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Mechanisms of bone loss in rheumatoid arthritis

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Abstract Rheumatoid arthritis (RA) is an autoimmune inflammatory disease in which destruction of bone in the joints causes major morbidity. Recent research has shed light on the cell and molecular mechanisms that lead to this osteolysis, all due directly or indirectly to the chronic inflammation. The aspects of this research covered in this review include the alteration of cell proliferation and survival that results in growth of the RA synovium. This process depends upon an increase in angiogenesis and local blood flow, which is also a feature of increased bone turnover. In addition, the inflammatory environment increases expression of chemokines, which are involved in the recruitment of monocytic osteoclast precursors. Chronic inflammation also promotes an overall catabolic state, with increased osteoclast differentiation and resorptive activity, driven by dysregulation of receptor activator of NF- κ B ligand (RANKL) and the synergistic activity of inflammatory cytokines such as tumor necrosis factor- α and interleukin-1. Osteoclast survival is increased in this environment, but osteoblast differentiation and survival are decreased, with a consequent reduction in bone formation and a net loss of bone. Recognition of these processes and the factors involved will enable more effective and targeted treatments for RA.

Key words Bone resorption · Osteoclast · Osteolysis · Rheumatoid arthritis (RA)

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Introduction

In rheumatoid arthritis (RA), inflammation of the synovial tissues can result in the destruction of the joint through the breakdown of bone and cartilage. Although secondary osteoporosis is often seen,¹ the bone erosion seen in RA is largely localized to the inflamed joints.^{2,3} The systemic and localized bone loss is distinct from systemic hormonally regulated bone pathologies. Despite RA affecting approximately 2% of the world's population, we still have an incomplete understanding of the processes of this chronic systemic disease and, until recently, studies were largely confined to the inflammation that occurs in the soft tissues. However, recent advances in the understanding of bone metabolism have enabled investigation of the mechanisms involved in the bone and cartilage loss in this disease.

Physiological osteoclast formation

The bone remodeling unit (BMU)

In maturity, the integrity of bones is maintained by ongoing bone removal and replacement, a process termed bone remodeling.⁴ Bone remodeling occurs in the “bone remodeling unit (BRU)” or “basic multicellular unit (BMU),” terms coined originally by Frost⁵ to describe the structures in which bone removal by osteoclasts and its replacement by osteoblasts occurs in a highly coordinated sequence. The initiation or “activation” of BMUs requires signals that result in the recruitment of osteoclast precursors, and their differentiation and activation, to remove bone matrix.⁶ Both targeted and nontargeted (stochastic) remodeling are thought to occur, although the relative extent of either is not known. However, there is evidence that targeted remodeling takes place in response to bone matrix damage and fatigue loading of bone.^{7,8} The reason that this remodeling is targeted to a specific site is thought to be due to local loss of osteocyte viability.⁹ It has been argued¹⁰ that at the origin of each BMU a local “area code” is turned on that

provides the signals for preosteoclasts, circulating cells of the monocyte/macrophage lineage to exit the circulation at the appropriate site. The nature of such signals is not known, although osteocytic NO¹¹ and osteopontin¹² have been proposed as possible factors. A cascade of events can be proposed, such that signals elicited by osteocyte death, for example, might be further amplified by bone lining cells or vascular endothelial cells into signals that attract osteoclast precursors. Consistent with this, several cell-attractant molecules have been shown in vitro to attract osteoclast precursors. The chemokine SDF-1 has been reported to increase the recruitment of CXCR4-positive cells, including osteoclast progenitor cells, to regenerating bone.^{13,14} Likewise, chemokines that bind the receptor CCR1 (macrophage inflammatory protein [MIP]-1 α , regulated on activation of normal T cell expressed [RANTES], monocyte chemoattractant protein [MCP]-1, and MCP-3) have been implicated in the recruitment of osteoclast precursors and in their differentiation (depicted schematically in Fig. 1A).¹⁵⁻¹⁷ In diseases such as RA, some of these processes appear to be overridden, while production of molecular factors that promote bone turnover are dysregulated.

Factors required for osteoclast formation

Once recruited to sites of bone remodeling the monocyte osteoclast precursors differentiate in response to local signals. Formation of osteoclasts in vitro was shown to require contact between cells of the osteoblast lineage and osteoclast precursors and the action of pro-resorptive stimuli, such as parathyroid hormone, interleukin (IL)-1, IL-11, prostaglandin E, and 1,25(OH)₂ vitamin D, is mediated by osteoblastic cells.¹⁸ These factors all increase the production and cell-surface expression of receptor activator of NF- κ B ligand (RANKL), by osteoblastic cells. RANKL interacts with its receptor, RANK, located on the surface of osteoclast precursor cells and mature osteoclasts.¹⁹ Experimentally, osteoclast formation occurs in vitro upon exposure of osteoclast precursors to soluble RANKL, a process that also requires the presence of macrophage colony-stimulating factor (M-CSF).²⁰ A secreted RANKL binding protein, osteoprotegerin (OPG),^{21,22} is also produced by osteoblastic cells and inhibits both the formation and the activity of osteoclasts by binding to RANKL and blocking its binding to RANK. There is now a great deal of evidence that the local RANKL/OPG ratio determines the effective activity of RANKL to promote osteoclast formation, which supports the concept that RANKL and OPG occupy a central place in the regulation of bone turnover.^{23,24} Interestingly, OPG has also been shown to inhibit monocyte migration towards RANTES and MCP-1.²⁵ RANKL promotion of osteoclast differentiation requires the presence of M-CSF; however, a large number of other factors have been identified that can modulate the activity of RANKL. These include tumor necrosis factor (TNF)- α ,²⁶ molecules such as DAP12 and FcR γ providing ITAM-dependent costimulatory signals,²⁷ IL-3, IL-10, and osteoclast inhibitory lectin (OCIL).²⁸ This review focuses on those factors

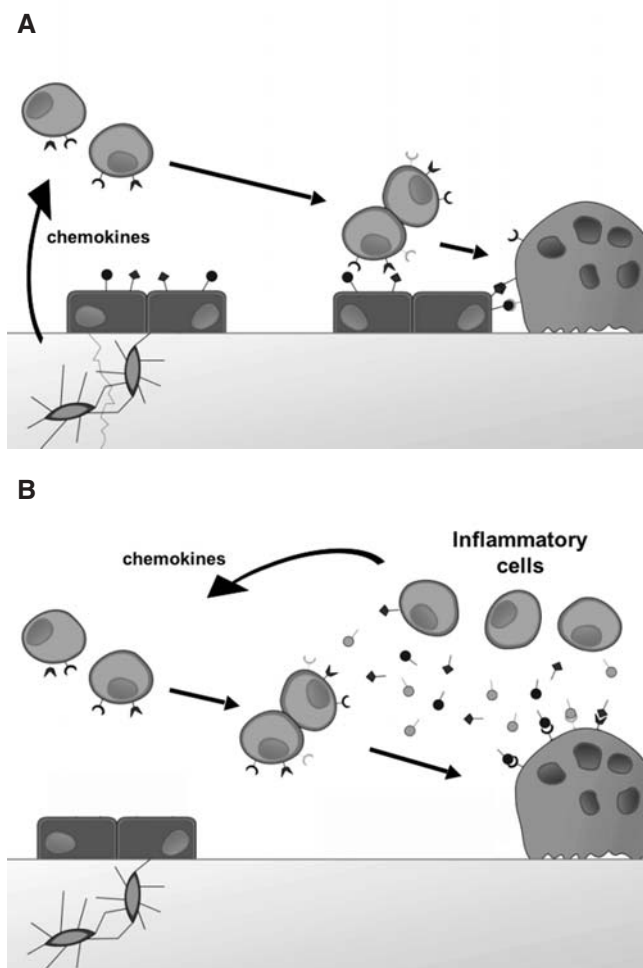


Fig. 1. **A** Proposed initiation of events in physiological bone resorption. Structural damage in the bone matrix results in osteocyte-initiated signals, which stimulate local release of cytokines and chemokines such as tumor necrosis factor (TNF)- α , interleukin (IL)-1 β , and macrophage inflammatory protein (MIP)-1 α . These factors stimulate expression by osteoblasts and/or lining cells of receptor activator of NF- κ B ligand (RANKL), while suppressing osteoprotegerin (OPG) production. In addition, chemokines such as regulated on activation of normal T cell expressed (RANTES) and monocyte chemoattractant protein (MCP)-1 are produced that recruit monocyte osteoclast precursors to sites of bone to be repaired, promote their fusion to multinuclear cells, and bring them into proximity to RANKL-expressing osteoblastic cells for maturation to functional osteoclasts. **B** Proposed initiation of events in pathological bone resorption. In inflammatory conditions, such as rheumatoid arthritis, there is prolonged production of inflammatory cytokines and chemokines such as TNF- α , IL-1 β , and MIP-1 α . The source of these molecules is the inflammatory cells, involving macrophages and lymphocytes as part of an immune response, rather than resident bone cells as part of a bone repair mechanism. In addition, RANKL is produced by the inflammatory cells and OPG is suppressed, resulting in osteoclast formation in the inflammatory tissues in the absence of osteoblasts

that regulate inflammatory-mediated osteolysis, in particular osteolysis in RA. Evidence is presented that dysregulated bone turnover that results in the net loss of bone can be due to RANKL expression by cells not of the osteoblastic lineage. In addition, the inflammatory environ-

ment can upregulate osteoclast formation, activity and survival while inhibiting osteoblastic bone formation. Since excellent reviews are available covering a number of aspects of this topic,^{29–32} this review describes the intimate relationship between inflammation and osteolysis in RA, with particular emphasis on some of the more newly recognized factors that may mediate bone resorption in inflammatory environments.

Evidence for involvement of RANKL in RA-induced bone loss

RANKL in animal models of RA

Elegant experiments by Redlich et al.³³ showed that bone erosion, in at least one model of RA, is mediated by osteoclasts. These authors crossed transgenic mice that over-express human TNF (hTNFtg) and that develop a severe and destructive arthritis, with osteopetrotic, *c-fos*-deficient mice (*c-fos*^{-/-}) completely lacking osteoclasts. The resulting mutant mice (*c-fos*^{-/-}/hTNFtg) developed a TNF-dependent arthritis, with paw swelling and reduction of grip strength, but were fully protected against bone destruction. Other animal models of RA provide good evidence for a role for RANKL in the osteoclast-mediated joint destruction of this disease. For example, in collagen-induced arthritis (CIA) in mice, cells expressing RANK increased in abundance with the progression of arthritis.³⁴ In areas of focal bone erosion where RANK-positive cells were abundant, TRAP-positive multinucleated osteoclast-like cells are also present suggesting differentiation of RANK-positive osteoclast precursor cells that are present in the inflamed synovial tissues. In addition, sites of RANK-expressing cells correlated with sites of RANKL expression. There is strong evidence that RANKL is important in the bone destruction in mouse models of RA. Firstly, RANKL gene knock out mice were protected from bone erosions in RA in a serum transfer model of arthritis.³⁵ It has been found that RANKL expression is increased in activated T cells but not in quiescent cells.³⁶ Kong et al.³⁷ reported that both soluble and membrane-spanning forms of RANKL are produced by activated T cells. Secondly, blocking of RANKL with OPG treatment at the onset of induced RA in mice prevented bone and cartilage destruction but had little effect on inflammation.³⁷ Protection of the cartilage was thought to be secondary to the protection of the subchondral bone. Thus, although none of the mouse models of RA perfectly reflect human RA, these data are nevertheless consistent with a central role for RANKL in the bone erosion side effect of this disease.

RANKL in human tissues in RA

Examination of human RA tissues also provides evidence for a key role for RANKL produced in the inflamed synovial tissues in destructive RA. RANKL appears to be over-expressed in pannus tissue from active RA.^{38,39} This

finding is consistent with the ready proliferation of macrophage-like cells from these tissues and their rapid differentiation into multinucleated bone resorbing cells.^{38,40} In addition, osteoclasts formed from circulating precursors from RA patients produced significantly greater bone resorption than cells from controls, which was ascribed to increased functional activity rather than osteoclast formation.⁴¹ Importantly, cells other than, or in addition to, osteoblasts may be a source of RANKL in inflammatory tissues. Using antibodies directed against RANKL and OPG, several studies have shown “ectopic” production of RANKL by several cell types in a variety of inflamed tissues adjacent to sites of bone resorption.^{39,42–45} In particular, RANKL protein was predominant within inflammatory cells in the inflamed tissues adjacent to pathological bone loss in RA.³⁹ It is important to note that the expression of RANKL in the tissues adjacent to bone loss was significantly greater than in the relevant controls in many of these diseases.^{39,42,43,46} Investigation of RANKL expression in human RA pannus tissue showed strong RANKL staining in the inflamed pannus and dual staining showed expression by both CD3 cells and macrophages.³⁹ These data are consistent with other reports that CD3-positive lymphocytes in synovial tissue of active RA patients are the predominant cell type expressing RANKL protein.^{39,46} Moreover, lymphocytes isolated from inflammatory arthritis pannus produce RANKL^{47,48} and importantly, there is evidence that activated T cells may regulate osteoclast activity by producing RANKL.⁴⁹ RANKL was also associated with monocyte/macrophages, and multinucleated cells expressing CD68 in pannus tissue from patients with rheumatoid arthritis as well as in inflammatory periodontal disease and in peri-prosthetic osteolytic lesions. In addition, a recent report demonstrated RANKL mRNA expression in macrophages and multinucleated cells in diseased tissues in an animal model of periodontitis associated with localized bone loss.⁵⁰ Since these cells may also express RANK it is possible that macrophages can independently “auto-stimulate” themselves to become osteoclasts. RANKL is also reported to be expressed by synovial fibroblasts⁵¹ and isolated microvascular endothelial cells.⁵² Fibroblasts may contribute to the formation of osteoclasts in inflammation. However, the absence of RANKL protein in endothelial cells residing in a variety of inflamed tissues suggest that endothelial cells may not have a role in directly stimulating osteoclast formation in inflammatory diseases.^{43–45} These studies suggest that most of the RANKL present in inflamed tissues associated with osteolytic lesions is produced by activated leucocytes that are recruited to sites of inflammation.

Although it is established that OPG exists in both monomeric and dimeric forms,⁵³ the significance of this and the roles of the different forms are not yet known. Using antibodies that differentiate between these forms, dimeric OPG was only detected in blood vessels (on factor VIII-positive endothelial cells), whereas monomeric OPG was predominantly detected in the synovium (on CD68 positive type A synoviocytes) in RA and other tissues.⁴³ The function of OPG produced by endothelial cells or type A synoviocytes is yet to be determined. However, it may be significant that

the expression of OPG was markedly reduced in the blood vessels and synovium of inflamed RA joint tissues in active disease compared to the corresponding tissues in disease remission or the appropriate control tissues. Endothelial OPG may be important in enhancing endothelial cell survival^{54,55} and may have a role in regulating angiogenesis in inflammation. OPG also has a potential role in inflammatory disease by providing modulation of the immune system.^{30,31} While RANKL has been shown to prolong the survival of dendritic cells and cause proliferation of T cells,⁵⁶ OPG can abolish these effects. Thus OPG can decrease the production of cytokines such as IL-6 by RANKL-stimulated dendritic cells and also decrease the generation of cytokines by proliferating T cells.⁵⁷

Factors that stimulate RANKL, or amplify the effectiveness of RANKL/RANK signaling

Together with RANKL, M-CSF also has a central role in the bone destruction of RA. M-CSF is expressed abundantly in the RA joint and is a necessary cofactor for RANKL promotion of osteoclastogenesis. Exogenous addition of M-CSF to animal models of RA exacerbates the disease.⁵⁸ Other cytokines, such as TNF- α , that stimulate osteoclast formation *in vitro* are also present in high levels in RA tissues⁵⁹⁻⁶⁸ and this cytokine environment is likely to augment the RANKL/RANK pathway in the bone of affected joints. Although TNF- α , like other inflammatory cytokines stimulates both OPG and RANKL expression,⁶⁹⁻⁷² it has also been reported that TNF- α acts synergistically with RANKL by sensitizing osteoclast precursors to low levels of RANKL²⁶ and may stimulate osteoclast formation directly in the absence of RANKL.⁷³ Tumor necrosis factor- α antagonists and antibodies are now an accepted treatment regime for active rheumatoid arthritis. Inhibition of TNF- α activity not only reduces the inflammatory response⁷⁴ but may also inhibit osteoclast bone resorption by reducing the activity of RANKL present in inflammatory tissues.

Arachidonate metabolites are important inflammatory mediators produced during inflammation, and these molecules are reported to regulate osteoclast formation and activity. The role of prostaglandin (PG) E2 has been studied extensively and its effects can vary, possibly depending on the stage of osteoclast differentiation.⁷⁵ Another inflammatory mediator which has been implicated in destructive arthritis is the T-cell product, IL-17.⁷⁶ Interleukin-17 is present in abundance in active RA joints and is a potent inducer of RANKL like TNF- α and IL-1. Interleukin-17 is likely to be important in inflammatory mediated osteolysis in those situations characterized by a major contribution by T cells. The actions of IL-18 seem to depend on the environment in which it is expressed. In noninflammatory situations, both *in vivo* and *in vitro*, IL-18 inhibits osteoclast formation by inducing naïve T cells to produce inhibitors such as granulocyte (G)M-CSF and interferon (IFN)- γ .²⁸ However, in inflammatory animal models IL-18 can augment osteoclast formation. Dai et al.⁷⁷ have examined the actions of IL-18 in human osteoclast precursors incubated

with normal peripheral blood T cells or synovial T cells isolated from RA synovium. In this model, IL-18, IL-1, and TNF- α all increased both soluble RANKL production and membrane bound RANKL expression in PHA prestimulated T cells or RA synovial T cells, but not in resting control T cells.⁷⁷ Upregulation of RANKL was accompanied by increased osteoclastogenesis and increased bone resorption *in vitro*.

A number of chemokines are abundant in inflammatory conditions and are likely to be important in recruitment of monocyte osteoclast precursors into the inflammatory pannus and eventually to sites of bone resorption. They may also potentiate the effects of RANKL in osteoclastogenesis. Monocyte chemotactic protein 1 (MCP-1) is a CC chemokine commonly found at sites of RA bone degradation and other inflammation-induced bone loss.⁷⁸ MCP-1 is expressed by mature osteoclasts and its expression is regulated by NF- κ B.^{79,80} Kim et al.¹⁷ identified MCP-1 by microarray as a RANKL-induced molecule during osteoclast differentiation. Osteoclasts were not formed when production of MCP-1 expression was suppressed by GM-CSF or when RANKL signaling through NFAT-c1 was blocked with cyclosporin A. Furthermore, in the presence of RANKL MCP-1 increased osteoclast formation whereas MCP-1 treatment of monocytes in the absence of RANKL promoted the formation of multinucleated TRAP-positive cells that could not resorb bone.¹⁷ These proposed actions of chemokines are depicted in Fig. 1B.

Role of angiogenic factors in RA

The development of new vasculature is essential to sustain the growth of the rheumatoid pannus and angiogenesis has therefore been proposed as a target in RA management.⁸¹ Inflammatory mediators that have been shown to induce angiogenesis at the site of the inflamed synovium include vascular endothelial growth factor (VEGF), fibroblast growth factor (FGF), integrin alpha(V)beta(3), TNF- α , IL-1, IL-8, IL-18, hypoxia and angiopoietin, and prostaglandins.^{81,82} Significantly, it has also been proposed that RANKL has angiogenic activity.⁸³ The likely role of angiogenic agents in RA disease progression is suggested by a study, which showed that serum VEGF levels at presentation in patients with early RA correlated with the development of radiographic damage after one year.⁸⁴ Based on the likely association between angiogenesis inhibition, and the reduction of signs and symptoms of RA by antiangiogenic drugs, these agents have been considered for the treatment of proliferative synovitis. These include angiostatin, endostatin, IL-1 receptor antagonist, thalidomide, fumagillin, and others.⁸¹

The vasculature is also important in bone metabolism and bone turnover. The elegant work of Reeve and coworkers persuasively demonstrated a relationship between blood flow and bone turnover⁸⁵ and indicated that bone can regulate its own blood flow. They developed a method of measuring blood flow in bone that depends on the complete

extraction of ^{18}F in a single passage through bone. They found that skeletal blood flow (SBF) in untreated Paget's disease of bone was strikingly increased in proportion to the biochemical severity of the disease. Treatment with calcitonin brought a rapid return to normal,⁸⁵ which was postulated to explain the rapid pain relief often reported on starting treatment with calcitonin. Likewise, treatment with sodium etidronate reduced the SBF and the responses of patients correlated well with their reductions in serum alkaline phosphatase.⁸⁵ These data indicate that active bone resorption may enhance local blood flow to the region and this could in turn further enhance synovial inflammation.

Regulation of cell survival: relationship to bone erosion in RA

Agents that alter cell survival can change the balance between anabolic and catabolic processes in tissues⁸⁶ and can lead to the unregulated proliferation and overgrowth of tissues such as the rheumatoid pannus. The proapoptotic molecules Fas and Fas-ligand have been implicated in RA,⁸⁷ since expression of these molecules is elevated in synovial and other cells in active RA disease. The inflammatory environment in RA is likely to influence a cell's susceptibility to Fas/Fas-L and inactivating mutations in these proteins in autoimmune disease have also been associated with proliferative arthritis.⁸⁸ It has also been suggested that regulation of apoptosis by TRAIL and TRAIL receptor expression may also be a factor that maintains inflammation in arthritis.^{89,90} Abnormal TRAIL receptor expression, principally upregulation of TRAIL decoy receptors, may be an important factor in determining the activity of TRAIL. The fact that TRAIL can inhibit the formation of osteoclasts *in vitro*⁹¹ suggests that TRAIL receptors on the precursors of osteoclasts may be important in regulating bone loss in RA.

In bone, prolonged survival of osteoclasts may be one mechanism for increased bone resorption, and perhaps a net loss of bone. Several positive and negative regulators of osteoclast life span have been identified, including hormones, cytokines, and growth factors.⁹² RANKL,⁹³ M-CSF,⁹⁴ IL-1,⁹⁴ and TNF- α ⁹⁵ all support osteoclast survival *in vitro*, and, as discussed above, these factors are all elevated in the RA joint. The mechanism by which RANKL, TNF- α , and IL-1⁹⁵ increase osteoclast survival is thought to involve the activation of Akt and the inactivation of Bad and caspase 9, thus preventing activation of the apoptosis cascade. M-CSF-mediated osteoclast survival also works through Akt⁹² in a different way, resulting in increased expression of the anti-apoptotic genes, Bcl-2, Bcl-xL,⁹³ and XIAP.⁹⁴

Bone loss in RA would be exacerbated if osteoblast differentiation and survival were reduced in the inflammatory environment. This is not a well-researched area but there are data on the regulation of osteoblast apoptosis in relation to this.⁹² The Wnt family of proteins are involved in cell differentiation and tissue morphogenesis. These mol-

ecules interact with other types of anabolic molecules, such as BMPs, and have received close attention recently as a group of agonistic and antagonistic molecules in bone formation.⁹⁶ Interest was stimulated by the discovery of a mutation in the low-density lipoprotein receptor-related protein 5 (LRP5), which was identified as causal for an autosomal dominant high bone mass trait in a single human family. The same mutation results in increased transcription of OPG in response to loading and may also result in a reduction in osteoclastogenesis. The identification of LRP5/Wnt signaling in bone mechano-sensing has resulted in a new paradigm for understanding bone formation. Since some members of the Wnt (Wnt1 and Wnt5a) and the functionally related frizzled proteins (Fz5) families have been found at elevated levels in RA synovium, it has been proposed that these may play a causal role in synovial hyperplasia and inflammation.⁹⁷ A number of Wnt antagonists have also been identified, including the Dickkopf (DKK) family of proteins. While there is no direct experimental evidence as yet for these molecules as suppressors of bone formation in RA, a potential anti-anabolic role has been proposed for Dkk1 in osteolytic lesions in multiple myeloma,⁹⁸ another disease characterized by increased bone resorption and decreased bone formation.

Inflammatory cytokines may also play an anti-anabolic role in RA. There is abundant literature on the likely pathological roles of TNF- α in pathological bone loss and, while it is recognized that inflammatory cytokines have a role in physiological bone tissue repair and turnover,³² high and/or sustained levels of these may have a net catabolic and anti-anabolic effect on bone tissue.⁹⁹ There is also evidence that TNF- α , for example, might reduce bone formation, by inhibiting the differentiation of osteoblasts.¹⁰⁰ A possible mechanism for this is the inhibition by TNF- α of expression of the transcription factor RUNX2/Cbfa1, a critical regulator of osteoblast differentiation.¹⁰⁰ In RA, and in other states of TNF- α excess, TNF- α may therefore decrease the phenotype progression of precursor cells to the osteoblast pathway. Since TNF also promotes osteoclastogenesis (see above), TNF- α suppression of RUNX2/Cbfa1 could decrease bone formation while simultaneously increasing resorption.

Treating bone loss in RA

There are now a large number of different drugs available to treat RA; however, most of these treatments are aimed at reducing inflammation and very few treatments target the debilitating bone loss seen in RA. Now that the key factors regulating osteoclast resorption in RA are known, new therapies specifically targeting bone destruction are being developed. The RANK-RANKL interaction in the formation of osteoclasts is an ideal target for therapy because it is a point at which numerous pathways for osteolysis converge. Accordingly, treatments based on the inhibition of RANKL inhibition by its natural inhibitor, OPG,³⁷ or soluble RANK,¹⁰¹ have been successful. Combinations of

therapies that target several factors, such as TNF- α , IL-1 β , and RANKL inhibitors have also been used in an animal model of RA¹⁰² and may be more successful than single therapies. Like anti-TNF treatment, methotrexate, sulfasalazine, and IL-4 have recently been shown to inhibit human osteoclastogenesis, and can inhibit RANKL mRNA expression by RA synoviocytes.¹⁰³ Selective blockade of potassium channels may also reduce osteoclast formation in T-cell driven osteolysis in inflammatory diseases. These recent animal studies are promising, and preliminary studies based on selective blockade of RANK-RANKL interaction demonstrate the importance of RANKL in bone destruction in RA.¹⁰⁴

Bisphosphonates, a group of drugs currently used to inhibit osteoclast activity in osteolytic bone tumors and osteoporosis, have also been used to inhibit osteolysis in animal models of osteolysis.¹⁰⁵ In a recent study,¹⁰⁶ zoledronic acid was tested as a potential treatment for the joint destruction in the transgenic mouse RA model produced by over-expression of human tumor necrosis factor (hTNF α). Untreated, these mice developed severe destructive arthritis as well as osteoporosis. In contrast, bone erosion was retarded by a single dose of zoledronic acid and was almost completely blocked by repeated administration of the drug. Although cartilage damage was partly inhibited, synovial inflammation was not affected by zoledronic acid treatment. These findings are in agreement with several other studies,^{107,108} although the success of bisphosphonates in preventing joint damage may depend on the disease model or on the particular bisphosphonate used. For example, in adjuvant-induced arthritis in rats, it was found that while clodronate decreased bone deformation, alendronate did not.¹⁰⁹ However, both bisphosphonates prevented systemic bone loss and strength reduction in this study. In the hTNF α model, calcitonin had no effect on synovial inflammation, bone erosion, cartilage damage, or systemic bone mass, whereas anti-TNF treatment entirely blocked synovial inflammation, bone erosion, synovial osteoclast formation, and cartilage damage, but had only minor effects on systemic bone mass.¹⁰⁶

As alluded to above, other potentially rewarding approaches in treating RA may include the use of antiangiogenic agents, as has been trialed in other diseases of bone destruction. For example, in multiple myeloma it has been demonstrated that patients with active disease have an increase in bone marrow angiogenesis that correlates with the progression of disease and an adverse prognosis. Accordingly, therapeutic approaches to target VEGF, as well as a range of other angiogenesis-related factors, are being investigated for this disease.¹¹⁰ Vascular endothelial growth factor also shows promise as a target in solid tumors. A humanized anti-VEGF monoclonal antibody, bevacizumab, resulted in tumor growth inhibition of human tumor cell lines implanted into nude mice and changes in vascular functions were frequently reported, including decreased vessel diameter, density, and permeability in response to treatment.¹¹¹ Safety studies with bevacizumab in cynomolgus monkeys showed that this agent is generally well tolerated, supporting similar studies in RA.

Conclusions

Bone destruction in the joints of RA patients occurs through similar mechanisms as in other inflammatory diseases. The RANKL/RANK pathway of osteoclast differentiation is central to the formation of osteoclasts at these sites of inflammation and is therefore an important target for the pharmaceutical regulation of bone loss in several common inflammatory diseases. Other inflammatory cytokines are likely to augment the RANKL/RANK pathway of osteoclast formation and activity, and some of these factors may also prolong the survival of osteoclasts and decrease the differentiation and survival of osteoblasts. In addition, increased blood flow to the inflamed pannus is likely to promote bone resorption and also represents a potential target in this disease.

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