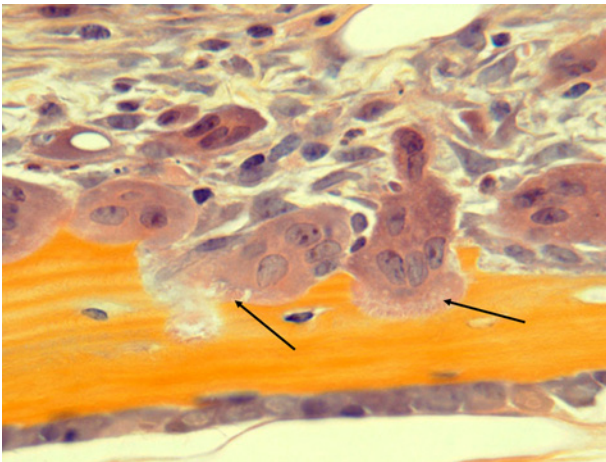
There are 5 main stimulatory NF-κB transcription factor proteins: RelA, p50, Rel B, p52, and c-Rel. In many publications, NF-κB refers to RelA/p50 heterodimeric signaling in the canonical NF-κB pathway. RelB/p52 heterodimers activate non-canonical signaling. RANKL activates both canonical and non-canonical signaling, while TNF activates predominantly canonical signaling (DiDonato *et al.*, 2012). One of the earliest responses of OCPs to RANKL is recruitment of RelA/p50 and NFATc2 to the promoter of *NFATc1*, which has been called the master regulator of osteoclastogenesis. This results in NFATc1 transiently auto-amplifying its own expression (Takayanagi *et al.*, 2002), accompanied by down-regulation of the expression of constitutively active repressors of RANK signaling (Zhao and Ivashkiv, 2011), which allows osteoclastogenesis to proceed.

NFATc1 is subsequently activated by c-Fos through TRAF6/NF-κB and CCAAT/enhancer binding protein-α (Chen *et al.*, 2013) signaling to complete OCP differentiation and subsequent OC activation. These early events in osteoclastogenesis are associated with a number of epigenetic phenomena in response to RANKL. These include: histone acetylation and methylation (Takayanagi *et al.*, 2002) as well as histone demethylation on the NFATc1 promoter (Yasui *et al.*, 2011); inhibition of histone acetylation of ATF3 by Jun dimerization protein 2 (Maruyama *et al.*, 2012); and up-regulation of miR-146a, miR-21, and miR-155 to induce OCP differentiation (Delgado-Calle *et al.*, 2012). In this respect, OCs and their precursors are similar to many other types of cells, and no doubt other forms of epigenetic regulation of OC functions will be identified.

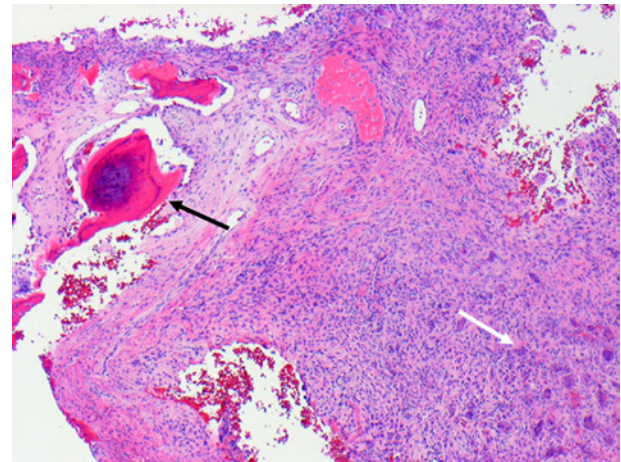
Non-canonical NF-κB signaling by RANKL entails degradation of TRAF3, proteasomal processing of NF-κB p100 to p52, and translocation of RelB/p52 heterodimers to nuclei (Yao *et al.*, 2009). It is not required for basal OC formation because NIK<sup>-/-</sup>, RelB<sup>-/-</sup>, and p100<sup>-/-</sup> mice have normal OC numbers and no osteopetrosis. However, it does appear to be required for localized osteolysis in pathologic states (Boyce, 2013), and TNF-transgenic mice deficient in p100 develop more severe joint inflammation and erosion than do controls, indicating that p100 limits TNF-induced OC formation and inflammation (Yao *et al.*, 2009). A peptide inhibitor of the NF-κB essential modulator (NEMO) reduced inflammatory arthritis in mice (Jimi *et al.*, 2004), but it has not been studied in humans to date. Some factors have been shown to stimulate OC formation independent of RANKL, particularly in pathologic conditions, including 1,25(OH)<sub>2</sub> Vitamin D<sub>3</sub>, TGFβ, IL-6, TNF, APRIL, LIGHT, NGF, and IGFs I and II (Hemingway *et al.*, 2011).

## REGULATION OF OSTEOCLAST ACTIVATION

Bone resorption begins with OCs attaching to bone surfaces by αVβ3, the integrin vitronectin receptor, resulting in recruitment of Src tyrosine kinase by standard outside-in signaling (Teitelbaum, 2011). This leads to activation of several Src-dependent signaling pathways. For example, Src phosphorylates Syk, which recruits DAP12, a co-stimulatory ITAM protein, and Slp76, and these form an adapter protein complex that activates small Rho family GTPases, including Rac. αVβ3 and the M-CFSR interact physically, and inside-out signaling through αVβ3 leads to integrin activation (Teitelbaum, 2011). RANK also forms a complex with αVβ3 through Src, thus activating Syk, Slp-76, Vav3, and Rac, similar to αVβ3/Src interaction.



**Figure 3.** Osteoclasts actively resorbing bone. Osteoclasts with ruffled borders (black arrows) are actively resorbing bone. Note also numerous other mononuclear cells in the marrow adjacent to the osteoclasts. These include immune and stromal cells and osteoclast precursors. H&E, orange G, and phloxine.



**Figure 4.** Giant cell reparative granuloma of bone. Biopsy tissue from a lytic lesion in the mandible of a 14-year-old girl, showing reactive new bone (black arrow), adjacent fibrous tissue, and numerous osteoclasts (white arrow) far removed from the bone surface. H&E.

These interactions lead to the formation of the ruffled border membrane by fusion of lysosomal secretory vesicles with the cytoplasmic membrane (Teitelbaum, 2011). Vesicle fusion is a complex process mediated by several proteins, including Rab7, and synaptotagmin VII, and others involved in autophagy and extracellular protein secretion (Hocking *et al.*, 2012).

H<sup>+</sup> ions are pumped through the ruffled border and, along with Cl<sup>-</sup>, form HCl, which demineralizes bone, and cathepsin K is secreted to degrade the matrix. Cathepsin K inhibitors are being studied in clinical trials as novel anti-resorptive therapy, and a recent study suggests that cathepsin K expressed by OCs negatively regulates osteoblastic RANKL expression (Lotinun *et al.*, 2013). Src expression is required in osteoclasts for resorption, and its over-expression in many cancers enhances proliferation, invasion, and metastasis. Thus, Src-targeted drugs could inhibit both OCs and tumor cells in metastatic bone disease. Saracatinib is a small molecular Src inhibitor under investigation as an adjuvant to standard chemotherapy in metastatic prostate cancer (Boyce and Xing, 2011). Proton pump inhibitors have been developed, but none of these is in clinical trials.

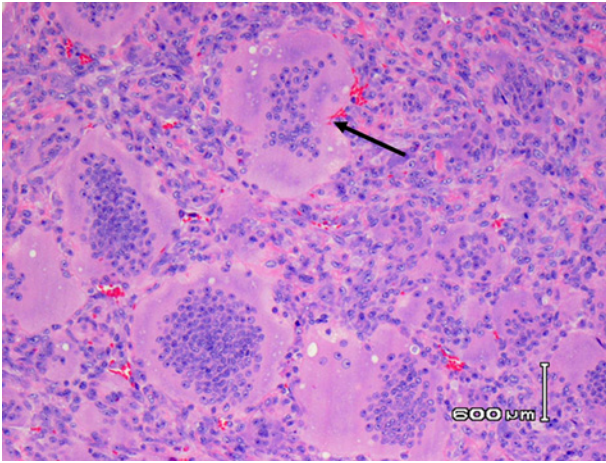
During bone formation, numerous proteins, including TGFβ, BMPs, FGFs, and IGFs, are laid down along with collagen and are released subsequently by OCs during resorption when they can become activated and available locally to promote the resorptive process in normal and pathologic states, such as metastatic bone disease (Kakonen and Mundy, 2003). For example, TGFβ, BMPs, FGFs, and IGFs can induce OC formation by direct and/or indirect (*e.g.*, by modulating expression of RANKL or OPG) mechanisms (Hemingway *et al.*, 2011). TGFβ can also attract osteoblastic cells to resorption sites to act as a coupling factor (Tang *et al.*, 2009).

OCs function most effectively as large multinucleated cells (Fig. 3), such as those seen in osteolytic conditions, including Paget's disease, giant cell tumors, and reparative granuloma of bone. Their formation does not require them to be attached to bone matrix. For example, only a fraction of the OCs in reparative granuloma or giant cell tumors (Figs. 4, 5) is on the bone

surface, and OCs are seen occasionally in primary breast, pancreatic, and other cancers (Fig. 6), where their roles and the mechanisms driving their formation have yet to be determined. In these conditions, tumor cells presumably express RANKL or other osteoclastogenic cytokines (Hemingway *et al.*, 2011). Several proteins, including dendritic-cell-specific transmembrane protein (DC-STAMP), Atp6v0d2, OC-STAMP, and CD9, are involved in the fusion process, and their expression is regulated by PU.1, MITF, c-Fos, and NFATc1 (Fig. 1) (Yagi *et al.*, 2007; Mellis *et al.*, 2011). Vitamin E (α-tocopherol) activates mitogen-activated protein kinase 14 and MITF to induce DC-STAMP expression and OCP fusion (Fujita *et al.*, 2012). These investigators found that doses of α-tocopherol taken by humans as dietary supplements increased OC formation and reduced bone volume in rats, suggesting that excessive Vitamin E consumption could cause osteoporosis.

## OSTEOIMMUNOLOGY AND CO-STIMULATORY SIGNALING

NF-κB and NFATc1 signaling in OC formation, coupled with their required roles in lymph node formation, immune responses, and inflammatory arthritis, spawned the new field of osteoimmunology, in which interactions between bone and immune cells are studied (Takayanagi, 2012). Co-stimulatory signaling is a component of normal and aberrant immune responses. It enhances OC formation and activation by activating NFATc1, which, like NF-κB, regulates immune responses, but it also controls neuronal, cardiovascular, muscle, and other cell functions (Hogan *et al.*, 2003). Co-stimulatory signaling activates NFATc1 through ligand binding to immunoglobulin-like receptors, such as TREM-2 (triggering receptor expressed in myeloid cells-2) and osteoclast-associated receptor (OSCAR) (Takayanagi *et al.*, 2002). The ligands for most of these receptors on OCPs have not been identified, but OSCAR is activated by specific parts of collagen, which become exposed during resorption (Barrow *et al.*, 2011). These receptors recruit adapter



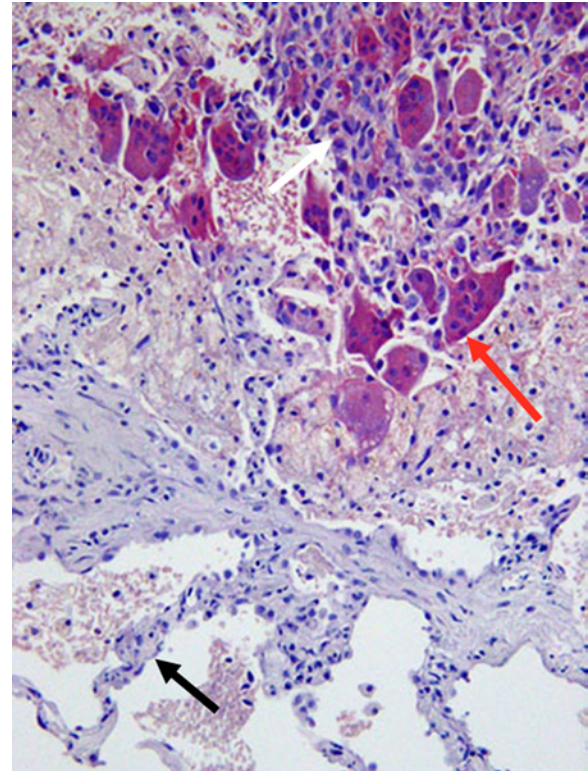
**Figure 5.** Giant cell tumor of bone. Numerous enormous multinucleated osteoclasts from a giant cell tumor of bone from the distal radius of a 25-year-old man, with associated mononuclear precursors and stromal cells with no bone matrix associated with them. H&E.

molecules, including Fc receptor common  $\gamma$  subunit (FcR $\gamma$ ) and DAP12, resulting in phosphorylation of immunoreceptor tyrosine-based activation motifs (ITAMs) within these adapters and the activation of downstream signaling. RANK and co-stimulatory signaling activate phospholipase C $\gamma$  (PLC $\gamma$ ) and calcium-calmodulin signaling, resulting in the release of calcium from stores within the cytoplasm and dephosphorylation of NFATc1 by the calcium-dependent phosphatase, calcineurin, and NFATc1 translocates to nuclei (Takayanagi *et al.*, 2002). Thus, co-stimulatory and RANK signaling likely synergize through NFATc1 activation to enhance OC formation and activation, making NFATc1 a strong candidate for therapeutic intervention in inflammatory bone diseases.

NFATc1 activation can be prevented by cyclosporine-A, a calcineurin/NFATc1 inhibitor used clinically as an immunosuppressive drug and which prevented bone loss in a mouse model of RA (Koga *et al.*, 2005). Interestingly, NFATc1 activation through either RANK or OSCAR increases OSCAR expression on OCPs in a positive feedback loop (Asagiri and Takayanagi, 2007). OSCAR and RANKL expression is increased in the synovium from joints of patients with RA (Crotti *et al.*, 2012), but to date there have been no reports of OSCAR expression in inflamed gingival tissues in patients.

NFATc1 was also discovered to positively regulate the expression of osterix, a transcription factor with essential functions downstream of Runx2 in OB precursor differentiation. Treatment of normal mice with cyclosporine A resulted in osteoporosis because of its inhibitory effects on NFATc1-induced OB formation (Koga *et al.*, 2005), suggesting that it has a more important role in OB than OC differentiation during normal bone remodeling. In inflammatory bone diseases, bone formation typically is inhibited by a variety of mechanisms, including TNF- and dickkopf-1-mediated inhibition of osteoblast functions (Boyce *et al.*, 2012), and in these diseases the major observed effects of NFATc1 inhibitors may be reduced bone resorption.

T and B cells are present in inflamed joints of RA patients and in gingival tissues of patients with periodontitis. They also



**Figure 6.** Osteoclasts associated with a sarcoma in the lung. Multinucleated osteoclasts (red arrow) are present in close association with a spindle-cell primary sarcoma in the lung of a 67-year-old woman. Lung alveolar wall (black arrow). Tartrate-resistant acid phosphatase, counterstained with hematoxylin.

express RANKL to increase osteoclastogenesis in these conditions, but their effects are complex. For example, CD3<sup>+</sup> T-helper (Th) cells express RANKL, and Th17 cells induce RANKL expression through IL-17, but T regulatory cells (Tregs) inhibit OC formation in part through expression of IL-4 and IL-10, and Th1 cells express INF $\gamma$ , which has both inhibitory and stimulatory effects on OC formation, depending upon the stage of OCP differentiation (Zhao and Ivashkiv, 2011). Tregs are present in crevicular fluid from RA patients, but levels of IL-4 and IL-10 were low, and patients with severe periodontitis had the lowest levels (Berthelot and Le Goff, 2010). Overall, the effects of T cells in inflammatory bone disease may be neutral, with synoviocytes being the main source of RANKL (Schett and David, 2010; Takayanagi, 2012). More B cells than T cells are detected in the pericrevicular soft tissues of patients with periodontitis, but there are also conflicting data about the B-cell expression of RANKL (Okamoto and Takayanagi, 2011), and thus further study will be required to determine their roles in inflammatory bone loss.

## NEGATIVE REGULATION OF OSTEOCLAST FORMATION AND ACTIVITY

### Osteoblastic-cell-mediated

OC formation and/or activity is limited by numerous mechanisms, including calcitonin secretion by thyroid C cells, but few

of these were identified before the discovery of RANKL, which facilitated the generation of sufficiently high numbers of highly purified OCPs for gene expression profiling. Osteoprotegerin (OPG), which binds to RANKL to inhibit interaction with RANK, is the major negative regulator of all aspects of bone resorption (Appendix Table 1) (Asagiri and Takayanagi, 2007; Boyce *et al.*, 2012). Loss-of-function mutations of *TNFRSF11B*, the gene encoding OPG, account for most cases of juvenile Paget's disease, in which unopposed RANKL induces osteoporosis (Whyte *et al.*, 2002). This is similar to the phenotype of OPG<sup>-/-</sup> mice.

Osteoblastic cells regulate OC formation through additional mechanisms, some of which regulate OPG expression. For example, canonical Wnt/ $\beta$ -catenin signaling, which is required for OB formation, also promotes OPG expression by OBs (Hill *et al.*, 2005). In contrast, Wnt 5a-induced non-canonical signaling promotes OC formation through receptor tyrosine kinase-like orphan receptor proteins expressed in OCPs (Maeda *et al.*, 2012). In addition, Jagged1/Notch1, which also promotes osteoblast differentiation, increases the OPG/RANKL ratio in stromal cells to inhibit OC formation (Appendix Table 1) (Boyce *et al.*, 2012), and *Zfp521* inhibits OC formation by reducing RANKL expression and suppressing Ebf-1-induced osteoclastogenesis (Kiviranta *et al.*, 2013).

Semaphorins (Semas), which are expressed widely as secreted and membrane-associated proteins, also regulate OC/OB interactions. *Sema3A* is secreted by OBs and OCs and inhibits RANKL-induced OC formation by inhibiting ITAM and RhoA signaling. *Sema4D* is expressed by OCs and inhibits OB differentiation and function (Appendix Table 2) by activating RhoA-ROCK, which inhibits IGF-1 signaling. In contrast, *Sema6d* induces OC formation through TREM-2/DAP12/PLC $\gamma$ -induced NFATc1 activation and promotes OC activation by inducing podosome formation through Rac-GTP generation in OCs (Kang and Kumanogoh, 2013). Semaphorins signal through plexins and neuropilins, and knockout mice lacking some of these proteins have either osteopetrosis or osteoporosis, supporting important roles for them in bone cell functions.

RANKL also limits OC formation through several mechanisms. For example, although c-Fos mediates NFATc1-induced OC formation, it also promotes secretion of interferon- $\beta$ , which in turn binds to its receptor on OCPs, leading to reduced c-Fos protein levels (Asagiri and Takayanagi, 2007). c-Fos/NFATc1 signaling also increases the expression of ephrinB2 on the surfaces of OCPs. Ephrins control axon, endothelial, and immune cell functions during embryonic development through direct interaction with Eph receptors on adjacent cells and bi-directional signaling (Davy and Soriano, 2005). Ephrin B2 binding to Eph4 on osteoblastic cells down-regulates c-Fos and NFATc1 expression through reverse signaling to reduce OC formation, and forward signaling through Eph4 promotes OB precursor differentiation by inhibiting the small GTPase, RhoA (Appendix Tables 1, 2) (Zhao *et al.*, 2006).

Many questions remain about exactly how and where in BRUs these various interactions take place and which ones are between OCs and OBs or their precursors. Intuitively, it would make sense to have OB precursors positively regulating OC

formation and fusion at the resorbing edges of BRUs and also promoting OC apoptosis in reversal sites where OCPs could enhance OB precursor differentiation. In contrast, mature OB secretion of OPG and expression of other negative regulators of OC formation would keep OCs away from bone-building sites.

### Constitutive Transcriptional Repression of RANK Signaling

There are also constitutive mechanisms to inhibit basal OC formation. For example, in the absence of RANKL, Bcl6 is recruited to the NFATc1, cathepsin K, and DC-STAMP promoters to inhibit osteoclastogenesis. RANKL induces the removal of Bcl6 from these promoters and its replacement by NFATc1 to mediate osteoclastogenesis (Miyachi *et al.*, 2010). Interferon regulatory factor-8 (IRF-8), Eos, and v-maf musculoaponeurotic fibrosarcoma oncogene family protein B are other constitutively expressed transcriptional repressors (Zhao and Ivashkiv, 2011), and their expression is down-regulated by RANK signaling. They are also direct targets of B-lymphocyte-induced maturation protein-1, deletion of which in OCs results in osteopetrosis due to up-regulation of Bcl6 and impaired osteoclastogenesis (Miyachi *et al.*, 2010). In contrast, Bcl6<sup>-/-</sup> mice have increased OC formation and severe osteoporosis. Thus, RANKL/RANK activation of NFATc1 in OCPs not only promotes osteoclastogenesis directly, but also facilitates it indirectly by repressing expression of these negative regulators.

### Pro-inflammatory Cytokines

Cytokines, such as TNF, are important inducers of bone resorption and inflammation in numerous disorders, including RA and periodontal disease (Boyce *et al.*, 2012), but they also activate mechanisms to restrict the destruction. For example, most of the factors that induce expression of RANKL also induce expression of OPG, albeit to a lesser degree, the net effect being increased bone resorption (Kearns *et al.*, 2008). Similarly, although TNF induces resorption through and independently of RANKL (Yao *et al.*, 2009), it also limits OC formation by several mechanisms. These include preventing the degradation of non-canonical NF- $\kappa$ B inhibitory proteins (Yao *et al.*, 2009) and inducing expression of IRF-8 and the Notch-induced DNA-binding molecule, RBP-J $\kappa$ , in OCPs (Zhao and Ivashkiv, 2011). TNF also promotes secretion by OCPs and OCs of TNF-stimulated gene 6, which synergizes with OPG to limit OC activity through an autocrine mechanism (Mahoney *et al.*, 2011). IL-10, an anti-inflammatory cytokine, which functions to resolve inflammation, limits OC formation by inhibiting expression of c-Fos, c-Jun, TREM-2, and NFATc1 in OCPs. IL-4 limits bone resorption by promoting OPG expression and suppressing expression of RANKL, RANK, NF- $\kappa$ B, c-Fos, NFATc1, MAPK, and calcium signaling during OC formation (Zhao and Ivashkiv, 2011). During co-stimulatory immune reactions, ITAM-bearing proteins partner with proteins containing immunoreceptor tyrosine-based inhibitory motifs (ITIMs), and these negatively regulate osteoclastogenesis (Mori *et al.*, 2008).

## OSTEOCLAST APOPTOSIS AND SURVIVAL

OCs undergo apoptosis at reversal sites in BRUs where new bone is laid down by OBs. Estrogen maintains bone mass in part by promoting OC apoptosis mediated by TGF $\beta$  (Boyce, 2013) and increasing Fas-ligand expression in OCs, but also by inhibiting expression of genes regulating OC activity, without affecting OCP proliferation or fusion (Nakamura *et al.*, 2007). Bisphosphonates induce OC apoptosis, in part by inhibiting the activity of enzymes in the mevalonate pathway and promoting caspase cleavage of Mammalian Sterile 20-like (Mst) kinase 1 (Rogers *et al.*, 2011), although some amino-bisphosphonates can inhibit bone resorption without inducing OC apoptosis (Matsumoto *et al.*, 2011). Denosumab (Hanley *et al.*, 2012) and raloxifene (Krum *et al.*, 2008) also induce OC apoptosis, but calcitonin and cathepsin K inhibitors do not (Boonen *et al.*, 2012).

OC survival is enhanced by cytokines, including M-CSF, RANKL, TNF, IL-1, and VEGF-A, through up-regulation of Rho family small G-protein Ras/Rac1/Erk and PI3 kinase/mTOR/S6K signaling (Tanaka *et al.*, 2006), while cytokine withdrawal leads to reduced expression of the anti-apoptotic protein, Bcl-2, and rapid OC apoptosis (Tanaka *et al.*, 2010). An early effect of RANKL signaling is NF- $\kappa$ B p65-mediated prevention of Bid- and caspase 3-induced OCP apoptosis (Vaira *et al.*, 2008). M-CSF prevents OC apoptosis by several mechanisms, including: activation of MITF, which increases Bcl-2 expression (Tanaka *et al.*, 2006, 2010); increased degradation of Bim by c-Cbl, an ubiquitin ligase; up-regulation of Bcl-X<sub>L</sub> expression, which inhibits cleavage of procaspase-9; and inhibition of caspases 3 and 9, which initiate apoptosis. Bim is a pro-apoptotic Bcl-2 family member whose expression is down-regulated by IL-3 signaling through the Raf/Erk and/or PI3K/mTOR pathways. Bim<sup>-/-</sup> mice have decreased OC activity, despite increased OC survival (Tanaka *et al.*, 2010). However, enhanced OC survival overall is associated with increased bone resorption.

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