Distinct roles for the NF-κB1 (p50) and c-Rel transcription factors in inflammatory arthritis

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Rheumatoid arthritis (RA) is a complex disease, with contributions from systemic autoimmunity and local inflammation. Persistent synovial joint inflammation and invasive synovial pannus tissue lead to joint destruction. RA is characterized by the production of inflammatory mediators, many of which are regulated by the Rel/NF-κB transcription factors. Although an attractive target for therapeutic intervention in inflammatory diseases, Rel/NF-κB is involved in normal physiology, thus global inhibition could be harmful. An alternate approach is to identify and target the Rel/NF-κB subunits critical for components of disease. To assess this, mice with null mutations in c-rel or nfkbia were used to examine directly the roles of c-Rel and p50 in models of acute and chronic inflammatory arthritis. We found c-Rel–deficient mice were resistant to collagen-induced arthritis but had a normal response in an acute, destructive arthritis model (methylated BSA/IL-1 induced arthritis) suggesting c-Rel is required for systemic but not local joint disease. In contrast, p50-deficient mice were refractory to induction of both the chronic and acute arthritis models, showing this subunit is essential for local joint inflammation and destruction. Our data suggest Rel/NF-κB subunits play distinct roles in the pathogenesis of inflammatory arthritis and may provide a rationale for more specific therapeutic blockade of Rel/NF-κB in RA.


Introduction

Rheumatoid arthritis (RA) is a complex autoimmune disease that results in chronic inflammation of synovial joints. A common paradigm for the pathogenesis of RA is that major histocompatibility-restricted T-cell activation in response to an unknown joint-derived antigen results in production of autoantibodies and systemic disease induction. Rheumatoid synovitis then occurs, characterized by the infiltration of inflammatory cells into the synovial compartment and the production of inflammatory mediators, many of which are thought to be regulated by the Rel/nuclear factor-κB (NF-κB) transcription factor family (1, 2). Collagen-induced arthritis (CIA) is a widely used murine model of RA (3). Recent studies on CIA in Rag-1–deficient mice, which lack mature T and B cells and yet still develop CIA, support the concept of two components of disease development (4). One is the systemic immune component outlined above, whereas a second involves local synovial cell activation that may be less dependent on a systemic immune response. This concept is further supported by the observation that synovial cells from patients with RA are intrinsically invasive when engrafted into SCID mice (5). The formation of synovial pannus tissue is a characteristic feature of RA and leads to destruction of underlying articular cartilage and bone.

The Rel/NF-κB family of transcription factors are homo- and heterodimeric proteins comprising subunits encoded by a multigene family related to the c-rel proto-oncogene. Five mammalian Rel/NF-κB proteins have been described to date: NF-κB1 (p50, p105), NF-κB2 (p52, p100), RelA (p65), RelB, and c-Rel subunits, encoded by the nfkbia, nfkbi, rela, relb, and c-rel genes, respectively (see refs. 6 and 7 for reviews). In unstimulated cells, the majority of Rel/NF-κB dimers are retained in the cytoplasm as an inactive complex bound to inhibitor proteins (IκBs). In response to a wide variety of stimuli, IκB proteins are phosphorylated by the IKK complex (8), targeting the IκBs for intracellular degradation (9, 10). Rel/NF-κB dimers then translocate to the nucleus and bind to decameric DNA sequences (κB elements) required for the transcription of many genes, including cytokines, chemokines, and adhesion molecules (1). Functional effects of the different Rel/NF-κB dimers may be dependent on binding preferences for particular κB sequences, as well as varying expression profiles of the individual subunits in tissues (11, 12).

Much of our understanding of the function of the Rel/NF-κB family members developed from in vitro experiments, but more recent studies with gene-knockout mice have illuminated the in vivo functions of individual subunits. Lack of RelA causes embryonic lethality due to apoptosis of fetal hepatocytes (13), whereas...
c-Rel (14), p50 (15), and RelB (16, 17) are required for normal hemopoietic and immune cell function. Although c-Rel–deficient (c-rel–/) and p50-deficient (nfkb1–/) mice develop normally, their mature B and T cells exhibit selective activation defects in response to certain mitogenic stimuli (14, 15).

Rel/NF-κB has been considered a potential target for therapeutic intervention in inflammatory disease. Indeed, it has now been realized that at least part of the anti-inflammatory effects of widely used drugs, such as dexamethasone and aspirin, is through inhibition of Rel/NF-κB (18–20). However, because Rel/NF-κB plays such a critical role in immune regulation, nonspecific inhibition might compromise normal host defenses. An alternative approach is to identify and target the Rel/NF-κB subunits critical for disease development.

In this study, we employed mice homozygous for null mutations in genes encoding the Rel/NF-κB subunits, c-Rel and p50, to directly examine the role of these transcription factors in inflammatory arthritis. Two models were examined: CIA, a model of chronic systemic autoimmune arthritis, and methylated BSA/IL-1–induced arthritis, an acute monoarticular arthritis. Each of these models has features of human RA. We report that Rel/NF-κB is an essential mediator of both chronic autoimmune and acute inflammatory arthritis. Both c-rel–/– and nfkbid−/− mice had impaired cellular and humoral immunity to type II collagen (CII) and largely failed to develop CIA. However, c-Rel–deficient mice had a normal response in the acute arthritis model, suggesting c-Rel is not required for the destructive phase of joint disease. In contrast, p50-deficient mice were refractory to the induction of acute arthritis, showing that this Rel/NF-κB subunit is essential for local joint inflammation and destruction. Our data suggest that Rel/NF-κB subunits play distinct roles in the pathogenesis of inflammatory arthritis and may provide the basis for more specific therapeutic intervention in RA.

Methods

Mice. Mice deficient in the c-rel gene (c-rel–/–), were derived as described previously (14) and backcrossed onto the C57BL/6 strain for 10 generations before use. The p50-deficient mice (nfkbid−/−) (15), were further backcrossed onto C57BL/6 mice for eight generations before use. C57BL/6 mice, obtained from The Walter and Eliza Hall Institute animal services (Kew, Victoria, Australia), were used as wild-type (WT) controls in all experiments. All mice were aged 8 to 13 weeks.

Collagen-induced arthritis. Chick CII (Sigma Chemical Co., St. Louis, Missouri, USA), dissolved in 10 mM acetic acid at a concentration of 2 mg/mL, was emulsified in an equal volume of CFA containing 5 mg/mL heat-killed Mycobacterium tuberculosis (strain H37Ra; Difco Laboratories, Detroit, Michigan, USA), as described previously (21, 22). Arthritis was induced by injecting mice i.d. in several doses at the base of the tail with 100 μL emulsion at day 0 and 21. Individual experiments contained at least nine mice per group, and all experiments were performed two or more times.

Animals were assessed for swelling of limbs, and a clinical score was allocated to each mouse two to three times per week, for up to 60 days. The scoring system was as described previously (23), where 0 indicates normal, 1 indicates slight swelling, 2 indicates extensive swelling, and 3 indicates joint distortion and/or rigidity. The maximum score per mouse was 12. Mice were considered to have arthritis when two consecutive positive evaluations were obtained. Where swelling was restricted to digits, the maximum score for the affected limb was 1, regardless of severity. At sacrifice, paws were removed, fixed, decalcified, and processed for paraffin embedding. Hematoxylin and eosin–stained sections (5 μm) were evaluated for arthritis as described previously (21), where mild indicated minimal synovitis with cartilage and bone erosions confined to discrete foci; moderate indicated synovitis and erosions with intact joint architecture; and severe indicated extensive erosions and disrupted joint architecture. Clinical and histological evaluations were assessed by two independent investigators blinded to the experimental groups.

Acute monoarticular arthritis. Acute monoarticular arthritis was induced as described (24, 25). On day 0, mice were anesthetized and the skin over the knees incised to expose the stiffe joints. Two hundred micrograms of methylated BSA (mBSA; Sigma Chemical Co.) was injected into the left knee joint, and the contralateral joint received 10 μL saline. The incisions were then closed with wound clips. Human recombinant IL-1β (250 ng) in 0.5% (vol/vol) normal mouse serum (in saline) was injected subcutaneously into the left footpad on days 0, 1, and 2. Mice were sacrificed at day 7 and their rear limbs removed and processed, as above, for histological assessment of hematoxylin and eosin–stained frontal sections of the stiffe joints. Coded joint sections were assessed histologically for five features of inflammatory arthritis, each on a scale of 0 (normal) to 5 (severe), as described (24). These were exudate (presence of inflammatory cells or fibrinlike material in the joint space), synovitis (thickening of the synovium, infrapatella fatpad, and peristemeum, but excluding pannus), pannus (synovial hyperplasia immediately adjacent to cartilage or bone), cartilage degradation, and bone degradation. Sections were assessed by two independent observers without knowledge of the treatment groups. There were six to ten mice per experimental group.

ELISA for detection of Abs to CII. ELISAs were performed to detect Abs to CII as previously described (21). Horse-radish peroxidase–conjugated goat anti-mouse IgG (Sigma Chemical Co.) or IgM (Southern Biotechnol-oogy Associates, Birmingham, Alabama, USA) antisera were used as detection Abs. Standard curves for anti-CII IgG and IgM were constructed from sera of CII-hyperimmunized mice using arbitrary units.
**T-cell proliferation assay.** A single-cell suspension of inguinal lymph node cells (2 \( \times 10^6 \) cells/mL in RPMI containing 50 \( \mu \)M 2-mercaptoethanol and 5% FBS), prepared from mice (n > 3) at least 10 days after CIA immunization, was incubated at 37°C (5% CO\(_2\)) for 72 hours with 0–50 \( \mu \)g/mL of denatured CII (boiled for 10 minutes). Cultures were performed in 96-well round-bottomed plates (Becton Dickinson Labware, Franklin Lakes, New Jersey, USA) with 0.2 mL per well (4–6 wells per group). For the final 6–12 hours, 5 \( \mu \)Ci/mL \([^{3}H]\)thymidine (Amersham International, Amersham, United Kingdom) was added and cells harvested with an Inotech cell harvester (Rockville, Maryland, USA). A microplate scintillation counter (Canberra Packard, Victoria, Australia) was used to determine radioactive incorporation as a measure of T-cell proliferation.

**Electrophoretic mobility shift analysis.** Synovial tissue was dissected from acutely arthritic joints (n = 20), 7 days after injection of mBSA, and a single-cell suspension was prepared by digestion for 1 hour at 37°C in RPMI containing 2.4 mg/mL dispase II (Boehringer-Mannheim GmbH, Mannheim, Germany), 1 mg/mL type II Clostridial collagenase (Sigma Chemical Co.), and 0.1 mg/mL bovine pancreatic DNase I (Boehringer-Mannheim GmbH). After overnight incubation in RPMI plus 10% FBS, nuclear extracts were prepared from the washed, adherent synovial cells for electrophoretic mobility shift analysis (EMSA) as described (26). The Ig\( \kappa \) probe (5'-GTACGAGGGGCTTTCCGA-3') was used in these experiments. For supershift analyses, extracts were incubated for 1 hour on ice with 0.2 mg/mL rabbit Ab to mouse p65, c-Rel, and RelB (Santa Cruz Biotechnology, Santa Cruz, California, USA), and p50 (gift from N. Rice, National Cancer Institute, Frederick, Maryland, USA) before addition of radioactively labeled probe. Bands were visualized using a Phospho-Imager with Image Quant v5.0 software (Molecular Dynamics, Sunnyvale, California, USA).

**Statistics.** For CIA clinical scores and acute monoarticular arthritis histological scores the Mann-Whitney \( t \)-test was used to determine the level of significance between means of groups. For data per- two-sample rank test was used to determine the level of significance between means of groups. For data per-

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**Table 1 Reduced incidence and clinical severity of CIA in c-rel\(^{-/-}\) and nfkbf1\(^{-/-}\) mice**

<table>
<thead>
<tr>
<th>Group</th>
<th>Incidence (%)( ^{a} )</th>
<th>Range( ^{b} )</th>
<th>Mean ± SEM( ^{d} )</th>
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<tr>
<td>WT</td>
<td>23/37 (62)</td>
<td>1–12</td>
<td>3.1 ± 0.6</td>
</tr>
<tr>
<td>c-rel(^{-/-})</td>
<td>4/35 (11)( ^{f} )</td>
<td>1–4</td>
<td>0.3 ± 0.1</td>
</tr>
<tr>
<td>WT</td>
<td>15/25 (60)</td>
<td>1–10</td>
<td>1.8 ± 0.5</td>
</tr>
<tr>
<td>nfkbf1(^{-/-})</td>
<td>0/24 (0)( ^{k} )</td>
<td>-</td>
<td>0</td>
</tr>
</tbody>
</table>

\(^{a}\) Pooled data from three (for c-rel\(^{-/-}\)) or two (for nfkbf1\(^{-/-}\)) separate experiments. \(^{b}\)jackknife CIA incidence (arthritic/total mice) at day 60 (percentage in parentheses). \(^{c}\) Range of clinical scores of responsive mice to day 60. \(^{d}\) Corresponds to maximum clinical scores attained for all mice to day 60. \(^{f}\) Significant compared with WT. \(^{k}\) P < 0.0001 compared to WT.

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**Results**

**c-rel\(^{-/-}\) and nfkbf1\(^{-/-}\) mice are resistant to CIA.** To examine the role of Rel/NF-kB in a model of autoimmune arthritis, WT mice and mice homozygous for null mutations either in the c-rel or nfkbf1 genes were compared for incidence and severity of CIA induced by intradermal immunization with chick CII in CFA, followed by a boost injection 21 days later. We showed previously that this procedure successfully elicits arthritis in C57BL/6 mice (21) and is dependent on T and B cells (22). Paws were serially assessed and a clinical score given for each mouse according to the procedure outlined in Methods. Figure 1 shows the kinetics of the response in c-rel\(^{-/-}\) and WT mice both in terms of incidence (Figure 1a) and clinical severity (Figure 1c).

Only four of 35 (11%) c-Rel–deficient mice showed clinical signs of arthritis compared with 23 of 37 (62%) WT mice (\( P < 0.001 \)) by day 60 (Table 1). The clinical scores of the c-rel\(^{-/-}\) mice ranged from 1 to 4 compared with 1 to 12 in WT mice and were significantly lower (\( P < 0.001; 0.3 ± 0.1 \) vs. 3.1 ± 0.6, mean ± SEM for c-rel\(^{-/-}\) and WT mice, respectively). Therefore, c-Rel–deficient mice had a lower incidence and reduced severity of CIA compared with WT controls. Compared with 15 of 25 WT mice, p50-deficient mice were completely resistant to CIA, with no clinical evidence of disease in a total of 24 mice tested (Table 1).

**Histological analysis of CIA.** To confirm the clinical analyses, hematoxylin and eosin–stained sections of joints from c-rel\(^{-/-}\), nfkbf1\(^{-/-}\), and WT mice were assessed by histology. There was a strong concordance between the clinical and histological assessments. Joints of WT mice frequently (62% of joints examined) showed severe inflammation and joint tissue destruction with invasive pannus tissue associated with bone and cartilage degradation (Figure 1b and Table 2). In contrast, the majority (83%; see Table 2) of joints from the c-rel\(^{-/-}\) mice appeared normal (Figure 1d). However, in the few c-rel\(^{-/-}\) arthritic joints, the histological features were similar to that of the WT arthritic mice (not shown), with 6% of the joints examined showing severe pathology (Table 2). Therefore, c-Rel plays an important but not indispensable role in the full expression of CIA. In accordance with clinical assessments, none of the nfkbf1\(^{-/-}\) mouse joints showed any signs of arthritis.

**Altered humoral and reduced cellular immune responses to CII in c-rel\(^{-/-}\) and nfkbf1\(^{-/-}\) mice.** Studies were undertaken to determine why c-rel\(^{-/-}\) and nfkbf1\(^{-/-}\) mice are resistant to CIA. Since CIA is dependent on both humoral and cellular immune responses to CII (27), these were examined. The c-rel\(^{-/-}\) mice had a dramatically lower IgG response compared with WT mice at all time points (Figure 2a, upper panel). However, the c-rel\(^{-/-}\) IgM response, although lower at day 12, was comparable to WT mice at day 28.
Clinical and histological assessment of CIA in c-rel−/− versus WT mice. The incidence of arthritis (shown as cumulative percentage) (a) and mean clinical scores (± SEM) (c) of c-rel−/− (open circles, n = 35) and WT (filled circles, n = 37) mice are shown with time after primary immunization with CII. Data are pooled from three experiments. For statistical analysis see Table 1. At 60 days after primary immunization with CII, mice were sacrificed, their hind limbs removed, and the paws processed for histology (see Methods). Frontal sections of the interphalangeal joints of WT (b) and c-rel−/− (d) mice are shown. WT mouse joints frequently showed severe pathology with pannus invading into the subchondral bone (arrow). The majority of c-rel−/− mouse joints examined appeared normal, with intact articular cartilage (C) and no inflammatory cells in the joint space (J) or synovium (arrowhead). B, bone; M, bone marrow. Hematoxylin and eosin stained. x200.

Nuclear translocation of p50 but not c-Rel in acute arthritis synovial cells. To further investigate the role of Rel/NFκB subunits in acute inflammatory arthritis, EMSA and supershift analyses were performed on nuclear extracts of primary adherent synovial cells obtained from mBSA-injected joints of WT and c-rel−/− mice. Figure 5 shows two major bands were detected for each genotype; a slower migrating band supershifted by Abs to p50 and p65 (top arrow) and a faster migrating band supershifted by Abs to p50 only (bottom arrow). Abs to c-Rel and RelB had no effect on the electrophoretic mobilities of either band. These findings are consistent with the activation in WT and c-rel−/− arthritic joint synovial cells of p50/p65 heterodimers and p50 homodimers, but not of c-Rel- or RelB-containing complexes.

Humoral response to CII in c-rel−/− (a) and nfkβ−/− (b) mice. Serum levels of CII-specific Abs (IgG and IgM) were determined by ELISA in c-rel−/−, nfkβ−/−, and WT mice (n = 9–10) at different times after primary immunization with CII. Results show the mean + SEM values for IgG and IgM in arbitrary units per milliliter for individual experiments. *P < 0.001, *P < 0.01, *P < 0.05 compared with WT control mice.
Histopathological assessment of joints from c-Rel–deficient and WT mice

<table>
<thead>
<tr>
<th>Histological grade</th>
<th>Number of joints (%)</th>
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<tr>
<td></td>
<td>WT</td>
</tr>
<tr>
<td>Total joints</td>
<td>155</td>
</tr>
<tr>
<td>Normal</td>
<td>29 (19)</td>
</tr>
<tr>
<td>Mild</td>
<td>5 (3)</td>
</tr>
<tr>
<td>Moderate</td>
<td>25 (16)</td>
</tr>
<tr>
<td>Severe</td>
<td>96 (62)</td>
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</table>

َاJoints of the front and rear paws of clinically positive mice (n = 4 mice per group) were examined histologically. *Data show the number (percentage in parentheses) of joints distributed within each histological severity grade for the two groups of mice. ‡P < 0.001, compared with WT mice.

Discussion

The Rel/NF-κB family of transcription factors regulate a wide variety of inflammatory mediators, most of which are found in synovial joints of patients with RA (1, 2). Several immunohistochemical studies have reported Rel/NF-κB activation in synovial tissue from RA patients (28, 29) and in animal models of inflammatory arthritis (30, 31), thereby implicating Rel/NF-κB in disease pathogenesis. In this study we used mice homozygous for null mutations in genes encoding the Rel/NF-κB subunits, c-Rel or p50, to obtain direct evidence for Rel/NF-κB involvement in acute, monarticular, and chronic polyarticular inflammatory arthritis models. Our results show that Rel/NF-κB is essential for the development of inflammatory arthritis and, to our knowledge, provide the first evidence that selective blockade of Rel/NF-κB subunits may prevent it. We also show that the absence of different Rel/NF-κB subunits can have differential effects suggesting distinct roles, rather than redundancy, for the various Rel/NF-κB subunits in different stages of disease.

An interesting result from this study was the almost complete absence of CIA in c-Rel–deficient mice, but a typical response of these mice in the acute arthritis model. CIA is a widely used model of RA thought to be dependent on the establishment of both cellular and humoral immunity to CII (27), although recent evidence suggests that innate immune responses to CII (4) are dependent on p50 but not c-Rel. Deficiencies in other immune or inflammatory mechanisms may also limit CIA in c-rel−/− and nfkb1−/− mice. Both knockout mice had reduced T-cell proliferative responses to CII (Figure 3), consistent with previous findings that showed a diminished response to mitogens and/or altered production of cytokines by T cells (14, 15, 35–37). However, antigen-present-
ing cell defects could also be involved. Whereas \textit{c-rel}–/– and \textit{nfkb1}–/– mice exhibited both impaired humoral and cellular immune responses to CII that together could account for the dramatic reduction of CIA, other contributing influences could be the cytokine profile and activation state of macrophage populations (15, 38). Macrophages are important both as antigen-presenting cells (39) and as a source of inflammatory mediators in CIA (40, 41).

In contrast to the response of \textit{c-rel}–/– mice in CIA, these mice were indistinguishable from WT in the acute inflammatory arthritis model, which is dependent on IL-1 and CD4+ T cells, but not B cells. The knee sections of each showed infiltration of neutrophils and mononuclear cells, pannus formation, soft tissue inflammation, and cartilage and bone erosion. This is consistent with the histological features observed in the few CIA-affected joints of the \textit{c-rel}–/– mice. Taken together, these results suggest that c-Rel is required for establishment of the immune response in CIA but not for the destructive (effector) phase of inflammatory arthritis. In contrast to \textit{c-rel}–/– mice, \textit{nfkb1}–/– mice had a markedly diminished response in the acute arthritis model. There were few inflammatory cells in the joint space and connective tissues and a striking reduction in synovial hyperplasia. Since Rel/NF-κB regulates the transcription of adhesion molecules, prostaglandin-producing enzymes, and chemokines (1), this may explain the reduced cellular trafficking into the mBSA-injected joints of p50-deficient mice.

The effect of p50 deletion on T-cell function is unclear. There are conflicting reports on \textit{nfkb1}–/– T-cell proliferation and cytokine production (15, 36, 37). The p50-deficient mice were protected in a T cell–dependent model of allergic airway inflammation (37); antigen-specific T-cell responses were maintained in these mice, but IL-5 production was reduced. It is possible that different T-cell cytokine profiles in the \textit{c-rel}–/– and \textit{nfkb1}–/– mice may account for responses in the acute arthritis model. Both \textit{c-rel}–/– and \textit{nfkb1}–/– mice had antigen-specific T-cell proliferative responses, but these were reduced — to a similar degree — compared with WT mice (Figure 3). In spite of this, T cells from \textit{c-rel}–/– mice were able to elicit a normal response in the T cell–dependent acute arthritis model, but \textit{nfkb1}–/– mice showed almost no disease.

A recent study showed Rel/NF-κB inhibition enhanced synovial apoptosis in a rat model of arthritis (42), consistent with the potential involvement of Rel/NF-κB transcription factors in synovial hyperplasia. Once formed, pannus is thought to mediate cartilage and bone destruction in RA through matrix metalloproteinase (MMP) production (43). The transcription of MMPs appears to be regulated in part by Rel/NF-κB (44). Inhibition of Rel/NF-κB in human rheumatoid synovial cell cultures reduced proinflammatory cytokine

![Figure 4](image)

**Figure 4**
Acute inflammatory arthritis in Rel/NF-κB–deficient and WT mice. Monarticular arthritis was induced as described in Methods. Frontal sections of control (a) and mBSA-injected patello-femoral joints of WT (b), \textit{c-rel}–/– (c), and \textit{nfkb1}–/– (d) mice are shown (hematoxylin and eosin–stained). ×200. E, inflammatory exudate; F, femur; I, inflammatory cell infiltrate; P, pannus; Pt, patella. Arrow indicates synovial lining. Coded joint sections were assessed histologically for five features of inflammatory arthritis, each on a scale of 0 (normal) to 5 (severe). Results (e and f) compare the mean ± SEM values for each of the histological features of \textit{c-rel}–/– (n = 6) (e) and \textit{nfkb1}–/– (n = 10) (f) mice (open bars) to those of WT mice (filled bars). Control (saline) joints scored 0 for all categories (not shown). *P < 0.005.
production, together with MMP-1 and MMP-3 (45). In IL-1β-stimulated rabbit synovial fibroblasts, p50 bound to a regulatory element in the distal promoter of MMP-1 (46). These findings, together with the results presented here, suggest p50 is involved in pannus formation and perhaps in regulating joint tissue destruction through effects on MMP gene transcription.

The importance of p50 in synovial cell gene transcription was confirmed by EMSAs, which showed p50/p65 heterodimers and p50 homodimers in nuclear extracts of cells isolated from the acutely inflamed joints of WT and c-rel−/− mice (Figure 5). In contrast, c-Rel was not involved in the activated synovial cell response. Because primary cells were used, the Rel/NF-κB complexes could have been derived from macrophage-like (type A) or fibroblast-like (type B) synovial cells, or both. Using immunohistochemistry, p50 and p65 have been found in type A cells of rheumatoid synovium (28, 29) and in passaged rheumatoid synovial fibroblasts by EMSA (47, 48). We have now also identified p50 homodimers in synovial cell nuclei. Whereas regulation of gene transcription by p50-containing dimers is complex — indeed it has been suggested that p50 homodimers can function as transcriptional repressors (11) — our data clearly demonstrate that the net effect of p50 deletion is a marked reduction of inflammatory joint disease.

In summary, our results show that Rel/NF-κB is essential for the development of inflammatory arthritis and provide strong evidence that selective blockade of Rel/NF-κB subunits may prevent major components of the disease. We also show that the absence of different Rel/NF-κB subunits can have differential effects in acute inflammatory arthritis suggesting distinct roles, rather than redundancy, for the various Rel/NF-κB subunits. Several strategies have been used to block global Rel/NF-κB activity in various disease models (31, 42, 49–51). However, targeting a specific Rel/NF-κB subunit, especially within a given anatomical compartment, may cause fewer side effects than those that inhibit all Rel/NF-κB functions. Such a specific approach with p65 antisense phosphorothioate oligonucleotides was used to prevent experimental colitis (52). Our results support several previous studies that indicate total Rel/NF-κB blockade would prevent inflammatory arthritis (31, 42, 49). However, we have demonstrated that specific Rel/NF-κB subunits may play distinct roles at different stages of disease. Consequently, local inhibition of p50 may prevent joint inflammation and the synovial tissue responses that lead to joint destruction. We conclude that p50 and genes regulated by p50 are targets for rational drug design in inflammatory arthritis.

Acknowledgments

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