

Review Article

Role of SOCS proteins in inflammation and autoimmune diseases

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Various cytokines are involved in the regulation of the immune system and inflammation. Dysregulation of cytokine signaling can cause a variety of diseases, including allergies, autoimmune diseases, inflammation, and cancer. Most cytokines utilize the so-called Janus kinase-signal transducers and activators of transcription (JAK-STAT) pathway. This pathway is negatively regulated by suppressors of cytokine signaling (SOCS) proteins. SOCS proteins bind to JAK or certain cytokine receptors, thereby suppressing further signaling events. Studies using conditional knockout mice have shown that SOCS proteins are key physiological pathological regulators of inflammation as well as immune homeostasis. Recent studies have also demonstrated that SOCS1 and SOCS3 are important regulators of adaptive immunity, especially helper T cell differentiation.

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The JAK/STAT pathway and SOCS proteins

Cytokines play several essential roles in the development, differentiation, and function of myeloid and lymphoid cells. Some of them, including interleukins, interferons (IFNs), and hematopoietic growth factors, activate the Janus kinase (JAK)-signal transducers and activators of the transcription (STAT) pathway^{1,2)}. In this pathway, cytokine binding results in receptor oligomerization, which initiates the activation of JAK kinases (JAK1, JAK2, JAK3, and Tyk2). The activated JAKs phosphorylate the

receptor cytoplasmic domains, which creates docking sites for SH2-containing signaling proteins, including STATs. It is now known that a large number of cytokines, growth factors, and hormonal factors activate JAK and/or STAT proteins. For example, pro-inflammatory cytokine IL-6 binds to the complex of the IL-6 receptor α chain and gp130, which mainly activate JAK1 and STAT3. IFN γ receptors utilize JAK1 and JAK2, then mainly activate STAT1. Interestingly, the anti-inflammatory cytokine IL-10 also activates STAT3. IL-6- or IL-23-mediated STAT3 activation, IL-12-mediated STAT4, and IL-4-mediated

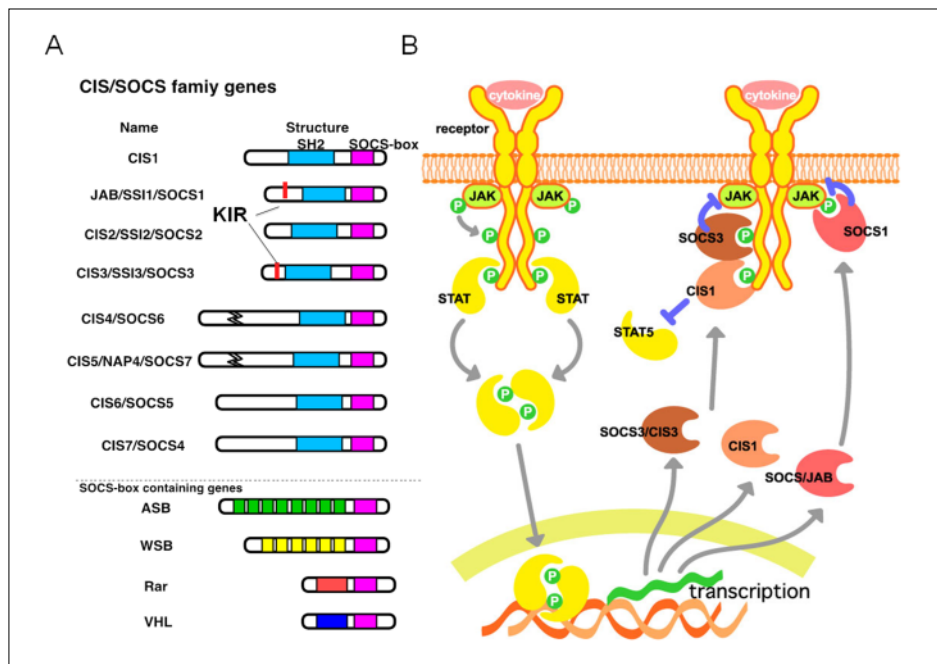


Fig.1 The structure and function of SOCS proteins

(A) Schematic structure of the CIS/SOCS family proteins. The SOCS box is conserved in all CIS/SOCS family proteins. SOCS1 and SOCS3 contain a unique kinase inhibitory region (KIR) immediately upstream of the central SH2 domain, which is proposed to function as a pseudosubstrate. Other SOCS box-containing proteins are illustrated.

(B) Mechanism of suppression by CIS, SOCS1, and SOCS3. All of these are induced by cytokine stimulation. CIS binds to the STAT5 activating receptors, thereby suppressing further activation of STAT5 and inducing degradation of the receptor. SOCS1 binds to JAKs, and SOCS3 binds to the receptor through the SH2 domain, but both inhibit JAK activity through KIR. These complexes may be degraded by ubiquitination and proteasomal degradation recruited through the SOCS box.

STAT6 activation have been demonstrated to be essential for Th17, Th1, and Th2 differentiation, respectively³. IL-2-mediated STAT5 activation is essential for T cell proliferation as well as regulatory T cell (Treg) development⁴. The action of STAT5 also appears to be very direct, as STAT5 binds the *Foxp3* gene, which is the master regulator of Tregs (see Fig.2). IL-6/STAT3 and IL-4/STAT6 inhibits *Foxp3* gene expression, although the mechanism remains controversial^{5,6}.

Although our understanding of the intracellular signaling molecules that mediate the functional outcome of cytokine-receptor activation in the immune system has increased significantly, recent research has placed increasing emphasis on the mechanisms for the termination of signals and cross-talks with other signaling pathways. The CIS/SOCS family of proteins is one of the major mechanisms for such regulations⁷⁻⁹. At the time of their discovery, the SOCS proteins were recognized as an important

mechanism in the negative regulation of the cytokine-JAK-STAT pathway, but recent studies using gene-disrupted mice have revealed that they play additional unexpected and important roles in many immunological processes¹⁰⁻¹³, atherosclerosis¹⁴, metabolism^{15,16}, and cancer^{13,17-19}. In this review, we will focus on the recent progress of SOCS studies on inflammation and helper T cell differentiation.

The CIS/SOCS family

The suppressor of cytokine signaling (SOCS) proteins and cytokine-inducible SH2-containing (CIS; also known as CISH) protein molecules comprise a family of intracellular proteins^{7,11,20}. There are eight CIS/SOCS family proteins: CIS, SOCS1, SOCS2, SOCS3, SOCS4, SOCS5, SOCS6, and SOCS7, each of which has a central SH2 domain, an amino-terminal domain of variable length and sequence, and a carboxy-terminal 40-amino-acid

module known as the SOCS box (Fig. 1A)²¹. The SOCS box is also found in other miscellaneous proteins. The SOCS box interacts with elongin B and elongin C, Cullins, and the RING-finger-domain-only protein RBX2 (which recruits E2 ubiquitin-transferase)^{22,23}. VHL (von Hippel-Lindau) gene product, whose gene product is the principal negative regulator of hypoxia-inducible factor has been shown to bind to SOCS1 and induces the degradation of Jak2. Chuvash polycythemia-associated VHL mutants have altered affinity for SOCS1 and do not engage with and degrade phosphorylated JAK2²⁴. These results indicate that CIS/SOCS family proteins, as well as other SOCS box-containing molecules, function as E3 ubiquitin ligases and mediate the degradation of proteins that are associated with these family members through their N-terminal regions (Fig. 1A).

The central SH2 domain determines the target of each SOCS and CIS protein. The SH2 domain of SOCS1 directly binds to the activation loop of JAKs²⁵. The SH2 domains of CIS, SOCS2, and SOCS3 bind to phosphorylated tyrosine residues on activated cytokine receptors⁸. SOCS3 binds to gp130-related cytokine receptors, including the phosphorylated tyrosine 757 (Tyr757) residue of gp130, the Tyr800 residue of IL-12 receptor β 2, and Tyr985 of the leptin receptor²⁶⁻³⁰. Thus, SOCS3 in the brain has been implicated in leptin-resistance^{15,31}. SOCS molecules bind to several tyrosine phosphorylated proteins, including Mal (TLR signaling)³² and IRS1/2 (insulin signaling)¹⁶. Thus, SOCS proteins generally induce the degradation of the target molecules by binding through the SH2 domain and ubiquitination through the SOCS box (Fig. 1B).

In addition, both SOCS1 and SOCS3 can inhibit JAK tyrosine kinase activity directly through their kinase inhibitory region (KIR). KIR has been proposed to function as a pseudosubstrate that is essential for the suppression of cytokine signals²⁵. The SH2 domain of SOCS3 does not have a high affinity to the activation loop of JAKs yet the KIR of SOCS3 has a higher affinity to the kinase domain of JAK2 than that of SOCS1³³. Because the receptors to which SOCS3 binds mostly activate STAT3, SOCS3 is an inhibitor that is relatively specific to STAT3. SOCS3 also inhibits STAT4, which is activated by IL-12²⁹. However, because SOCS3 does not bind to the IL-10 receptor, SOCS3 cannot inhibit IL-10 signaling. Therefore, IL-10 induces a robust and prolonged STAT3 activation, whereas IL-6-mediated STAT3 activation is transient in macrophages. This is an important mechanism to distinguish the anti-inflammatory activity of IL-10 and inflammatory activity of IL-6²⁰.

SOCS1 and inflammation

Although *SOCS1* knockout (KO) mice are normal at birth, they exhibit stunted growth and die within 3 weeks of birth, with activation of peripheral T cells, necrosis of the liver, and macrophage infiltration of major organs³⁴. The neonatal defects exhibited by *SOCS1*^{-/-} mice appear to occur primarily as a result of unbridled IFN γ signaling, since *SOCS1*^{-/-} mice that also lack the IFN γ gene or the IFN γ receptor gene do not die neonatally. Since *SOCS1/Rag2*-double knockout (DKO) mice survived much longer, *SOCS1* has been thought to be an important negative regulator of Th1. This is confirmed by generating T cell-specific *SOCS1*-conditional KO (cKO) mice³⁵.

In addition, *SOCS1* has been demonstrated to be involved in the suppression of inflammation by regulating innate immune cells and nonimmune cells. Using liver-specific *SOCS1*-cKO mice, we demonstrated that *SOCS1* deletion in hepatocytes enhanced concanavalin A (ConA)-induced hepatitis, because proapoptotic signals, including STAT1 and JNK activation, were enhanced in *SOCS1*-deficient liver³⁶. *SOCS1* deletion in NKT cells also enhanced sensitivity to ConA-induced hepatitis. However, the number of iNKT cells was drastically decreased but that of type II NKT cells was increased by *SOCS1* deficiency³⁷. This mechanism remains to be clarified. Deficiency of *SOCS1* in macrophages cells resulted in hyper-responses to lipopolysaccharide (LPS)³⁸⁻⁴⁰ and *SOCS1*-deficient dendritic cells (DCs) promoted hyperactivation of Th1, lupus-like autoimmune diseases, and anti-tumor immunity^{41,42}.

SOCS1 has been implicated in the mechanism of glucocorticoid-mediated STAT1 suppression^{43,44}. *SOCS1* is also highly upregulated by *M. tuberculosis* infection and reduced responses to IL-12, resulting in an impaired IFN- γ secretion by macrophages that in turn accounts for deteriorated intracellular mycobacterial control. Thus, *SOCS1* expression by macrophages hampered *M. tuberculosis* clearance early after infection *in vivo* in an IFN- γ -dependent manner. On the other hand, at later time points, *SOCS1* expression by non-macrophage cells protected the host from infection-induced detrimental inflammation⁴⁵. We have demonstrated that *SOCS1* plays an essential role in intestinal immune homeostasis by regulating prostaglandin E2 (PGE2)-mediated DC and macrophage suppression¹². Although *SOCS1/Rag2* DKO mice did not die neonatally, these mice developed severe colitis at 2 to 6 months of age, mostly due to impairment of the PGE2-mediated anti-inflammatory mechanism⁴⁶.

SOCS3 and inflammation

In contrast to *SOCS1*, the role of *SOCS3* in innate inflamma-

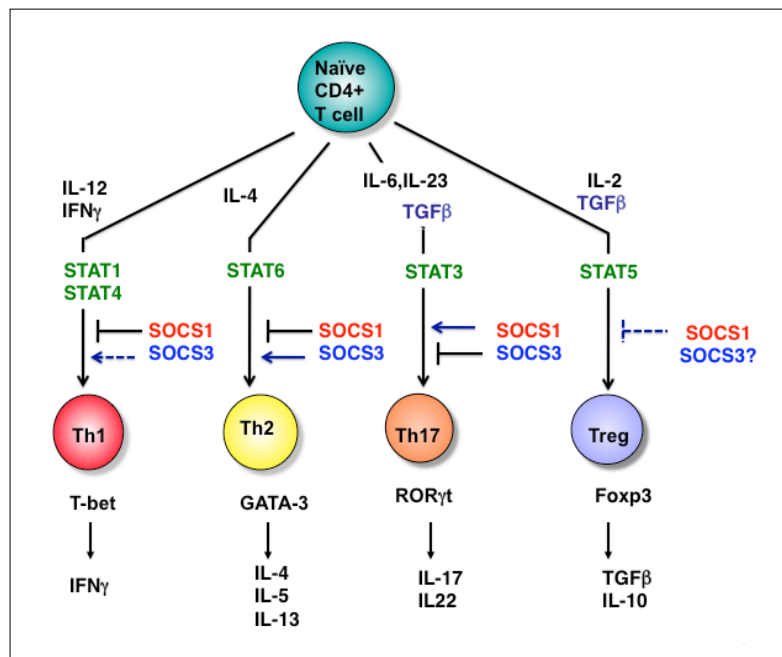


Fig.2 Role of STATs and SOCS proteins in helper T cell differentiation

T cell differentiation from naïve cells into the various functional subtypes (e.g., Th1, Th2, Th17, and Treg cells) primarily depends on the action of cytokines as indicated. SOCS1 and SOCS3 regulate the cytokine pathways indicated, and thereby dictate CD4⁺ T cell differentiation. Note that SOCS3 inhibits Th1 through overexpression by suppressing IL-12-mediated STAT4 activation. However, SOCS3 deletion also inhibits Th1 by promoting the production of IL-10 and TGF- β . In Tregs, SOCS1 deletion promotes Treg expansion; however, these are functionally impaired. Thus, SOCS1 negatively regulates the Treg number, but is necessary for the function of Tregs. The role of SOCS3 in Tregs has not yet been clarified. For Th17, ROR γ t induction by TGF- β has been demonstrated to be independent of Smad2/3/4.

tion is complex. SOCS3 deficiency in macrophages protects mice from endotoxemia, because of the reduced production of inflammatory cytokines, which is due to the enhanced anti-inflammatory effect of STAT3²⁰. Furthermore, macrophage-specific SOCS3-cKO mice have reduced IL-12 responses and succumb to toxoplasmosis. In the absence of SOCS3, macrophages are hypersensitive to the anti-inflammatory properties of IL-6. Thus, SOCS3 plays a critical role in suppressing IL-6 signals and promoting immune responses to control *T. gondii* infection⁴⁷. Macrophage-specific SOCS3-cKO mice exhibited resistance to the tumor transplantation model because of reduced tumor-promoting cytokines such as TNF α and IL-6 and enhanced production of anti-tumorigenic chemokine MCP2/CCL8¹³. Thus, SOCS3 is an important modulator of macrophage phase and functions.

SOCS3^{-/-} DCs exhibited constitutive activation of STAT3 and expressed low levels of MHC class II molecules, co-stimulatory molecules, and IL-12⁴⁸. Adoptive transfer of SOCS3^{-/-} DCs suppressed EAE. SOCS3^{-/-} DCs produced a higher amount of TGF- β than WT DCs, resulting in a selective expansion of forkhead box P3 (FoxP3)-positive regulatory T cells (Tregs). However, SOCS3-transduced DCs also expressed low levels of MHC class II and CD86 molecules and produced high levels of IL-10 but low levels of IL-12, IFN γ , and IL-23 p19⁴⁹. STAT3 activation was suppressed by SOCS3 overexpression. Although the mechanism has not yet been clarified, SOCS3-transduced DCs efficiently induced Th2-cell differentiation and suppressed Th17 *in vitro* and *in vivo* and the adoptive transfer of *Socs3*^{-/-} DCs suppressed EAE, just like SOCS3-Tg DCs⁴⁹. These results suggest

that the status of STAT3 activation levels may determine the balance between Th2 and Tregs induced by DCs.

In certain situations, SOCS3 suppresses inflammatory reactions by inhibiting STAT3. STAT3 activation is found in epithelial and lamina propria cells in the colon of mice with intestinal bowel disease (IBD), as well as in human ulcerative colitis and Crohn's disease patients⁵⁰ and in synovial fibroblasts of RA patients⁵¹. Forced expression of either SOCS3 or a dominant negative form of STAT3 in mouse arthritis models suppressed the induction/development of the disease, indicating that SOCS3 in non-immune cells is probably anti-inflammatory⁵¹. These findings are consistent with the idea that the IL-6- and IL-6-related cytokines-STAT3 pathway promotes chronic disease progression and SOCS3 is part of this negative-feedback loop. This idea is supported by a recent finding that the JAK inhibitor CP-690550 is a potent therapeutic agent for the autoimmune arthritis model by suppressing the IL-6/STAT3 amplification⁵². However, if STAT3 plays a protective role, such as in ConA-induced hepatitis, deletion of SOCS3 is anti-inflammatory¹⁸.

In addition, SOCS3 is an important negative regulator of granulopoiesis because SOCS3 negatively regulates the G-CSF receptor signaling^{53,54}. Mice in which the SOCS3 gene was deleted in all hematopoietic cells developed a spectrum of inflammatory pathologies with hyper-neutrophilia. SOCS3-deficient mice developed inflammatory neutrophil infiltration into multiple tissues and consequent hind-leg paresis⁵⁵.

Overview of helper T cell differentiation

Helper T cells play essential roles in adaptive immune responses and chronic inflammation. After emigrating from the bone marrow, thymocyte progenitors enter the thymus. Following positive selection, single positive (CD4⁺ or CD8⁺ SP) cells migrate to the periphery as naïve T cells. Naturally occurring CD4⁺CD25⁺ Foxp3⁺regulatory T cells (nTregs) also develop in the thymus from immature CD4⁺ T cells. After exiting the thymus, naïve T cells are activated by antigen-presenting cells (APCs), and develop specialized properties and effector functions, Th1, Th2, and Th17 (Fig.2)⁵⁶.

Th1 cells, which evolved to enhance the eradication of intracellular pathogens (e.g., intracellular bacteria, viruses, and some protozoa), are characterized by their production of IFN γ a potent activator of cell-mediated immunity. IL-12 is an inducer of Th1. Immune pathogenesis that results from dysregulated Th1 responses to self or commensal floral antigens can promote tissue destruction and chronic inflammation, macrophage activation, and granuloma. Th2 cells, which evolved to enhance the

elimination of parasitic infections (e.g., helminths), are characterized by the production of interleukin (IL)-4, IL-5, and IL-13, which are potent activators of B-cell immunoglobulin (Ig)E production, eosinophil recruitment, and mucosal expulsion mechanisms (mucous production and hypermotility). IL-4 is necessary to induce Th2. Dysregulated Th2 responses can cause allergies and asthma. Th17 cells secrete a distinctive set of immunoregulatory cytokines, including IL-17A, IL-17F, IL-22, and IL-21. These cytokines collectively play roles in inflammation and autoimmunity and in the elimination of extracellular bacterial and fungal pathogens. Th17 cells have been demonstrated to play essential roles in autoimmune models, such as experimental autoimmune encephalitis (EAE) and collagen-induced arthritis (CIA). The Th17 differentiation of naïve T cells is initiated by IL-6 and TGF- β . In addition, IL-23, as well as IL-21, is thought to be a key cytokine for the maturation and/or maintenance of Th17 cells. IL-6, IL-21, and IL-23 all activate STAT3, which is thought to be essential for Th17 differentiation^{57,58}. It has also been reported that STAT3 plays a critical role in the induction of the orphan nuclear receptor, ROR γ t, which directs Th17 cell differentiation by inducing the IL-23 receptor⁵⁹. TGF- β also induces the differentiation of naïve T cells into Foxp3⁺ Tregs (iTregs) in the peripheral immune compartment, and iTreg and Th17 are reciprocally regulated^{60,61} (Fig.2).

SOCS1 and helper T cell differentiation and functions

We have recently demonstrated that SOCS1 is an essential regulator for helper T cell differentiation. Most SOCS1^{-/-}CD4 naïve T cells differentiated into Th1, even under Th2 or Th17 skewing conditions, whereas Th17 differentiation was strongly suppressed³⁵. This was also dependent on IFN γ , because Th17 was normally developed in SOCS1^{-/-}IFN γ ^{-/-} T cells. As a result, T cell-specific SOCS1-deficient mice developed autoimmune inflammatory diseases with age¹⁰ and were very sensitive to dextran sulfate sodium (DSS)-induced colitis⁶² and ConA-induced hepatitis (Th1 type disease)³⁷, but were resistant to experimental autoimmune encephalomyelitis (EAE), a typical Th17 type disease³⁵.

Th17 suppression by SOCS1 deficiency is probably due to the hyperproduction and signal transduction of IFN γ . Indeed, STAT1 activation in SOCS1^{-/-} T cells was upregulated and strong Th1 skewing was corrected under STAT1^{+/-} conditions³⁵ (and unpublished data). Interestingly, STAT3 activation was reduced in SOCS1-deficient T cells, mostly due to the upregulation of SOCS3 gene expression, which can account for reduced IL-6

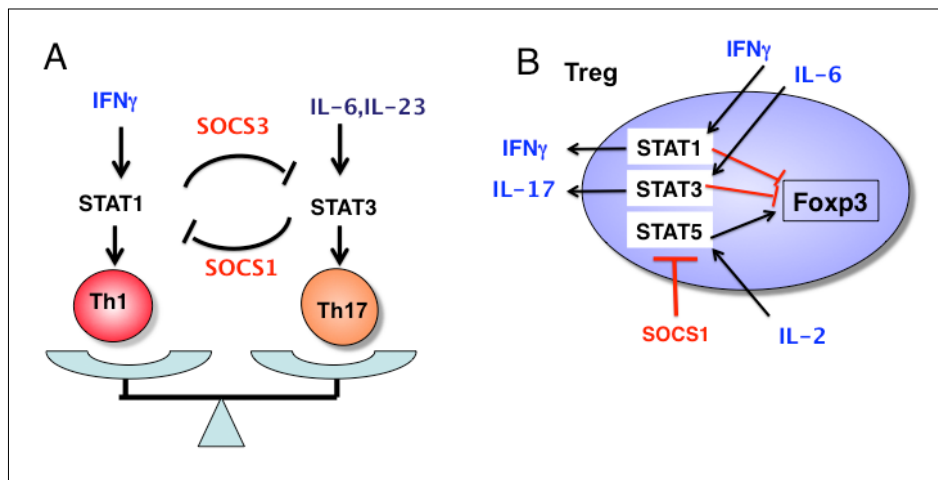


Fig.3 Role of SOCS1 and SOCS3 in Th1/Th17 balance (A) and in nTreg integrity (B)

(A) In Th1 conditions, activated STAT1 induces SOCS3 expression, which in turn inhibits STAT3 activity, thereby suppressing Th17 development. In Th17 conditions, SOCS1 is highly expressed and inhibits IFN γ -mediated Th1 development.

(B) In nTregs, the SOCS1 level is constantly high, and suppresses STAT1, STAT3, and STAT5 in inflammatory conditions (high levels of cytokines such as IL-2, IL-6, and IFN γ). The higher activation of STAT5 results in an increase in the number of Tregs, because STAT5 directly upregulates Foxp3 expression. The higher activation of STAT1 and STAT3 induces loss of Foxp3 expression by unknown mechanisms and promotes IFN γ and IL-17 production, respectively. Thus, SOCS1 is a “guardian” of Tregs.

responses and Th17 differentiation. Indeed, SOCS3-tg mice were resistant to EAE, and Th17 differentiation of SOCS3-tg T cells was suppressed. The reciprocal regulation of Th1 and Th17 by SOCS1 and SOCS3 is illustrated in Fig.3A. In addition, SOCS1^{-/-} T cells were less responsive to TGF- β , although the mechanism has not yet been clarified. Reduced STAT3 activation and TGF- β signaling may explain the suppression of Th17 differentiation in SOCS1-deficient T cells. Our microarray analysis revealed that T-bet, Eomesodermin, and Gfi-1 were upregulated in SOCS1-deficient Th17 cells, all of which have been reported to suppress Th17 differentiation^{63, 64}.

SOCS1 also plays an important role in the regulation of regulatory T cells. Higher numbers of Tregs are observed in the thymus and spleen of T cell-specific SOCS1-deficient mice⁶⁵. This is probably due to higher IL-2 responses, because IL-2 enhances the proliferation of Tregs (Fig.3B). Importantly, SOCS1 has been shown to be a target of miR-155 in Tregs⁶⁵. During thymic differentiation, the upregulation of Foxp3 drives the high expression of miR-155, which in turn promotes the expansion of Treg cells by targeting SOCS1.

However, SOCS1 has recently been found to play more important functional roles in Tregs. Various studies have suggested that Tregs may become harmful effector T cells in inflammatory conditions. Lu et al. observed that SOCS1 deletion specifically in Tregs induced the development of spontaneous dermatitis, splenomegaly, and lymphadenopathy, suggesting a defective Treg function in these mice⁶⁶. The defective suppression activity of SOCS1-deficient Tregs was confirmed through the failure to suppress colitis in *Rag2*^{-/-} mice by the co-transfer of naïve T cells and Tregs¹⁰. In the absence of SOCS1, Tregs easily lost Foxp3 expression, and became pathogenic T cells that induced severe colitis⁶⁶. In addition, SOCS1 plays an important role in preventing inflammatory cytokine production from Tregs. Normally, Tregs do not secrete inflammatory cytokines even in inflammatory conditions. In the absence of SOCS1, Tregs secrete IFN γ and IL-17 by hyperactivation of STAT1 and STAT3, respectively¹⁰. Thus, SOCS1 is a “guardian” of Tregs, since SOCS1 inhibits loss of Foxp3 and conversion of Tregs to Th1- or Th17-like cells (Fig.3B).

Role of SOCS3 in helper T cells

The degree to which SOCS3 expression in T cells is increased is correlated to the severity of human allergic diseases such as asthma and atopic dermatitis⁶⁷. The enhanced action of SOCS3 may promote allergic responses, since transgenic SOCS3 expression in T cells inhibits Th1 development and promotes Th2 development⁶⁷. Enhanced Th2 development may be due to the suppression of Th1 because IL-12 mediated Th1 differentiation by SOCS3 overexpression. Therefore, SOCS3-tg mice were sensitive to *L. Major* infection, where Th1 is necessary for eradication of this microbe⁶⁸. As described before, SOCS3-expressing T cells differentiated into Th17 cells less efficiently than WT T cells³⁵. In contrast, mice lacking SOCS3 in T cells result in reduced allergen-induced eosinophilia in the airways⁶⁹(Fig.3A). SOCS3 silencing with small interfering RNA (siRNA) in primary CD4⁺ T cells attenuated the Th2 response *in vitro* and *in vivo*⁷⁰. SOCS3 deficiency promoted Th17 differentiation in T cells⁷¹. Using VavCre-SOCS3 conditional KO mice, Wong et al. reported that the IL-1-induced inflammatory joint disease model was severely deteriorated in the absence of SOCS3 accompanying the enhanced IL-17 production from CD4⁺ T cells⁵⁵. SOCS3 deficiency in T cells reduced atherosclerotic lesion development and vascular inflammation, which was dependent on IL-17, whereas the overexpression of SOCS3 in T cells reduced IL-17 and accelerated atherosclerosis¹⁴. The absence of SOCS-3 in helper T cells therefore generally inhibits Th1 and Th2 by producing IL-10 and TGF- β , but had dramatic pro-inflammatory effects under Th17 conditions¹¹. The paradoxical effect of SOCS3 on T cell regulation is mostly due to the dual function of STAT3; it promotes the production of both inflammatory IL-17 and anti-inflammatory IL-10 and TGF- β .

In the LCMC clone 13 infection model, SOCS3 is highly induced in T cells, and T cell-specific SOCS3-deficient mice exhibit a profound augmentation of immunity and are protected from severe organ pathology, with an increase in the number of virus-specific CD8⁺T cells and an increase in the ability of CD4⁺T cells to secrete TNF- α and IL-17. This T cell-intrinsic SOCS3 induction has been implicated as a major factor contributing to immunological failure in the setting of chronic active infection⁷².

SOCS and human inflammatory diseases

It has been estimated that more than 20% of all malignancies are initiated or exacerbated by inflammation; for example, most human hepatocellular carcinomas (HCCs) are a consequence of hepatitis C virus (HCV) infection⁷³. The expression of *SOCS1* is often silenced in these tumors by hypermethylation of CpG

islands including HCCs. We found that silencing of *SOCS1* was frequently observed even in pre-malignant HCV-infected patients¹⁹. Liver injury is associated with hyperactivation of STAT1 and reduced activation of STAT3. Therefore, the reduced expression of SOCS1 may enhance tissue injury and inflammation through the hyperactivation of STAT1, promoting the turnover of epithelial cells and enhancing their susceptibility to oncogenesis. Therefore, SOCS1 is a unique anti-oncogene that prevents carcinogenesis by suppressing chronic inflammation^{36, 74}.

SOCS3 may also be involved in the development and progression of malignancies. SOCS3 expression levels were reduced in tumor areas of patients infected with HCV compared with non-tumor regions¹⁸. Hyperactivation of STAT3 by SOCS3 repression may contribute to tumorigenesis by inducing multiple tumor-promoting genes¹⁷.

As mentioned before, levels of SOCS3 in T cells are correlated to allergic diseases⁶⁷. Several genomic SNPs in the human *SOCS1* gene were found to be associated with serum IgE levels⁷⁵, asthma⁷⁶, and leukemia⁷⁷. SOCS1 mutations were found in human lymphomas⁷⁸.

Concluding remarks

Over the past decade, following the discovery of the SOCS protein families, we have extended our understanding of the structure and function of these proteins. SOCS proteins act as simple negative-feedback regulators, and they also play a part in the fine tuning of the immune response and inflammation. Therapeutic trials using SOCS anti-sense oligonucleotides, shRNA, and peptide mimetics are currently underway in animal models. SOCS1 and SOCS3 are ideal therapeutic targets for autoimmune diseases and inflammatory diseases, including cancer.

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Disclosure

We have no conflicts of interest.

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