

THE destruction of joints caused by rheumatoid arthritis and osteoarthritis is characterized by an imbalance of enzyme catalysed cartilage breakdown and regeneration. A complex cytokine network perpetuates joint conditions by direct regulation of metalloproteases, by indirect recruitment of cells that secrete degradative enzymes, and by inhibition of reparative processes. The destructive action of cytokines such as interleukin-1, interleukin-6 and tumour necrosis factor- α can be modulated at multiple points associated either with cytokine production or with cytokine action. Potential agents for cytokine reduction include selective anti-cytokine antibodies, anti-cytokine receptor antibodies, cytokine receptor antagonist proteins, and soluble and chimeric cytokine receptor molecules. Pharmacologic regulation of IL-1 and TNF α remain primary targets for treatment of arthritis, and results of early clinical trials are promising. However, the results of long-term clinical trials will be required to support the value of anti-cytokine therapy in treatment of arthritis.

Key words: Acute phase proteins, Adhesion molecules, Antiarthritic molecules, Cartilage matrix, Interleukin-1, Interleukin-6, Metalloproteases, Osteoarthritis, Rheumatoid arthritis, Tumour necrosis factor- α

Cytokine reduction in the treatment of joint conditions

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Introduction

Slow, steady destruction of painful, swollen joints is the hallmark of connective tissue disorders such as rheumatoid arthritis (RA) and osteoarthritis (OA). RA, an autoimmune disease, involves T and B cell infiltration of the synovial lining and extensive proliferation of synovial lining cells, resulting in the formation of pannus and influx of polymorphonucleocytes (PMNs) and monocytes into both the synovium and synovial fluid. There may be considerable bone loss, particularly at the margins of the synovial lining, and joint deformity.^{1,2} OA is primarily characterized by loss of articular cartilage. Synovitis may play a role, particularly in painful joints, but bone loss and pannus formation are uncharacteristic. However, cartilage and bone parameters are not normal either in RA or OA. RA and OA are at opposite ends of an inflammation spectrum, in that RA affects multiple joints of the body, involves a large-scale systemic response and is immunologically driven. Cartilage loss in RA proceeds from the invading edge of the pannus. In OA cartilage loss appears to be primarily driven by the cartilage itself. In both cases, joint destruction is characterized by an imbalance of enzyme catalysed cartilage breakdown and regeneration.

While both OA and RA are characterized by increased amounts of metalloproteases in synovium and cartilage, the synovial involvement is far more pronounced in RA. Thus the pathway of joint destruction in RA is thought to overlap the events causing OA, but to be broader in scope and to engage more members of the cytokine network.^{3,4} The aetiology of OA is thought to involve precipitating events ranging from crystal deposition, joint neuropathy, sub-chondral sclerosis, mechanical defects, metabolic abnormalities, to mechanical trauma. However, alterations in chondrocyte metabolism are thought to be at the root of OA of various origins.⁵

Various members of the cytokine network have been implicated in joint destruction both by direct regulation of metalloproteases, by indirect recruitment of cells that secrete degradative enzymes, and by inhibition of reparative processes. Among the most well characterized are IL-1, IL-6, TNF and chemokines such as IL-8. The destructive action of cytokines can be modulated at multiple points associated either with cytokine production, including transcription, translation, secretion and degradation; or with cytokine action, including inhibition of receptor binding and signal transduction. In this review, the potential of cytokine modification for treatment of the

destruction of joints accompanying OA and RA will be considered.

Conditions of joints in osteoarthritis and rheumatoid arthritis

Osteoarthritis (OA) is defined as a complex of interactive degradative and repair processes in cartilage, bone and synovium, with secondary components of inflammation. The aetiopathologic processes involved are complex, and their relative importance continues to be debated.

By currently held concepts, two general pathways lead to OA.⁶ The first involves fundamentally defective cartilage with biomaterial properties directly or indirectly leading to OA. Thereby, the cartilage matrix fails under normal loading of the joint. A recently described type II collagen defect well exemplifies this pathway;⁷ following biomechanical failure osteoarthritis ensues. The second, and by far most prevalent, concept of the cause of OA is based on the major role that physical forces play in causing damage to normal articular cartilage matrix. First, there is direct injury of the matrix; secondly, chondrocytes embedded in the matrix are injured by the same forces.⁸⁻¹¹ In the course of time, these chondrocytes react to injury by elaborating degradative enzymes and developing inappropriate repair responses.^{11,12} Recent research implicates the enzymatic breakdown of cartilage as a key feature of disease progression.¹¹

OA is characterized by the increasing degeneration of articular cartilage, a thickening of the subchondral bone, and the formation of marginal osteophytes. Biochemical and histological studies indicate that there is focal loss of extracellular ground substance in the matrix of OA lesions. As the disease progresses, there are ulcerations in the cartilage and, finally, the entire articular surface is lost.¹³ In OA, changes not only involve the cartilage, but other joint structures such as the synovial membrane.¹⁴

Synovial inflammation is responsible for some OA symptoms and is also likely to play an important role in the pathological process by interacting with, and thereby accelerating, catabolism.¹¹ The order in which the biochemical changes take place during the destructive phase of this disease is not yet clear. However, one of the main biochemical changes in the articular cartilage appears to be related to alterations in the proteoglycan structure. These macromolecules undergo quantitative and qualitative changes.^{13,15} There is a progressive depletion of cartilage proteoglycan, which parallels the severity of the disease.¹⁶ At a certain stage, the chondrocytes appear unable to fully compensate for the proteoglycan depletion resulting in a net loss of matrix. The structural changes of the proteoglycan macromolecules include a decrease in hyaluronic

acid content, a diminution in the size of proteoglycan aggregates and monomers, and a decrease in the aggregation properties of the monomers.¹⁶⁻¹⁸ The latter changes seem likely to reflect the degradation of the proteoglycan monomer core protein in which cleavage has occurred in several areas, including the hyaluronic acid binding region (HABR).¹⁹ Although the content of type II collagen remains unchanged in OA cartilage, the increased cartilage hydration and the ultrastructural changes of the collagen fibres²⁰ represent important alterations in the collagen fibre network. The increase in minor collagen types such as type I collagen, particularly in the pericellular area, suggests a change in chondrocyte metabolism.²¹ These changes in the proteoglycan content of the matrix, together with the damaged collagen structure, lead to a functional deterioration of the cartilage, making it less resistant to compression and other mechanical stress, which lead to the appearance of progressive cartilage lesions.

In OA, mechanical factors and enzymatic pathways are both involved in cartilage matrix degradation.¹¹ The enzymatic process appears related not to a unique system, but rather to a cascade of events.¹¹ In contrast to rheumatoid arthritis (RA), where the synovium is probably the most important source of degradative enzymes, chondrocytes seem to be the most important enzymatic source responsible for OA cartilage matrix metabolism. Currently, the enzyme families that have been identified as playing a significant role in OA pathophysiology are the metalloproteases, the serine proteases and the thiol proteases.

The main metalloproteases involved in cartilage matrix degradation are collagenase, stromelysin and gelatinase.²²⁻²⁴ Collagenase appears to be responsible for the breakdown of the collagen network in OA cartilage. An increased collagenase level has been identified *in situ* in human OA cartilage²⁵ as well as in the experimental dog OA model.²⁰ In addition, the collagenase level was found to correlate with the severity of OA cartilage lesions.²⁵ Stromelysin has been identified in human articular cartilage, and its level also correlates with the severity of OA lesions.^{23,24} An increased level of collagenase and stromelysin mRNA has been found in OA chondrocytes,²⁶ and their levels enhanced in OA synovial fluid.²⁷ Furthermore, active stromelysin can mimic *ex vivo* and *in vitro* the breakdown of the proteoglycan monomer core protein, including HABR cleavage seen in OA.^{17,18,28} Histochemical studies²⁴ have also revealed a correlation between the level of stromelysin and the degradation of pericellular proteoglycan. Degradation of types II and IX collagen was also reported to occur through stromelysin. Moreover, stromelysin may play a dual role in OA pathophysiology; on the one hand, by degrading matrix macromolecules itself and, on the

other, by activating procollagenase.²⁹ The 92 kDa gelatinase is selectively expressed in fibrillated cartilage and is also likely involved in OA cartilage degradation.²² The biological activity of metalloproteases is controlled by both physiological inhibitors and activators. At least two tissue inhibitors of metalloproteases (TIMP-1 and -2) are known to exist in humans. In OA cartilage, there is an imbalance between the synthesis and activity of TIMP and metalloproteases corresponding to a relative deficit in the amount of the inhibitor,^{30,31} favouring an increased level of active metalloproteases and secondarily matrix degradation.

Serine and thiol dependent proteases, including the plasminogen activator (PA)/plasmin system and cathepsin B respectively, have both been suggested as possible activators of metalloproteases.^{32,33} The plasmin system was shown to be involved in the activation of latent metalloproteases during *in vitro* studies, demonstrating that plasmin activated RA synovial collagenase.³³ However, a later study showed that complete collagenase activation requires, in conjunction with plasmin, the presence of active stromelysin.²⁹ To date, very few studies have addressed the involvement of PA/plasmin in OA pathophysiology. A recent report indicates that OA cartilage contained an increased level of plasmin associated with an increased synthesis in PA (urokinase type).³⁴ This study also revealed that a correlation exists between the level of plasmin and active collagenase in OA cartilage showing severe lesions.³⁴ Moreover, one of the major physiological inhibitors of plasminogen activators, PAI-1, was found to be markedly decreased in OA cartilage.³⁴ These findings, together with the increased level of PA, may partly explain the increased level of biologically active metalloproteases in OA tissue. Degradation of the extracellular matrix macromolecules often occurs in the pericellular area around the chondrocytes, where the matrix pH is in the acid range. At first, cathepsin D was thought to be the prime candidate for causing matrix degradation. Although cathepsin D is elevated in OA cartilage, it does not seem to be involved in cartilage resorption.³⁵ However, cathepsin B, another lysosomal enzyme, is likely to play an important role in cartilage degradation through its direct degradative effect on both collagen and proteoglycans, and also by activating metalloproteases.³⁶ Although cathepsin B is maximally active at pH 6.0, this enzyme can also exert proteolytic activity for a limited time at neutral pH.³⁶ As in several other human enzyme systems, the proteolytic effect of cathepsin B is regulated by specific protease inhibitors. Two such inhibitors, with molecular weights of 67 kDa and 13–16 kDa, have been found in articular cartilage.^{37,38} It appears that the small inhibitors are forms of cystatin and the large inhibitor is a kininogen.^{39,40} In OA cartilage, the

cathepsin B level is increased, showing higher activity in cartilage lesions with a concomitant decrease in cysteine protease inhibitory activity.³⁷ This imbalance between cathepsin B and cysteine protease inhibitor levels may be an important contributing factor in OA cartilage degradation.

Rheumatoid arthritis (RA) is an autoimmune disorder characterized by a chronic, erosive synovitis of joints.⁴¹ The cause of RA is unknown, and it is in fact possible that there are several causes for the disease. Infectious agents remain a focus of suspicion; however, there is no direct evidence for their involvement. An interrelationship between infectious agents, genetics and autoimmunity has also been suggested.⁴²

In RA, the disability is due to damage to joint structures such as the capsule, ligaments and erosion of cartilage and bone. The initial pathologic changes in early RA happen at the synovial microvascular level with an activation and swelling of endothelial cells.⁴³ Plasma exudation which follows is reflected by the development of oedema in the subsynovial lining tissue. The cells in the synovial lining cell layer become activated, and their numbers are greatly increased. This lining consists of three cell populations: phagocytic cells of the monocyte–macrophage lineage, dendritic cell and fibroblast-like cells.⁴⁴ A small number of polymorphonuclear leukocytes are also observed at the superficial layer of the synovium. A large accumulation of mononuclear cells is seen around blood vessels and in the sublining synovial tissues. The sublining tissue is infiltrated with nodular collections of mononuclear cells, particularly around blood vessels. Activated T lymphocytes are predominant in focal aggregates and plasma cells at the periphery of the nodule. More diffuse collections of mononuclear cells consist of macrophages, T cells, B cells and plasma cells.⁴⁵ At chronic stages of the disease, the synovium becomes hypertrophic and villous projections of synovial tissue invade the joint cavity. The mass of inflammatory cells may invade over the surface of the articular cartilage (pannus) or may burrow into the subchondral bone. Joint destruction occurs predominantly in areas adjacent to the margin of the invading pannus.⁴⁶

The destruction of the joint probably results from the production of an excessive amount of degradative enzymes. The major producer of enzymes is likely to be the synovial fibroblast at the synovial lining level. Polymorphonuclear leukocytes can also release several proteinases, and probably contribute to the proteolytic activity found in inflammatory synovial fluids.⁴⁷ Activation of neutrophils also results in the release of reactive oxidants which contribute to inflammation and injury of joints. The neutrophil proteases include elastase (serine protease), gelatinase and a collagenase

(metalloprotease); the latter is distinct from the collagenase synthesized by fibroblasts.^{48,49} In the synovial lining cells, both the macrophages and fibroblast-like cells express collagenase and stromelysin.⁵⁰⁻⁵²

Interactions of cells with the cartilage matrix molecules are also important in the regulation of cellular activity in inflammation. The production of cytokines such as IL-1 is increased when mononuclear cells are exposed to several types of collagens such as types II, III and IX.^{53,54} Fragments of type I collagen as well as other constituents of the bone matrix can stimulate production and release of IL-1 by monocytes.

Autoimmunity to cartilage matrix molecules may also play a role in cartilage destruction. Antibodies to cartilage collagens (types II, IX and XI) are commonly present in patients with rheumatic diseases.^{55,56} The collagen fragments released from cartilage can possibly be recognized by the immune system as foreign proteins. Immune complexes containing antibodies to type II collagen have been observed in the superficial zone of articular cartilage in patients with RA.⁵⁶ The appearance of these antibodies occurs after articular cartilage damage has begun. The loss of cartilage may be accelerated by the deposition of immune complexes in the superficial layers of the cartilage, which favours invasion of the pannus.⁵⁷ Neuropeptides may play a role in the modulation of the inflammatory response.⁵⁸ Substance P, a peptide located in peripheral nerves, has been shown to induce the production of cytokines (IL-1, TNF α and IL-6) by mononuclear cells⁵⁹ and metalloproteases and prostaglandins by synovial fibroblasts.⁶⁰ This finding emphasizes the complexity of factors involved in the pathogenesis of RA.

It is likely that important interactions occur between monocyte/macrophages and synoviocytes resulting in the regulation of enzyme release, cytokine formation and cell proliferation. It is suggested that while immune mechanisms may initiate early stages of the disease, synoviocytes and macrophages are independently capable of maintaining a destructive phenotype and this may contribute to the chronic nature of RA.

Role of cytokines in the destruction of joints

The spectrum of factors responsible for the altered function of synovium and cartilage in conditions such as OA and RA has not been fully defined. Prevailing theory dictates that the perpetuation of the above arthritic diseases is likely to be controlled by a complex cytokine network, in which three of them, IL-1, TNF α and IL-6 appear to be of major importance. These cytokines are soluble molecules that transmit information between cells. IL-1, TNF α and IL-6 have been detected in synovial fluid, synovium

and cartilage from RA patients, and IL-1 and IL-6 in the latter tissues from OA patients. The major source of these cytokines in synovium is believed to be monocytes/macrophages; however, current evidence suggests that synovial lining cells (synoviocytes) also produce these cytokines. Findings suggest that IL-1 may be secreted by chondrocytes, whether this represents autocrine and/or paracrine stimulation is not yet known. It may be that this cytokine diffuses through the synovial fluid into the cartilage thus inducing chondrocytes to produce IL-1. Also, IL-6 is believed to be produced by chondrocytes.⁶¹

The three cytokines, IL-1, IL-6 and TNF α are synthesized and released as part of the response of their cells of origin to specific signals, and they influence the response and function of their target cells, largely by exerting a positive or negative influence on gene expression (Fig. 1). One important principle is reflected in their ability to cause multiple effects, overlapping and synergizing with other cytokines. The action of IL-1 and TNF α on joint tissues are multifaceted, with many different gene products being influenced either by stimulation or suppression. The association of these cytokines with tissue damage arises from their propensity to stimulate the proteolytic pathways of extracellular matrix degradation and, at the same time, subdue the synthetic pathways leading to new matrix formation.

It is likely that IL-1 is responsible for increasing the protease synthesis in diseased synovium, as metalloprotease production correlates with the severity of synovial inflammation⁶² and the latter with the level of IL-1 in the synovial fluid⁶³. The capacity of IL-1 produced by the inflamed synovium to stimulate the production of collagenase and PA by synovial fibroblasts has been well documented.⁶⁴⁻⁶⁶ In synovium, protease inhibitors, such as TIMP or PAI-1, are either suppressed or unaffected by the cytokine IL-1. Mediators of the inflammatory process such as prostaglandin E₂ and IL-6 are both stimulated by IL-1. As synoviocytes secrete IL-1, it is tempting to speculate that autocrine stimulation may also play a role in the regulation of synoviocyte enzyme synthesis. IL-1 may also contribute to the fibrosis observed in arthritic synovium, as it increases the synthesis of types I and III collagens by synovial fibroblasts.^{67,68}

Many of the effects of TNF α on synovium overlap with those of IL-1. In synovial tissue TNF α stimulates collagenase and prostaglandin E₂ production, displays synergism with other cytokines and, in certain circumstances, induces IL-1 production.

The exact role of IL-6 in arthritic synovium has not yet been clearly defined. The ability of IL-1 and TNF α to induce IL-6 protein and mRNA in synovial fibroblasts⁶⁹ suggests that this cytokine may be an important intermediate signal in the induction of other cellular responses to these cytokines. How-

Potential target sites for treatment of joint conditions

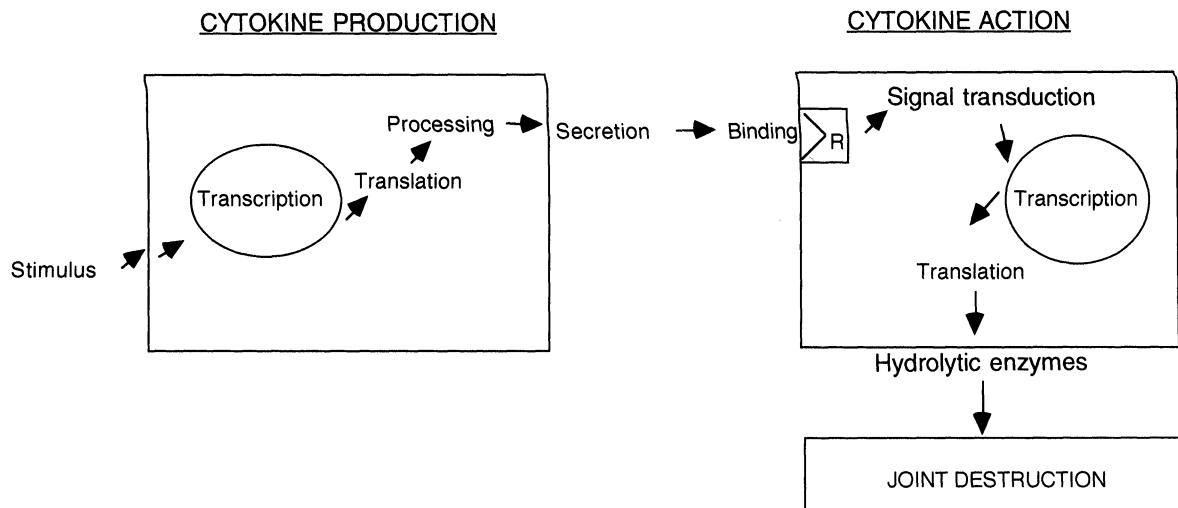


FIG. 1. Diagrammatic representation of potential sites at which the destructive activities of cytokines can be modulated. The biological effects of cytokines can be modulated at points associated with cytokine production, including transcription, translation, secretion and degradation; or at points associated with cytokine action, including inhibition of receptor binding, signal transduction or inhibitors of hydrolytic enzymes.

ever, IL-6 has no direct effects on the synthesis of proteases, prostaglandins or matrix proteins, but stimulates synthesis of TIMP.⁷⁰ It is suggested that IL-6, by its *in vivo* activation of B cells, may contribute to the immunologic phenomenon; its role in inflammation may be via regulation of changes in the concentrations of acute phase proteins.

Another important effect of cytokines on the development of synovitis is their contribution to the expression of adhesion molecules, which in turn control the accumulation of leukocytes. Evidence exists that IL-1 and TNF α are involved in the early events affecting the joints by their contribution to the migration of cells to the site of inflammation and their stimulation of a variety of responses in endothelial cells. At present, four families of adhesive molecules have been described:⁷¹ the immunoglobulin supergenes, the integrins, the cadherins and the syndecans. The cytokines IL-1 and TNF α were recently shown to upregulate the expression of some of the members of the first family, the ICAM, present on fibroblast and endothelial cells.⁷² It is likely that increased ICAM expression plays an important initial step in the binding of leukocytes to synovial endothelial cells in RA joints⁷³ and facilitates the subsequent entry of these cells into the inflammatory synovial fluid. Yet no effect by these cytokines has been demonstrated on members of the other adhesion molecule families. In synovial fluid, as well as in the synovial tissues, polymorphonuclear leukocytes are attracted and are believed to play a role in tissue destruction by their release of cytokines and degrading proteases. Evidence is mounting that a new cytokine superfamily, known as the intercrine (or chemokine), the members of which have the ability

to mediate the recruitment of leukocytes, is likely to play an important role in inflammation.^{74,75} As these intercrines exhibit differing patterns of specificity for various leukocyte populations, this makes them attractive candidates as important components of the inflammatory process. Members of this family are classified as two groups according to the position of the first two cysteines in the conserved motifs. Among the first group are IL-8 and MCP-1; RANTES belongs to the second group. The members of the first group appear to be chemotactic for neutrophils, but not for mononuclear leukocytes, whereas the second group members attract mostly mononuclear cells and granulocytes. Recently, it was shown the MCP-1 and IL-8 are expressed in monocyte/macrophages, fibroblasts and endothelial cells in response to the cytokines TNF α and IL-1.⁷⁶⁻⁸¹ Similarly, RANTES expression in fibroblasts was also found to be enhanced by these two latter cytokines.⁸² Although they are present in the synovial environment of RA and OA,^{81,83-85} their diversity of function and their mutual interaction have led to considerable confusion. However, it is important to recognize that they are generated only after specific cell-cell interaction and their activity is limited to the microenvironment immediately surrounding the cell that produced it.

The action of cytokines is not unique to synovial membranes, they also compromise articular cartilage functions. In articular cartilage, IL-1, TNF α and IL-6, appear among the known cytokines to play a major role in the pathological process. In addition to the above-mentioned effects on the synovium, IL-1 suppressed the synthesis of collagen types characteristic of hyaline cartilage (type II and type IX), while

promoting the synthesis of these collagen types characteristic of fibroblasts⁸⁶ (type I and type III), thereby causing the decreased repair of cartilage matrix. The synthesis of the aggregating proteoglycan (aggrecan) characteristic of hyaline cartilage is also suppressed, although those proteoglycan molecules that are synthesized appear to undergo normal post-translational processing.⁸⁷ It has been suggested recently that the effect of IL-1 on the inhibition of proteoglycan synthesis may be mediated by IL-6 in human cartilage, and that IL-1 induces the synthesis and secretion of IL-6 by human chondrocytes.^{88,89} Interestingly, the action of IL-1 on different connective tissue cells does not always produce the same response, i.e. stimulation or suppression. For example, in contrast to the effect of IL-1 on chondrocytes, IL-1 stimulates glycosaminoglycan synthesis by synovial fibroblasts.⁹⁰ This is probably due to the differential expression of aggrecan in synovial fibroblasts and chondrocytes. In cartilage, by far the greatest proportion of glycosaminoglycan is synthesized as aggrecan. However, in synovium, aggrecan is not significantly expressed, but decorin is abundant, and the latter gene is upregulated by IL-1.⁹¹ The effects of IL-1 on cartilage proteins are not limited to inhibition of synthesis, as prostaglandin E₂ production is stimulated by this cytokine.⁹² This stimulation of prostaglandin production is of particular interest because of the role it may play in exacerbating joint inflammation, stimulating bone resorption and modulating the immune response. Finally, IL-1 may be involved in osteophyte formation, by stimulating the proliferation of human osteoblast-like cells causing increased bone formation.

In cartilage, TNF α appears to induce many effects analogous to those generated by IL-1, although the former is generally less potent in its effect than either form of IL-1.⁹³ Although the effect of TNF α on chondrocytes has been less well studied than that of IL-1, it is clear that TNF α , like IL-1, can stimulate the production of proteolytic enzymes such as collagenase, stromelysin, elastase and PA, as well as prostaglandins E₂ and IL-6. TNF α also has, however, no effect on TIMP production by articular chondrocytes. TNF α also modulates the synthesis of cartilage matrix and, like IL-1, suppresses aggregating proteoglycan synthesis⁹⁴ and selectively decreases the production of cartilage type I and type II collagens. It suppresses the expression of type II collagen, but increases type I and III collagen gene expression.⁹⁵

In contrast to its detrimental role of inhibiting proteoglycan synthesis,⁸⁸ IL-6 does not appear to influence the production of the metalloproteases, either alone or in combination with IL-1.^{96,97} Indeed, it appears to stimulate the production of TIMP.^{70,97} In this manner, IL-6 production would counteract the degradative potential of IL-1. In cartilage, the involve-

ment of IL-6 in the proliferation of chondrocytes as well as clones is plausible, as this cytokine has been shown to enhance human OA chondrocyte proliferation.⁹⁸

Finally, the process involved in the inhibition/activation process of metalloproteases in arthritic joint tissues could very well be modulated by IL-1. For instance, the imbalance in the TIMP-1 and metalloproteases levels³¹ in articular cartilage may be mediated by IL-1, as *in vitro* experiments showed that increasing concentrations of IL-1 produced decreased TIMP-1 synthesis in parallel with increased metalloprotease synthesis in articular cartilage and chondrocytes.^{30,34} PA synthesis is also modulated by IL-1. *In vitro* stimulation of cartilage chondrocytes with IL-1 showed a dose dependent increase of the PA, concomitant with a sharp decrease in PAI-1 synthesis.⁹⁹⁻¹⁰¹ The potent inhibitory effect of IL-1 on PAI-1 synthesis, in combination with a stimulatory effect on PA synthesis, is a powerful mechanism for regulation of the generation of plasmin and metalloprotease activation. In addition to its role as an enzyme activator, plasmin may also be involved in cartilage matrix degradation by direct proteolysis of the proteoglycan monomer.¹⁰²

Potential stages for cytokine reduction

Cytokine production is associated with disturbances of homeostasis ranging from acute conditions such as sepsis to chronic connective tissue disorders such as RA and OA. Distinctly different strategies of cytokine modification are required for blocking the effects of cytokines produced locally and chronically in joint conditions compared with those employed for short periods when relatively high systemic levels are produced, as in sepsis.¹⁰³ Local elevation of cytokines in joints during RA and OA may result in minimal increases in systemic blood levels. Thus TN α , IL-1 and IL-6, if detectable, are much lower than in sepsis, making direct measurement of cytokines difficult.¹⁰⁴

Cytokine production can be assessed indirectly by their biological effects. One of the best measured sequelae of cytokine production and action is a marked change in the pattern of hepatic synthesis of plasma proteins. Plasma proteins whose synthesis is responsive to cytokines (or secondarily to inflammation, trauma or sepsis) are known as acute phase proteins or acute phase reactants. Two inducible acute phase proteins, in particular, serum amyloid A (SAA) and C-reactive protein (CRP), whose synthesis is regulated by synergism of IL-1 and IL-6,¹⁰⁵⁻¹⁰⁷ have prognostic value in the clinical management of arthritis. The concentrations of IL-1 and IL-6 in plasma of RA patients are considerably lower than that predicted by *in vitro* studies to be required for stimulation of SAA and CRP production. Thus the nature of

the signals to liver for acute phase protein synthesis are not completely understood and are the subject of active investigation in several laboratories. One possibility is that the signals from inflamed joints may not be IL-1 and/or IL-6, but rather TNF α or another signal that stimulates IL-1 and/or IL-6 production by Kupffer cells. Hepatocytes adjacent to the cytokine-producing Kupffer cells are then able to respond. A second possibility is that inflammatory cells may travel from inflamed joints to the liver where they come into contact with hepatocytes and then locally produce the cytokines by which hepatic cells are stimulated. A third possibility is that mediators that are currently unknown may travel to the liver to stimulate SAA and CRP synthesis. Although the exact pathway by which inflamed joints signal changes in hepatic protein synthesis is unclear, it has been recognized for more than 20 years that elevated concentrations of acute phase proteins are associated with a poor clinical prognosis and more recently, that reduction of disease activity by anti-rheumatic therapy is accompanied by reduced plasma concentrations of acute phase proteins such as SAA and CRP.¹⁰⁸⁻¹¹³ Measurement of acute phase proteins thus provides a strategy by which therapies which inhibit cytokine synthesis or action in the treatment of joint conditions can be monitored.

Several cytokines have been detected in high concentrations in RA synovial fluid and synovial membrane including IL-1, IL-6, TNF α ,¹¹⁴⁻¹²³ GM-CSF,¹²⁴ IL-8⁸¹ and TGF β .^{118,125} IL-1, TNF α and IL-6 are not only found in RA synovium, but have been detected in serum and synovial fluid of arthritis patients.^{126,127} Both IL-1 and TNF α are thought to play a significant role in the pathogenesis of arthritis, but cannot totally account for the pathology of RA and OA.¹²⁸ The leukocytic infiltration into the synovial lining and the synovial fluid is thought to be due to production of other mediators such as GM-CSF and IL-8, secondary to IL-1 or TNF α . Cytokine production and action may play a role in arthritic joint conditions in a number of ways, including disease induction, chronicity and exacerbation. Thus cytokine blocking agents will be of broad value in treating joint conditions. There are several stages at which cytokine production can be blocked, including transcription, translation, secretion and action. Agents that reduce the synthesis and action of cytokines known to be involved in joint disease will be reviewed briefly in the following sections.

Tumour necrosis factor- α : RA is characterized by cellular activation and TNF α production.¹¹⁸ TNF α , originally identified for its anti-tumour activity and subsequently for its cachectic activity, is known to influence inflammation and cellular immune responses.¹²⁹⁻¹³³ Transgenic mice that produce large amounts of human TNF α develop arthritis,¹³⁴ and

TNF α administered into knee joints of animals causes an acute arthritis.^{135,136} TNF α , mainly derived from monocyte/macrophage derived cells, signals its responses through two distinct cell surface receptors, TNF-R55 and TNF-R75.¹³⁷⁻¹³⁹ While TNF-R55 and TNF-R75 differ in the primary structures of their cytoplasmic domains, both are thought to bring about cellular alterations by altering gene expression through activation of various nuclear transcription factors, which act as a link between cell membrane and nucleus. TNF-R are localized in synovial tissue and the cartilage-pannus junction in RA patients to a much greater extent than in OA patients.¹⁴⁰

The single 3 000 bp TNF α gene consisting of four exons is on chromosome 6. Induction of TNF α synthesis is under both transcriptional and post transcriptional regulation.¹⁴¹ Mature TNF α , 17 kDa, 157 amino acids in length, is derived from a 26 kDa transmembrane precursor protein¹⁴² by cleavage of a 76 amino acid region from the amino terminus; the amino terminal region is thought to anchor the TNF α precursor in the plasma membrane. Native biologically active TNF α is a trimer with a three-dimensional structure similar to viral capsids.

Because TNF α is produced locally at sites of inflammation and plays a pivotal role in the cytokine network, it is desirable to prevent TNF α production at the earliest stages. One promising technique is the application of ribozymes, which have been designed to cleave target viral RNAs including HIV-1, and are considered to show promise as *in vivo* therapeutic agents if obstacles to co-localization with their substrates can be overcome.¹⁴³ In particular, it has been thought that the compartmentalization of RNAs in cells would limit the efficacy of ribozymes due to reduced diffusion.¹⁴⁴ Local administration of ribozymes specific for TNF α to the joint has been proposed,¹⁴⁵ and preformed ribozymes have been shown in a model system to reduce TNF α mRNA and production by 90% and 85%, respectively.¹⁴⁵

Several approaches have been used to inhibit the interaction of TNF α with its receptors. Suramin, which has been used experimentally to block ligand-receptor interactions, has been found to inhibit the biological activity of human TNF α through direct action on the ligand, i.e. by dissociation of the quaternary structure of TNF α .^{146,147} Recently, a chimeric monoclonal antibody to TNF α was used to treat patients with RA¹⁴⁸ with significant clinical improvements and changes in laboratory parameters including decreased CRP, SAA and IL-6 concentrations. Since CRP and SAA both appear to be stimulated by synergism of IL-1 and IL-6 and TNF α alone is less potent, this finding is consistent with the concept that TNF α is an early mediator that acts by way of IL-1 and IL-6.³

TNF α inhibitory proteins identified in serum and urine, were found to be fragments of the extracellular

portions of TNF receptors.¹⁴⁹⁻¹⁵³ These TNF α binding proteins can reach concentrations of greater than 2 ng/ml and are thought to neutralize TNF bioactivity or to act as reservoirs from which TNF can slowly be released. Recombinant soluble TNF receptors prevent development of experimental collagen arthritis¹⁵⁴ as do anti-murine TNF α antibodies.^{154,155}

Interleukin-1: As discussed above, IL-1 is a mediator of arthritis with activity at many points in the pathogenesis of the disease, e.g. capable of inducing synthesis of metalloproteinases implicated in cartilage loss, adhesion molecules necessary for the migration of inflammatory cells into inflamed tissue, and synthesis of cytokines such as IL-6, IL-8 and GM-CSF. Animal studies have shown that IL-1 directly injected into knee joints will cause arthritis, will induce a flare of quiescent, pre-existing arthritis and will increase incidence and severity of arthritis in prearthritic animals.¹⁵⁶⁻¹⁵⁸

Two distinct forms of human IL-1 have been characterized, IL-1 α and IL-1 β (reviewed in References 159 and 160). The two forms have ~27% amino acid identity and are the products of separate genes. The primary IL-1 gene products are 31 kDa proteins of about 270 amino acids and are precursors of the active species designated IL-1 α (159 amino acid residues) and IL-1 β (153 amino acid residues). The IL-1 precursors lack classical hydrophobic leader sequences and the 17 kDa mature IL-1 carboxyl peptides are formed by action of specific proteases upon secretion.

The human IL-1 α gene is 10 kb in length and the IL-1 β gene is ~7 kb long. Both are on chromosome 2 and are comprised of seven exons and six introns. Pro-IL-1 gene expression is regulated both transcriptionally and post-transcriptionally.^{161,162} The IL-1 genes are under specific transcriptional control which is influenced by cell type and the inducing agent, and modulated by other cytokines. The differential expression of IL-1 α and IL-1 β may be explained in part by the marked structural differences between the two promoters.¹⁶³

The biological properties of IL-1 are similar to and overlap those of TNF. In some situations, potentiation or synergism between IL-1 and TNF occurs, an effect that seems to derive from signal transducing molecules rather than upregulation of receptors, since IL-1 down-regulates TNF receptors.^{164,165}

As reviewed by Fenton,¹⁶² transcription, translation and release of IL-1 are distinct, dissociable processes. Secretion and processing of pro-IL-1 β appear to be linked. There are redundant mechanisms for controlling IL-1 production, and, once produced, there are multiple mechanisms for regulating the effects of IL-1 (reviewed in Reference 160). These include the

number of and the presence of receptors on target cells, potential production of soluble receptors and synthesis of inhibitors.

There are two Rs for IL-1, the 80 kDa type I and the 60 kDa type II. In type II IL-1R, the cytoplasmic domain portion is shorter than for type I IL-1R. Antibodies to the type I IL-1R block the biological effects of IL-1 whereas type II IL-1R appears to be a decoy receptor.¹⁶⁶ In contrast to antibodies to TNF receptors, antibodies to IL-1Rs have not yet been associated with agonist activities.^{103,150}

Soluble shed type I IL-1R has not been described in nature; type II IL-1R is shed.^{166,167} The extracellular domain of type II IL-1R has been cloned and expressed and shown to bind both forms of IL-1. In terms of cytokine reduction, an advantage that sIL-1R (or antibodies to IL-1) would seem to have over IL-1RA is that quantities that are only stoichiometric to IL-1 would be required to reduce IL-1 actions, rather than the 1 000-fold excess that would seem to be required of a competitive inhibitor, as studies in animal models have indicated.¹⁶⁸ The naturally occurring IL-1RA is induced at high levels and competes with IL-1 for binding to its receptor,^{126,169} thereby serving to buffer the intensity of an inflammatory response. Cloned IL-1RA inhibits the activities of IL-1 α and IL-1 β .^{126,141} Production of IL-1 and IL-1RA are regulated differently (reviewed in Reference 103).

IL-1 is a mediator of rheumatoid synovitis and there are several stages at which its action can be blocked. Transforming growth factor- β (TGF β) is known to counteract the effects of IL-1, perhaps by reducing the number of IL-1 receptors or by stimulation of IL-1RA release.¹⁷⁰ In animal models of arthritis, IL-1RA has been shown to inhibit the flare of streptococcal arthritis induced by streptococcal cell walls, and anti-murine IL-1 antibody effectively reversed the inhibition of proteoglycan synthesis and loss of cartilage that accompany monoarticular antigen arthritis.¹⁷¹ An initial study of subcutaneously administered IL-1RA in RA has been reported¹⁷² with greater than 50% reduction in swollen joints and CRP concentration.

Antisense oligonucleotides form duplexes with their corresponding sense mRNAs and prevent transcription.¹⁷³ Antisense IL-1 α inhibits the programmed cell apoptosis of cultured endothelial cells.¹⁷⁴ Antisense inhibition of IL-1R expression is actively being pursued. Because of difficulties in transport, uptake and targeting of antisense oligonucleotides,¹⁷⁵ topical or local administration such as in the cases of psoriatic arthritis would seem the most promising for initial investigation.

Synovial fibroblasts have been transfected with IL-1RA mRNA. Cells constitutively producing IL-1RA have been injected into rabbit knee joints and shown to block an IL-1 induced synovitis.¹⁷⁶ IL-1 β production uniquely requires an enzyme termed pro-IL-1 β

converting enzyme (ICE). Specific inhibition of ICE appears to be a promising approach to arthritis treatment in situations in which only IL-1 β is involved. However, it remains to be determined if there is redundancy of IL-1 α and IL-1 β in the pathophysiology of arthritis.

Interleukin-6: The IL-6 gene is located on chromosome 7 and consists of five exons; expression is induced by agents such as IL-1 and TNF to yield a multifunctional cytokine that acts on many different types of cells.^{103,177,178} IL-6 signals target cells through 80 kDa cell surface receptors¹⁷⁹ which as a complex, IL-6/R, interacts with a 130 kDa protein to initiate signal transduction.¹⁸⁰ Both the 80 kDa R and the 130 kDa signal transducing protein are subject to regulation by inflammatory mediators. For example dexamethasone (Dex) stimulates, while high IL-6 levels down-regulate the 80 kDa IL-6R. In contrast, IL-6, Dex and the combination stimulate the 130 kDa signal transducing protein. A soluble derivative of the 80 kDa receptor found in urine,¹⁸¹ is thought to be generated by limited proteolysis (shedding) of the 80 kDa plasma membrane IL-6R.¹⁷⁹ The soluble IL-6 receptor consists of the extracellular region only and lacks the transmembrane and cytoplasmic region; soluble IL-6R/IL-6 complex acts as an agonist for *in vitro* acute phase protein synthesis. Thus in situations of sustained high IL-6 and down-regulated IL-6R, the soluble IL-6R may have an important physiologic role in modulating the activity of IL-6. The IL-6 receptor subunits belong to a superfamily which includes LIF, oncostatin M, CNTF and GM-CSF.¹⁷⁹ One IL-6 action, fever, is blocked by cyclooxygenase inhibitors indirectly through increased release of arachidonic acid.¹⁰³ Soluble p80R for IL-6 enhances activity.^{181,182}

Cytokine reducing effects of anti-arthritis drugs

The scope, target mechanisms and effectiveness of anti-rheumatic drugs in the pre-cytokine era were thoroughly reviewed by Bonta and colleagues.¹⁸³ The use of cyclooxygenase inhibitors to inhibit the production of prostaglandins has been a major element in the clinical management of arthritis for about 100 years, while our understanding of the role of prostaglandins in inflammation has been developed over the past 20 years.

Non-steroidal anti-inflammatory drugs: The pro- and anti-inflammatory action of PGE has been recognized for a number of years.¹⁸³ Prostaglandins affect inflammation by several mechanisms. They alter blood flow through inflamed areas, potentiate capillary leakage, potentiate sensory fibre pain transmission and, by inducing intracellular cAMP, regulate

synthesis of proteins with cAMP regulatory elements, such as TNF α and IL-2. NSAIDs (aspirin, indomethacin, fentiazac, naproxen, piroxicam) increase TNF and IL-2, cytokines whose synthesis is under regulation by PGE₂;¹⁸⁴⁻¹⁸⁷ however, acute phase protein concentration is usually unaffected by NSAIDs. PGE₂ and other prostanoids inhibit interferon- γ production whereas they are required for PA synthesis.^{186,188-189} NSAIDs appear to act by inhibition of the synthesis of prostaglandins. Recently, two cyclooxygenase enzymes have been identified. The constitutive enzyme is called Cox-I and the second enzyme, termed Cox-II, is induced by cytokines produced at the sites of inflammation. Selective Cox II inhibitors are being designed in the hope of greater selectivity in arthritic disease and decreased gastrointestinal side effects.¹⁹¹

Glucocorticoids: Glucocorticoids, like insulin, exert, at concentrations usually found in plasma, direct regulatory effects on inflamed areas such as joints.¹⁹² Steroid analogues of cortisol, the major active glucocorticoid in humans, such as hydrocortisone and dexamethasone, inhibit release of arachidonic acid from phospholipids of cell membranes by inhibiting the activation of phospholipase A₂. Glucocorticoids inhibit collagen synthesis, and are well known inhibitors of IL-1 production acting at both transcriptional¹⁹³ and post-transcriptional levels.¹⁹⁴⁻¹⁹⁶ Glucocorticoids have also been reported to have a general regulatory effect, inhibiting production and activity of cytokines such as IL-2, -4, -6, -8, GM-CSF and TNF α .¹⁹⁵⁻²⁰¹ IL-10 and dexamethasone inhibit IL-1 and TNF production from LPS-stimulated monocytes.^{194,195,202-206}

Immunomodulators and immunosuppressive agents: RA has been classified as an autoimmune disease since the pathologic manifestations of RA resemble a misdirected immune response. T cells have been implicated in establishment of chronic arthritic disease. Since antigen or mitogen stimulation of T cells leads to increased expression of IL-2R, it has been suggested that removal of activated IL-2R expressing T cells during active disease could result in selective depletion of those T cells involved in the disease process. Furthermore, since RA probably results from specific sets of antigen-responsive cells, selective inhibition of these sets of T cells could provide therapy that does not suppress general immune function. A trial of anti-IL-2R has been reported.²⁰⁷ Chimeric toxin-IL-2 fusion proteins may be useful in RA treatment. A recent study has used oral chicken collagen as treatment in an animal model of arthritis, a treatment based on our knowledge that oral exposure to antigen can result in long-lasting tolerance to that antigen.²⁰⁸ Results to date are limited to animal models for which the inciting antigen is defined.

Immunomodulatory and/or cytotoxic plant alkaloids, are under investigation for their capacity to control the production and action of pro-inflammatory cytokines such as IL-1 and TNF. Compounds derived from extract of plant roots have been tested including tetrandine²⁰⁹ and extracts of *Trypterygium wilfordii* Hook F²¹⁰ including triptolide and triptolide.²¹¹ These compounds are frequently potent in their activity, but remain to be extensively characterized.

Others: A synthetic analogue of fumagillin, AGM 1470 has been shown to inhibit experimental collagen arthritis, presumably by inhibiting endothelial cell growth and thus neovascularization.^{212,213} In animals models of OA, tetracycline shows promise for treatment.^{214,215}

Tenidap and IX-207-887, which have advanced into clinical trials, have been shown to inhibit IL-1 synthesis and/or action.²¹⁶⁻²²⁰ IL-1R in OA are upregulated, whereas TNF-R occur at only low levels²²¹ and tenidap has been shown to down-regulate IL-1R.²²² In *in vitro* studies, tenidap has been shown to inhibit IL-6 production by PBMCs to an even greater extent than IL-1 is inhibited; furthermore tenidap treatment has been shown to lower CRP and SAA concentrations.^{217,223} Tenidap treatment has recently been shown to lower CRP concentrations in OA patients.¹¹³ SKF-86,002 and chloroquine inhibit IL-1 synthesis and/or action.^{196,224} L-709,049 and SK&F 86002 interfere with IL-1 β secretion.²²⁵ Although there are no reports of biological or synthetic IL-8 antagonists, quinolydimethoxyphenylamine (ETH 615) inhibition of IL-8 biosynthesis has been reported.²²⁶

Summary and future perspectives

Based on animal studies, T cell involvement in RA may play a critical role in initiation and perpetuation of disease. As studies with cyclosporin A suggest, T cell selective agents provided one promising avenue of therapy. There is also little question that proteases act directly to cause cartilage destruction in both RA and OA. Selective inhibition of one or more of these enzymes may provide a second avenue for treatment of RA and OA. The involvement of cytokines, particularly IL-1 and TNF, has been strongly supported in animal models of arthritis by therapy using selective anti-cytokine antibodies, anti-cytokine receptor antibodies, cytokine receptor antagonist proteins, and soluble and chimeric cytokine receptor molecules. These studies suggest that IL-1 and TNF may play additive, if not synergistic, roles, since inhibition of either IL-1 or TNF α provides therapeutic benefit. Because of the plethora of potential pathological consequences of cytokine elevation, pharmacologic regulation of IL-1 and TNF α remain primary targets for treatment of arthritis. Early clinical trials of very

selective biological compounds support the therapeutic value of cytokine inhibitors. However, the results of long-term clinical trials of selective pharmacological cytokine inhibitors will be required to define precisely the pathological roles of cytokines in arthritis and the value of anti-cytokine therapy.

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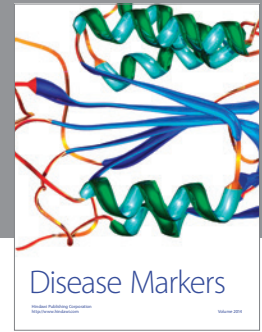
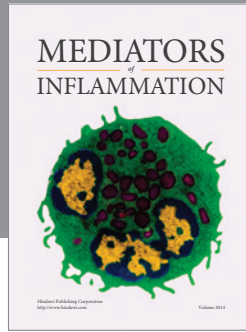
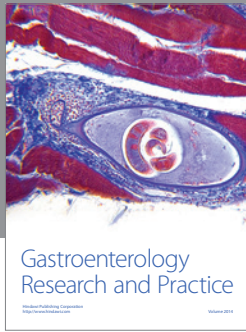
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