Rheumatoid arthritis (RA) is a chronic inflammatory disease characterized by autoimmunity and infiltration of joint synovium by activated inflammatory cells, including neutrophils, macrophages, lymphocytes, plasma cells, and osteoclasts (reviewed in ref. 1). Morbidity in RA occurs secondary to the destruction of cartilage and bone by an inflammatory mass termed the pannus, which consists predominantly of activated macrophages and proliferating synovial fibroblasts. Synovial inflammation is regulated by cytokines (1). Pro-inflammatory cytokines implicated in RA pathogenesis include TNF-α, IL-1, IL-6, IL-12, IL-15, IL-18, IFN-γ, and GM-CSF. A key role for TNF-α and IL-1 in RA pathogenesis has been confirmed by the success of TNFα and IL-1 neutralization in the treatment of RA. RA synovium is also characterized by the expression of anti-inflammatory cytokines and cytokine antagonists, including TGF-β, IL-10, the IL-1 receptor antagonist, and soluble TNF receptors. Production of these factors reflects an attempt to control inflammation and achieve homeostasis. The idea that the balance between pro-and anti-inflammatory factors is important in regulating the rate of progression, and thus the eventual severity and morbidity, of RA has gained broad acceptance among RA researchers (1).

### Inhibition of cytokine signaling by suppressor of cytokine signaling proteins

The current paradigm states that cytokine balance is determined by the relative levels of expression of pro-inflammatory and anti-inflammatory cytokines. However, cytokine action is modulated by feedback inhibition and signal transduction crosstalk among synergistic or opposing cytokines. Thus the regulation of cellular responsiveness to cytokines determines cytokine activity and the balance between opposing cytokines. Key regulators of cellular responses to cytokines are members of the suppressors of cytokine signaling (SOCS; also known as cytokine-induced SH2-containing protein, Jak binding protein, or STAT-induced STAT inhibitor) protein family, which plays an important role in feedback inhibition and cytokine cross-regulation (2, 3). SOCS were originally discovered, and are best known, as inhibitors of the Janus kinase–signal transducer and activator of the transcription (JAK-STAT) signaling pathway. SOCS inhibit JAK-STAT signaling by: (i) inhibiting JAK catalytic activity; (ii) competing with STATs for receptor docking sites, and (iii) targeting cytokine receptors for degradation by proteosomes. SOCS-1, SOCS-2, SOCS-3, and cytokine-induced SH2-containing protein expression is rapidly induced by many cytokines, including cytokines that activate the JAK-STAT pathway. Individual SOCS are capable of inhibiting multiple cytokines, and it is not clear how the specificity of inhibition by SOCS is regulated.

### JAK-STAT signaling and inflammatory arthritis

STATs are latent cytoplasmic proteins that are activated by tyrosine phosphorylation, which leads to dimerization, acquisition of DNA binding activity, and translocation to the nucleus and transcriptional activation of genes (4). Several cytokines implicated in RA pathogenesis activate STATs, namely, IL-6 family cytokines (activate STAT3), IL-12 (STAT4), IL-15 (STAT3 and STAT5), GM-CSF (STAT5), and IFN-γ (STAT1). STAT1 and STAT4 expression is elevated, and STAT1 and STAT3 are activated, in the synovium in RA and experimental arthritis (refs. 5–8, L. Ivashkiv and A. Koch, unpublished observations). Ablation of STAT3 activity using a dominant negative STAT3 mutant results in apoptosis of synovial fibroblasts (9), and gene transfer of a dominant negative STAT3 mutant to synovial macrophages and fibroblasts attenuates collagen-induced arthritis (6), demonstrating a role for STAT3 in the pathogenesis of inflammatory arthritis. Overexpression of SOCS-3 in joint macrophages and fibroblasts has been demonstrated to effectively block STAT3 activation during experimental arthritis and to attenuate disease severity (6). Thus, SOCS-3 overexpression is effective in attenuating arthritis, but the role of endogenous SOCS expression and the mechanisms by which SOCS-3 overexpression suppresses arthritis have not been elucidated. The paper by Egan and colleagues in this issue of the JCI (10) investigates the role of endogenous SOCS-1 in modulating acute arthritis induced by the injection of methylated BSA and IL-1 (mBSA/IL-1 arthritis).

### SOCS-1 and arthritis

mBSA/IL-1 arthritis is an acute arthritis that peaks seven days after induction and resolves by day 21. Similarities to RA include infiltration of joints by myeloid cells, pannus formation, and a combination of inflammation and tissue destruction. mBSA/IL-1 arthritis is independent of B cells or CD8+ T cells, but is dependent upon...
CD4+ T cells, and is modulated by TNF-α, GM-CSF, and IL-6. The authors found that mBSA/IL-1 arthritis was independent of IFN-γ. Mechanisms by which CD4+ T cells drive inflammatory arthritis and synovial infiltration and activation of macrophages in the absence of IFN-γ may be important in RA pathogenesis (11) and can be delineated using this model. The independence of mBSA/IL-1 arthritis from IFN-γ allowed the authors to address the role of SOCS-1 using SOCS-1–/– IFN-γ–/– double-knockout mice, which do not develop the severe spontaneous inflammation and perinatal lethality manifested by SOCS-1–/– mice (12–14). Deficiency of SOCS-1 resulted in increased severity but no change in the time course of arthritis. SOCS-1–/– IFN-γ–/– arthritic mice exhibited increased myeloid cell infiltration of synovium and adenopathy of draining lymph nodes, with hyperproliferation of T cells. Immunohistochemistry, using a SOCS-1 antibody and measuring the activity of a lacZ reporter gene that was integrated into the SOCS-1 locus, indicated that the SOCS-1 gene was expressed in synovial macrophages, lymphocytes, and fibroblasts, and in activated lymph node T cells, during the course of mBSA/IL-1 arthritis.

These results clearly identify SOCS-1 as an important negative regulator of acute arthritis in the mBSA/IL-1 arthritis model. Increasing SOCS-1 expression in the appropriate cells at the appropriate time may represent a promising new approach to the treatment of inflammatory arthritis. This study also raises a number of interesting questions, several of which can be readily addressed in the mBSA/IL-1 arthritis model. One important question is in which cells and tissues SOCS-1 exerts its antiarthritic activity. Previous work showed that SOCS-1–/– lymphocytes are both necessary and sufficient for the spontaneous lethal inflammatory syndrome manifested by SOCS-1–/– mice (14). Thus it is likely that suppression of T cell expansion and activation in draining lymph nodes is an important mechanism of SOCS-1 suppression of mBSA/IL-1 arthritis. This represents an interesting contrast to SOCS-3, which suppresses arthritis when overexpressed in synovial macrophages and fibroblasts (6). Thus, different SOCS proteins may suppress different components of autoimmunity and inflammation that are important in arthritis.

Another important question is which cytokine(s) need to be inhibited by SOCS-1 in order to attenuate arthritis? SOCS proteins are best known as inhibitors of JAK-STAT signaling, but they also interact with the adaptors protein Grb2, the phosphoprotein Vav, calcineurin, and IL-1 receptor–associated kinase, and thereby suppress signaling by Kit, the T cell antigen receptor, and Toll-like receptor 4 (TLR4), which shares a major signaling pathway with IL-1 (15–18). In addition, SOCS can potentiate Ras-MAPK pathways and suppress TNF-induced apoptosis (19, 20). Thus, SOCS-1 may suppress signaling by many different cytokines in mBSA/IL-1 arthritis (see Figure 1), and the cytokines that are suppressed will vary according to cell type and inflammatory microenvironment. In the draining lymph nodes, SOCS-1 most likely functions by suppressing signaling by cytokines important for T cell survival and proliferation, such as IL-6 and IL-2. In the synovium, SOCS-1 could potentially suppress signaling by multiple pro-inflammatory cytokines, including endogenous IL-1, IL-18, IL-6, GM-CSF, and endogenous TLR ligands, and could alter TNF signaling. In addition, because SOCS-1 binds to IL-1 receptor–associated kinase and suppresses TLR pathways that are also used by IL-1 (17, 18), SOCS-1 can inhibit the action of exogenous IL-1 during induction of mBSA/IL-1 arthritis. In summary, SOCS-1 may suppress inflammatory arthritis by inhibiting signaling by multiple cytokines.

SOCS-1 deficiency results not only in enhanced cellular responses to cytokines, but also in increased cytokine production in vivo, including elevated basal serum levels of IFN-γ and TNF-α (14, 19). Although Egan et al. (10) did not detect any changes in IL-1, IL-6, or IL-18 mRNA levels in synovium in SOCS-1–/– IFN-γ–/– arthritic mice relative to control arthritic mice, it will be important to carefully examine the effects of SOCS-1 deficiency on cytokine production in both synovium and draining lymph nodes. It is also interesting to speculate about which cytokines induce SOCS-1 expression during arthritis (Figure 1). Possibilities that come to mind include IL-6 and IL-10 in T cells in lymph nodes (21, 22), and IL-6, IL-10, TNF, IL-1, and endogenous TLR ligands in synovium. Of note, most of these are pro-inflammatory cytokines that induce SOCS expression as part of feedback inhibition, except for IL-10 that is anti-inflammatory. The roles of various cytokines in the induction of SOCS expression can be addressed using appropriate knockout mice.

In summary, this interesting paper (10) identifies SOCS-1 as an important suppressor of acute inflammatory arthritis and suggests novel approaches to arthritis therapy based on increasing SOCS-1 expression. The important role of SOCS-1 in modulating arthritis severity emphasizes the...
importance of feedback inhibition and signal transduction cross-regulation in synovial inflammation. Further study of mechanisms by which SOCS proteins regulate inflammatory arthritis are likely to yield important insights into RA pathogenesis and cytokine regulatory networks in complex inflammatory diseases.


See the related article beginning on page 907.

**Autonomic “myasthenia”: the case for an autoimmune pathogenesis**

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Of all neurological diseases, the one that virtually all third year medical students get right on final exams is myasthenia gravis (MG). MG is an autoimmune disorder that causes weakness and fatigue of skeletal muscles due to an antibody-mediated attack directed against AChRs at neuromuscular junctions (1). It is easy to learn and remember the pathogenesis, immunology, and treatment of MG because the pieces of the puzzle fit together nearly perfectly. In this issue of the JCI, Lennon et al. now provide intriguing evidence of another putative autoimmune disease of AChRs — autoimmune autonomic neuropathy (AAN) (2). They have shown that immunization of rabbits with a fragment of ganglionic ACHR induces a condition that mimics the human disease. AAN is very rare but can be life threatening. It is manifested by autonomic disturbances affecting the sympathetic and parasympathetic nervous systems: abnormal gastrointestinal motility with stalled traffic in the intestines, impaired contraction and striking dilatation of the bladder, and disturbances of blood pressure, sweating, salivation, pupillary reactions, and sexual function (3). Based on these experiments and other evidence (4), Lennon et al. suggest that: (a) an antibody-mediated attack directed against neuronal AChRs of autonomic ganglia may be implicated in at least some cases of AAN; and (b) the autoimmune response in some of these patients may be triggered by a remote neoplasm that incidentally expresses the autoantigen. These new findings provide interesting insights into the concepts of autoimmune and paraneoplastic diseases of the nervous system, and information that may be useful for practical treatment of AAN.

**Five criteria for recognizing antibody-mediated autoimmune disease**

The list of candidate autoimmune diseases affecting every level of the nervous system is long and rapidly increasing. However, it is not simple to prove that a given disease is autoimmune in nature, and even more difficult to identify the autoimmune mechanisms and molecular

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See the related article beginning on page 907.