Influence of LPS, TNF, TGF-β1 and IL-4 on the expression of MMPs, TIMPs and selected cytokines in rat synovial membranes incubated in vitro

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Abstract. Synovial membranes are formed by four main types of cells, i.e. fibroblasts, macrophages, epitheliocytes and adipocytes. To study the combined effect of various factors on these cell populations, synovial membranes dissected from rat knee joints were incubated in control medium or medium with lipopolysaccharide (LPS), TNF, TGF-β1 or IL-4 for 12 h. LPS stimulated TNF secretion and both agents increased secretion of IL-6. TGF-β1 slightly increased IL-6 secretion. LPS increased the mRNA levels of IL-6, IL-1β, TGF-β1, MMP1a, MMP3, MMP9, MMP13, TIMP14, TIMP1 and TIMP3 while the mRNA levels of MMP2, TIMP2 and TIMP4 were significantly decreased. Expression of IL-1β, MMP1a, MMP1b, MMP3, MMP9, MMP13 and TIMP1 increased after TNF treatment, while mRNA levels of MMP2, MMP14, TIMP2, TIMP3 and TIMP4 were decreased. TGF-β1 increased the mRNA levels of IL-1β, all MMPs, TIMP1, TIMP2, TIMP4 and increased mRNA levels of itself and TIMP3. IL-4 decreased mRNA levels of IL-1β, TGF-β1, MMP2, MMP9, MMP13 and all TIMPs. Only LPS decreased the amount and activity of MMP2. The effect of LPS and cytokines on most of the MMPs and TIMPs produced by whole synovial membrane was in good agreement with previous studies on their action on similar types of cells as those present in synovial membranes, but originating from other tissues. All tested agents decreased MMP2 mRNA expression levels and in the case of LPS also the protein level and its activity determined by zymography, contrary to previous observations on isolated cell populations. This indicates that the response of the organized tissue is an interplay of all components and cannot be deduced from the individual reactions.

Introduction

The synovial membrane defined in anatomy textbooks as the vascular connective tissue which produces the synovial fluid (1) is composed of the synovial lining and the subsynovium (2). The synovial lining, also called the synovial intima (3), consists of macrophage-like type A cells and fibroblast-like B cells (4-6). It is supported by the highly vascularized subsynovium, consisting of loose alveolar tissue, dense fibrous tissue and fat pad (2).

Most of the studies on the cells present in the synovial membrane deal with the cells forming the synovial lining which are traditionally called synoviocytes, although according to Wilkinson et al (3) the use of the term ‘synoviocyte’ should be restricted to the fibroblast-like cells since this is probably the only cell type characteristic of synovial tissue.

The population of mononuclear cells of hematopoietic origin isolated from synovium-rich rat tissues is dominated by two types of cells. The first type represents cells with the morphology and immunophenotype of macrophages that resemble type A synoviocytes. The second type belongs to indeterminate cells which under the influence of granulocyte macrophage colony-stimulating factor (GM-CSF) differentiate into mature dendritic cells (7,8). Interestingly, in an EM study, Nozawa-Inoue et al (9) divided the macrophage-like A cells into two types i.e. electron-lucent with prominent vacuoles and lysosomes, and thus corresponding to the macrophage phenotype, and electron-dense cells, in which these organelles were less marked. The latter cells could correspond to the indeterminate cells described by Moghaddami et al (7,8).

Synovial macrophages, similarly to macrophages from other tissues, originate from monocytes migrating from the bone marrow, express class II MHC molecules and are involved in antigen presentation (7,9,10). Moreover, in the rat, they are positive for intracellular ED1 antigen present in cells of macrophage/monocyte lineage, and for ED2 antigen characteristic for tissue macrophages (11). Synovial type A cells remove particulate matter, antigens and other proteins from the synovial fluid by phagocytosis and endocytosis (8,12-16). The proficiency for phagocytosis seems to be similar in peritoneal and synovial macrophages. However, some of the latter are considerably more endocytotic than peritoneal macrophages. This may be interpreted as an adaptation to the specific conditions in the joint, since
efficient removal of debris and waste from the joint fluid is functionally important (8).

Synovial type B cells show high uridine diphospho-glucose dehydrogenase (UDPGD) activity (3), express hyaluronan synthases (HASs) (17,18) and produce large amounts of hyaluronan (HA) which is deposited in the synovial lining (19) and is secreted continuously to maintain effective concentration within the joint (20). Moreover, synovial fibroblasts produce lubricin, a glycoprotein believed to be responsible for the low friction between the moving articular surfaces and which is not detected in other fibroblast-rich tissues (21). They also have a high content of prolyl hydroxylase essential for collagen synthesis (3,22) and produce other elements of the extracellular matrix, such as fibronectin (23) and laminin. The latter concentrates at the basal aspects of the fibroblast-like synoviocytes and forms discontinuous, basement membrane-like structures (24). The B cells are also the main source of the synovial lining VCAM-1 and ICAM-1 (22,25). Adhesion between type B cells is provided by cadherin-11 (26). Synovial type B cells also express heat shock protein Hsp25 which enables their immunocytochemical detection and together with scanning electron microscopy demonstration of their unique surface topography with intertwined slender processes covering the surface of synovial lining, thick and long processes extending into the joint cavity and with type A cells residing in the slits of this meshwork (27-29, cf. 30).

The functional characteristics of synoviocytes has usually been studied in cultures of fibroblasts and macrophages obtained from normal or pathological synovial membrane. Interaction of activated macrophages and fibroblasts is critical for the resulting inflammation and tissue damage in RA (cf. 31,32) and OA (33). The responses of macrophages and fibroblasts from healthy tissues to cytokines and other factors has been extensively studied (34,35). Since both types of synoviocytes produce cytokines, matrix metalloproteinases (MMPs) and other factors, they may influence each other, and thus their combined response in normal synovium may differ from that of the single cell type. Furthermore, cytokines produced by endothelial cells (36) or cytokines and hormones produced by adipocytes present in the subsynovium may also modify the synoviocyte response (37-40). Thus, studies concerning chosen populations of cells forming the synovial membrane may not be representative of the whole organ.

We recently demonstrated that synovial membrane dissected from rat knee joints responded to stimulation by pro- and anti-inflammatory cytokines by increasing production of HA (41). In this work we have studied the influence of LPS and of several cytokines on the expression of cytokines, MMPs, and tissue inhibitors of metalloproteinases (TIMPs) at the mRNA and protein level, in order to obtain information on the reaction of the whole synovial membrane to various stimuli.

Materials and methods

Animals. Synovial membranes were obtained from inbred, 3 month-old Lewis male rats. Animals, kept in pathogen-free conditions, were obtained from the Animal Unit of the

Mossakowski Medical Research Centre, Polish Academy of Sciences, Warsaw. The study and the methods were approved by the Animal Ethics Committee of the Medical University of Warsaw.

Preparation of rat synovial membranes. After opening of the knee joint cavity, the synovial membrane was excised together with the patella, the patellar ligament and the joint capsule. Next, the synovial membrane with the infrapatellar fat pad was separated from the other tissues according to the method previously described (42).

Incubation of synovial membranes. Synovial membranes were put into flat-bottomed 24-well plates (Corning, NY, USA) in 1 ml of RPMI medium with 1% antibiotic and antimitotic solution (Sigma-Aldrich Chemie, Düsseldorf, Germany) for 12 h of incubation at 37°C in 5% CO₂ in air on a slow-motion shaker. The synovial membrane from one knee joint served as the control to the synovial membrane from the opposite knee. Experimental synovial membranes were exposed to 10 μg/ml of LPS, 10 ng/ml TGF-β1, 20 ng/ml TNF (Sigma) or 10 ng/ml IL-4 (PromoKine, PromoCell GmbH, Heidelberg, Germany). After incubation, the media were collected and stored at -20°C. Synovial membranes were used as a source of RNA. In a few additional experiments performed for evaluation of the mRNA expression under exposure to LPS and TNF the time of incubation was shortened to 1, 2 or 3 h.

ELISA assays for IL-6 and TNF determination. Cytokine level in supernatants was evaluated with rat immunoassay kits (Invitrogen Corporation, Camarillo, CA, USA), according to the manufacturer's protocols.

Total RNA isolation. Two synovial membranes (from two rats) were pooled. RNA was isolated with NucleoSpin® RNA II kit (Macherey-Nagel, Düren, Germany), according to manufacturer's protocol. The quantity of the isolated total RNA was measured with Picodrop (Picodrop Limited, Saffron Walden, UK). The quality of the isolated RNA was checked by electrophoresis in 1% agarose denaturing gel containing 6% formaldehyde and buffered with MOPS (Sigma). After electrophoresis, the gel was scanned with digital imaging system GDS 9000 using the Grab-It 2.0 software (UVP, Cambridge, UK).

Reverse transcription. Reverse transcription was performed using the High Capacity cDNA Reverse Transcription kit (Applied Biosystems, Cheshire, UK), according to the manufacturer's protocol in Eppendorf Mastercycler gradient. cDNA samples were stored at -20°C.

Real-time PCR. Real-time PCR was performed in the ABI PRISM 7500 (Applied Biosystems) using 96-well optical plates. Each sample was run in triplicate and was supplied with an endogenous control (Rat GAPDH endogenous control VIC®/MGB Probe). For gene expression analysis, the following TaqMan expression assays were used: TIMP1-Rn00587558_m1, TIMP2-Rn00573232_m1, TIMP3-Rn00441826_m1, TIMP4-Rn01459159_m1, MMP1A-
ZnCl₂ and 2.5% Triton X-100 for 30 min and incubated overnight at 37˚C in reaction buffer containing 50 mM Tris-HCl and 2.5% Triton X-100 to remove SDS. For gelatin zymography the gels were washed twice for 30 min at room temperature in 2.5% Triton X-100 and lyophilised, resuspended in 30 μl sample buffer with 2-mercaptoethanol and separated by SDS-polyacrylamide gel electrophoresis (SDS-PAGE; 10% acrylamide). Separated proteins were transferred onto PVDF membranes by semidry blotting at 25 V for 30 min using the Trans-Blot SD apparatus (Bio-Rad Laboratories, Hercules, CA, USA). Rabbit anti-TIMP1, TIMP2, TIMP4, MMP2 (Abcam, Cambridge, UK) and TIMP3 (Abbiotec, San Diego, CA) primary antibodies were used. The incubation lasted 1 h. Biotinylated F(ab')₂ fragments of swine anti-rabbit immunoglobulins (Dako A/S, Glostrup, Denmark) served as the secondary antibody. Antibody binding was demonstrated by an amplified alkaline phosphatase detection system (Bio-Rad). To identify particular TIMPs, recombinant rat TIMP1 (R&D Systems, Inc., Minneapolis, MN, USA), rat TIMP2 and human TIMP4 (Abcam) were used. The relative molecular weight of the proteins were calculated in relation to the pre-stained SDS-PAGE (low-range) standard (Bio-Rad). Intensity of the bands was measured by semi-quantitative densitometric analysis using GelWorks 2D-Intermediate software (UVP, Cambridge, UK).

**Western blot analysis.** Culture supernatants (150 μl) were lyophilised, resuspended in 30 μl sample buffer with 2-mercaptoethanol and separated by SDS-polyacrylamide gel electrophoresis (SDS-PAGE; 10% acrylamide). Separated proteins were transferred onto PVDF membranes by semidry blotting at 25 V for 30 min using the Trans-Blot SD apparatus (Bio-Rad Laboratories, Hercules, CA, USA). Rabbit anti-TIMP1, TIMP2, TIMP4, MMP2 (Abcam, Cambridge, UK) and TIMP3 (Abbiotec, San Diego, CA) primary antibodies were used. The incubation lasted 1 h. Biotinylated F(ab')₂ fragments of swine anti-rabbit immunoglobulins (Dako A/S, Glostrup, Denmark) served as the secondary antibody. Antibody binding was demonstrated by an amplified alkaline phosphatase detection system (Bio-Rad). To identify particular TIMPs, recombinant rat TIMP1 (R&D Systems, Inc., Minneapolis, MN, USA), rat TIMP2 and human TIMP4 (Abcam) were used. The relative molecular weight of the proteins were calculated in relation to the pre-stained SDS-PAGE (low-range) standard (Bio-Rad). Intensity of the bands was measured by semi-quantitative densitometric analysis using GelWorks 2D-Intermediate software (UVP, Cambridge, UK).

**Gelatin and casein substrate zymography.** The MMP2, MMP9, MMP1 and MMP3 activity was detected by gelatin and casein zymography. Briefly, 10 μl of culture medium in 10 μl sample buffer (non-reducing conditions) was loaded onto gelatin or casein (both from Sigma) containing gels for SDS-PAGE (10% polyacrylamide with 1 mg/ml gelatin or 0.1 μg/ml casein). After electrophoretic separation, the gels were washed twice for 30 min at room temperature in 2.5% Triton X-100 to remove SDS. For gelatin zymography the gels were washed subsequently in buffer containing 50 mM Tris-Cl and 2.5% Triton X-100 (pH 7.5) for 30 min, kept in 50 mM Tris-Cl buffer (pH 7.5) with 5 mM CaCl₂, 1 μM ZnCl₂ and 2.5% Triton X-100 for 30 min and incubated overnight in 50 mM Tris-Cl with 5 mM CaCl₂, 1 μM ZnCl₂ at 37°C. For casein zymography the gels were incubated overnight at 37°C in reaction buffer containing 50 mM Tris-HCl buffer (pH 7.5) with 200 mM NaCl and 0.02% 23 lauryl ether [Brij-35]. The gels were washed with 0.5% Coomassie Blue R-250 dissolved in 30% (v/v) methanol and 10% (v/v) acetic acid in H₂O for 30 min. The excess dye was washed away with the same solvent. Proteolytic activities appeared as clear bands of lysis against a dark background of stained gelatin or casein. Gelatinase bands were identified according to their molecular weight with purified recombinant MMP2 and MMP9 (R&D System) as standards. The gels were scanned with the digital imaging system GDS 90000 using Grab-It 2.0 software. The images were subjected to semi-quantitative densitometric analysis using GelWorks 2D-Intermediate software (UVP), in which the intensity of specific bands was measured.

**Activity of TIMPs was determined by reversed zymography,** using 25 μl culture medium per lane. SDS-PAGE electrophoresis was done in 10% polyacrylamide gels with 1 mg/ml gelatin and 0.1 μg/ml MMP2, using the procedure described above.

**Results**

To assess the quality of the dissection and the preparation of the synovial membranes, differences in cytokine levels in the culture medium of the synovium from the left and the right knee joint of the same rat were evaluated. There were no significant differences in cytokine levels (data not shown).

**ELISA assays.** Synovial membranes were stimulated with LPS, TNF, or the anti-inflammatory cytokines, IL-4 and TGF-81, and the concentration of IL-6 and TNF in the culture medium was evaluated. LPS strongly stimulated production of both tested cytokines. TNF caused a considerably lower increase of IL-6 production than LPS (influence of TNF on TNF production was not studied). TGF-81 caused a low but statistically significant increase of IL-6 production. IL-4 did not exert any influence on the levels of the tested cytokines (Figs. 1 and 2).

**Figure 1.** IL-6 concentration (mean ± SE) in control medium (C) and in medium containing LPS, TNF, TGF-81 or IL-4 after 12 h incubation of the synovial membranes. In each group n=6. Differences between groups were significant, according to the Wilcoxon matched-pairs test at *p<0.05.

**Figure 2.** TNF concentration (mean ± SE) in control medium (C) and in medium containing LPS, TGF-81 or IL-4 after 12 h incubation of the synovial membranes. In each group n=6. Differences between groups were significant, according to the Wilcoxon matched-pairs test at *p<0.05.
Real-time PCR. The mRNA levels of all tested cytokines, MMPs, and TIMPs in the synovial membranes incubated in the control medium were compared to that of GAPDH mRNA (ΔCt) which served as an endogenous control. The mRNA levels of MMP2, MMP3, TIMP1, TIMP2 and TIMP3 were higher while that of TGF-β1 was approximately equal to the mRNA levels of GAPDH. Furthermore, the mRNA levels of IL-1β, IL-6, MMP13 and MMP14 were slightly lower and those of TNF, MMP1a, MMP1b, MMP9 and TIMP4 were markedly lower than those of GAPDH (Fig. 3).
LPS caused a statistically significant increase in the mRNA levels of IL-6, IL-1ß, TGF-ß1, MMP1a, MMP1b, MMP3, MMP9, MMP13, MMP14, TIMP1 and TIMP3, and a significant decrease in the mRNA levels of MMP2, TIMP2 and TIMP4. LPS did not influence the expression of TNF mRNA after 12 h (Fig. 4a), but additional experiments demonstrated a 100-fold increase of the mRNA level at 1 h, a 10-fold after 2 h and a 3-fold after 3 h.

TNF decreased the mRNA levels of MMP2 and MMP14 and increased those of MMP1a, MMP1b, MMP3, MMP9 and MMP13. Expression of TIMP2, TIMP3 and TIMP4 decreased after TNF treatment while the mRNA levels of TIMP1 and of IL-1ß mRNA increased. The mRNA levels of TGF-ß1, IL-6 and TNF remained unchanged (Fig. 4b). In an additional experiment, IL-6 mRNA expression was increased by 8-fold after 3 h of incubation.

IL-4 stimulation decreased the mRNA levels of IL-1ß, TGF-ß1, MMP2, MMP9, MMP13, MMP14 and of all TIMPs. The mRNA levels of IL-6, TNF, MMP1a, MMP1b and MMP3 were unchanged (Fig. 4c).

TGF-ß1 decreased the mRNA levels of IL-1ß, TIMP1, TIMP2, TIMP4 and of all MMPs, but increased the expression of TGF-ß1 and TIMP3 mRNA. The mRNA levels of IL-6 and TNF remained unchanged (Fig. 4d).

Western blot analysis. MMP2 in its pro-active and active forms was identified in control and in all tested supernatants. LPS caused a decrease in the amount of this enzyme (Fig. 5).

Zymography. Gelatin zymography analysis was used to assess whether LPS, TNF, IL-4 and TGF-ß1 influence the activity of MMP2 and MMP9 produced by synovial membranes. Gelatinolytic enzyme activity associated with proteins of molecular weights corresponding to 74 kDa and 72 kDa was detected indicating the presence of proMMP2, and the active
Cytokine expression and secretion. The production of cytokines, proteolytic enzymes and their inhibitors by a complex tissue like the synovial membrane, presumably represents the net result of the activities of particular cell types and can be modified by factors released by these cells and reciprocally influencing their function. In our work stimulation of the synovial membrane by LPS increased secretion of TNF and IL-6 (Figs. 1 and 2). Previous studies have shown that macrophages express TNF mRNA (43) and that LPS augments their TNF expression at both the mRNA and protein levels (44,45). Secretion of TNF by adipose tissue has also been described (37,46,47), but it presumably had originated from macrophages residing within the tissue (48). Thus, most, if not all of the TNF released from the synovial membrane is likely produced by macrophages, since T cells in normal rat synovial membrane occur in insignificant number (49). The level of TNF mRNA measured after 12 h of LPS stimulation, which is our standard procedure, was similar to control. Since, however, in other studies the expression of TNF mRNA was raised much earlier and it returned to the control levels after several hours (50), we performed an additional experiment with LPS stimulation lasting for 1-3 h and we observed a considerable increase in TNF mRNA expression.

Secretion of IL-6 after LPS stimulation was previously observed in synovial fibroblasts (34,51,52), in endothelial cells (34,51) and peritoneal macrophages (43,44,53). Adipocytes were also shown to express and release IL-6 (40,47,48). Thus, all four main cell types could contribute to the markedly increased expression of IL-6 at the mRNA and protein levels after LPS stimulation (Figs. 1 and 4a).

IL-18 transcript was present in the non-stimulated synovial membrane. Its level was considerably increased after LPS stimulation (Fig. 4a). We have previously observed that LPS stimulation increased the content of IL-1α in the synovial membrane lysates (42). Unstimulated rat monocytes or peritoneal macrophages do not express mRNA IL-18 (43), but IL-1α and IL-18 were shown to be produced after LPS stimulation (43,54,55). Moreover, LPS has been reported to stimulate IL-1 production by vascular cells (56), fibroblasts (34) and adipocytes (57). Thus, the IL-18 transcript could originate from all the main cell types present in the synovial membrane. LPS also increased the mRNA level of TGF-β1 (Fig. 4a). LPS has been shown to stimulate the release of TGF-β1 in endothelial cells (58) and the TGF-β1 expression in PMA-stimulated macrophages (59).

TNF also increased IL-6 secretion but with much less potency than LPS (Fig. 2). According to previous reports, TNF has no effect on IL-6 synthesis in monocytes and macrophages (60), but exerts a moderate stimulatory effect on the secretion of IL-6 by fibroblasts (61) and endothelial cells (51). Therefore, the increased production of IL-6 by the synovial membrane after TNF stimulation can be ascribed to the fibroblast-like synoviocytes and the endothelial cells. In spite of this increased secretory activity, the level of IL-6 mRNA was similar to the controls, but a rise was observed if stimulation was shortened to 3 h. The mRNA levels of IL-18 were moderately increased after TNF stimulation (Fig. 4b). Thus, our results are in accord with previous observations that after TNF stimulation, IL-1α and IL-18 are produced by monocytes, macrophages and vascular cells (43,54,56).

TGF-β1 exerted a weak stimulatory effect on IL-6 secretion by the synovial membrane, which, however, was unnoticeable at the mRNA level (Figs. 1 and 4d). TGF-β1 mRNA was previously found to be expressed by the fibroblast-like synoviocytes and the endothelial cells (34). TGF-β1 was shown to induce IL-6 mRNA and protein expression in human fibroblasts (62,63). There are, however, inconsistencies between reports on the influence of TGF-β1 on monocytes. In one study TGF-β1 was found to induce (64) and in another (65) to inhibit IL-6 production. Thus, the main effect of the stimulatory action of TGF-β1 on IL-6 secretion was probably due to its action on the fibroblast-like synoviocytes, while contribution of the synovial macrophages remains uncertain. TGF-β1 had no effect on TNF mRNA level (Fig. 4d). TGF-β1 was previously found to suppress TNF production by peritoneal macrophages (66), presumably inhibiting translation of TNF mRNA (67,68). Furthermore, TGF-β1 decreased the expression of IL-1β mRNA (Fig. 4d) in agreement with a previous report (69). In human PBMCs, TGF-β1 was shown to cause an increase in the mRNA levels of IL-1α and IL-1β (70). In our study, TGF-β1 markedly stimulated the expression of TGF-β1 mRNA (Fig. 4d) which is in good agreement with previous observations, that TGF-β1 can induce its own gene expression in fibroblasts (71), and other cell types (72,73).

IL-4 had no effect on TNF and IL-6 secretion or mRNA levels in the synovial membrane (Figs. 1, 2 and 4c). In previous studies IL-4 did not influence IL-6 production in rheumatoid synovial fibroblasts or various fibroblast cell lines (74). However, IL-4 stimulated the production of TNF by mononuclear phagocytes (75). Human umbilical vein endothelial cells stimulated by IL-4, in one study, produced
only low amounts of IL-6 and expressed low levels of the IL-6 transcript (76), while in another it increased IL-6 production (77). Thus, the lack of IL-4 stimulatory action on TNF and IL-6 production observed by us is in accord with studies on fibroblasts, but differs from that on mononuclear phagocytes (75). The influence of IL-4 on the secretion of TGF-ß1 was not studied, but a small, although significant decrease of TGF-ß1 mRNA levels was noted (Fig. 4c). In our work IL-4 slightly suppressed the mRNA levels of IL-1ß (Fig. 4c) which is in agreement with Te Velde et al (78) who described a decreased secretion of IL-1ß in human monocytes after IL-4 treatment, but in opposition to Lacroz et al (75) who observed increased production of IL-1 by mononuclear phagocytes.

MMPs. MMP2 (72 kDa gelatinase; gelatinase A, type IV-A collagenase), digests gelatin, fibronectin, laminin, collagen type V and to a lesser extent collagen type IV, cartilage proteoglycans and elastin (79). MMP2 is secreted as progelatinase and is activated by MMP14 (80). MMP2 is produced by synovial fibroblasts (79,81,82), by macrophages, particularly after stimulation with LPS (83), and by endothelial cells (84,85), particularly after stimulation with TGF-ß1 (86) or LPS (87). MMP2 is also produced by adipocytes (88). In our experiment, all agents tested, that is LPS, TNF, IL-4 and TGF-ß1, decreased the expression of MMP2 mRNA (Fig. 4a-d). Moreover, after LPS treatment the medium used for the synovial membrane contained less MMP2 protein as detected by Western blot analysis and less MMP2 activity as demonstrated in the zymography assay (Figs. 5 and 6).

MMP9 (92 kDa gelatinase; gelatinase B; type IV-B collagenase) digests gelatin, collagen types IV, V, XI, XVII, elastin, fibronectin, link proteins, aggrecan and several cyto-
kines (cf. 89). Typically, MMP9 is secreted by cells as a proenzyme in a noncovalent complex with TIMP1. Interaction of MMP9 with TIMP1 prevents its dimerization, formation of the complex with MMP1 and activation by MMP3 (89,90). Secretion of MMP9, the most abundant gelatinase in macrophages, has been reported to be stimulated 3- to 4-fold after exposure of human alveolar macrophages to LPS (83). In monocytes and in monocyte-derived macrophages TNF also increased the production of MMP9 (92,93). It is also noteworthy, that macrophages cultured on polymerized collagen, conditions which are roughly similar to those used for culturing synovial membrane, produced less MMP9 than when the cells were cultured on plastic, but strongly responded to TNF stimulation by increasing MMP9 expression (94). TNF has also been shown to stimulate the expression of MMP9 in endothelial cells (95).

Furthermore, MMP9 is expressed by adipocytes and is has been shown to be up-regulated in a macrophage-conditioned medium, presumably due to its TNF content (96). Production of the 92 kDa gelatinase has not been observed in synovial fibroblasts (82). In our work MMP9 expression was increased about 60-fold after LPS stimulation. TNF also strongly stimulated MMP9 expression but less potently than LPS. Its expression was decreased by TGF-ß1 and IL-4, similarly to another study on mononuclear phagocytes (75) (Fig. 4a-d). Media from experimental and control cultures of synovial membranes, however, did not contain MMP9 gelatinolytic activity in the zymography assay.

TIMPs. TIMPs are produced by a variety of cells and are present in most tissues, but their expression is individually regulated by cytokines and depends on the cell type. TIMPs 1, 2 and 4 are secreted into the medium, while TIMP3 is predominantly matrix-associated (98,110). LPS has been reported to stimulate TIMP1 expression and production by alveolar macrophages (91,104) or by human monocytes (93). TIMP1 mRNA expression was observed to be markedly increased in dermal fibroblasts after TNF treatment (111). In endothelial cells, TNF had no effect on TIMP1 and TIMP2 mRNA levels and did not induce TIMP3 or TIMP4 expression (105). IL-4 decreased TIMP1 production in monocytes (93). Production of TIMP2 in dermal fibroblasts at the mRNA and protein levels was shown to be stimulated
by IL-4, but not by TGF-81 (112). TGF-81 has been shown to induce expression of both TIMP1 and TIMP3 in human synovial lining cells (113). Moreover, increased expression of TIMP3, under influence of TGF-81, has been observed in fibroblasts (114).

In our work LPS increased the mRNA expression of TIMP1 and TIMP3 in the synovial membrane, while the expression of TIMP2 and TIMP4 decreased (Fig. 4a). Stimulation by TNF was much weaker but qualitatively similar to that produced by LPS, with the exception of the TIMP3 mRNA expression which decreased (Fig. 4b). TGF-81 considerably raised the expression of TIMP3 and decreased the expression of TIMP1, TIMP2 and TIMP4 (Fig. 4d). IL-4 slightly, but significantly, decreased the expression of all TIMPs (Fig. 4c). TIMP1, TIMP2 and TIMP4 were detected by Western blot analysis in similar amounts in supernatants from all control and experimental groups (Fig. 7). Only TIMP1 was detected by reversed zymography and no difference was observed between control and stimulated membranes.

In this study we attempted to accomplish the general characterization of the synovial membrane response to a few biologically active factors known to influence various cell types at least nominally analogous to those present in the synovial membrane. A precise comparison of cytokine expression at the mRNA and protein level in the rat synovial membrane with data for particular cell types presented by other authors is difficult, due to the difference in the origin of the cells (human vs. murine), their various modifications in culture, the differences in the period of observation and the dosage of the stimulating agents. Nevertheless, even with these limitations, it is seen that the responses of the synovial membrane to LPS, TNF, TGF-81 and IL-4 are essentially similar to those reported in the papers mentioned above and can be explained by the activity of all of the particular cell types present in the membrane, although their individual contribution is difficult to estimate. The expression, secretion and activity of MMPs and TIMPs in the synovial membrane represents a summary of the activities of its various cell populations but cannot be predicted on the basis of analysis of particular cell types forming the tissue. This is particularly evident in the case of MMP2, whose expression and activity according to data obtained by others working with pure cell populations should have been increased after LPS stimulation, but resulted in a decrease in our system of the whole synovial membrane.

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