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Stat5a Inhibits IL-12-Induced Th1 Cell Differentiation through the Induction of Suppressor of Cytokine Signaling 3 Expression

Hiroaki Takatori,† Hiroshi Nakajima,‡ Shin-ichiro Kagami,§ Koichi Hirose,† Akira Suto,† Kotaro Suzuki,§ Masato Kubo,‡ Akihiko Yoshimura,‡ Yasushi Saito,* and Itsuo Iwamoto*

In previous studies, we have shown that Th2 cell differentiation is diminished but Th1 cell differentiation is increased in Stat5a-deficient (Stat5a−/−) CD4+ T cells. In the present study, we clarified the molecular mechanisms of Stat5a-mediated Th cell differentiation. We found that enhanced Th1 cell differentiation and the resultant IFN-γ production played a dominant inhibitory role in the down-regulation of IL-4-induced Th2 cell differentiation of Stat5a−/− CD4+ T cells. We also found that IL-12-induced Stat4 phosphorylation and Th1 cell differentiation were augmented in Stat5a−/− CD4+ T cells. Importantly, the expression of suppressor of cytokine signaling (SOCS)3, a potent inhibitor of IL-12-induced Stat4 activation, was decreased in Stat5a−/− CD4+ T cells. Moreover, a reporter assay showed that a constitutively active form of Stat5a but not Stat6 activated the SOCS3 promoter. Furthermore, chromatin immunoprecipitation assays revealed that Stat5a binds to the SOCS3 promoter in CD4+ T cells. Finally, the retrovirus-mediated expression of SOCS3 restored the impaired Th cell differentiation of Stat5a−/− CD4+ T cells. These results suggest that Stat5a forces the Th1/Th2 balance toward a Th2-type by preventing IL-12-induced Th1 cell differentiation through the induction of SOCS3. The Journal of Immunology, 2005, 174: 4105–4112.

Over the last several years, significant progress has been made in the regulatory mechanisms of the transition of naive CD4+ T cells into mature Th2 cells (1–3). Whereas early studies have demonstrated that Th2 cell differentiation is essentially a Stat6-dependent process (4–6), recent studies have revealed that Stat6-independent pathways also participate not only in vitro Th2 cell differentiation (7) but also in vivo Th2 cell-mediated allergic airway inflammation (8, 9). Because the presence of IL-4-producing cells during T cell activation induces subsequent Stat6-dependent Th2 cell differentiation (1–3), it is inferred that Stat6-independent IL-4 production enhances the Stat6-dependent process of Th2 cell differentiation.

Regarding the Stat6-independent pathway, recent studies including ours indicate that Stat5a is involved in Th2 cell differentiation. We have previously shown that Ag-induced Th2 cytokine production and subsequent allergic airway inflammation are decreased in Stat5a-deficient (Stat5a−/−) mice (10). We have also shown that Th cell differentiation is biased toward a Th1-type at single cell levels in Stat5a−/− CD4+ T cells (11) and that the retrovirus-mediated expression of Stat5a restores the impaired Th2 cell differentiation of Stat5a−/− CD4+ T cells (11). Moreover, it has recently been demonstrated that the enforced expression of a constitutively active form of Stat5a induces IL-4 production in CD4+ T cells by regulating the accessibility of the IL-4 gene (12). These results suggest that the intrinsic expression of Stat5a within CD4+ T cells plays a critical role in Th2 cell differentiation and in the induction of allergic airway inflammation and that Stat5a may function as a direct inducer of IL-4 production. In addition, we have found that, by comparing Stat6−/− mice to Stat5a−/− and Stat6−/− double-deficient mice, Stat5a is indispensable in Stat6-independent Th2 cell differentiation of Stat6−/− CD4+ T cells (44). However, the molecular mechanisms underlying Stat5a-mediated Th cell differentiation are still largely unknown.

In the present study, we determined the molecular mechanisms underlying Stat5a-mediated Th cell differentiation. First, we found that IL-12-induced Stat4 phosphorylation and Th1 cell differentiation were enhanced in Stat5a−/− CD4+ T cells. Second, we found that the expression of suppressor of cytokine signaling (SOCS)3, a potent inhibitor of IL-12/Stat4 signaling (13, 14), was decreased in Stat5a−/− CD4+ T cells. Third, we found that Stat5a bound to SOCS3 promoter in CD4+ T cells and directly induced SOCS3 expression. Finally, we found that the retrovirus-mediated expression of SOCS3 restored the Th1/Th2 balance of Stat5a−/− CD4+ T cells. Taken together, our results indicate that Stat5a induces SOCS3 expression in CD4+ T cells and thus inhibits IL-12-induced Th1 cell differentiation, forcing the Th1/Th2 balance toward a Th2-type.

*Department of Allergy and Clinical Immunology, Chiba University School of Medicine, Chiba, Japan; †Laboratory for Signal Network, RIKEN Research Center for Allergy and Immunology (RCAI), Kanagawa, Japan; and ‡Division of Molecular and Cellular Immunology, Medical Institute of Bioregulation, Kyushu University, Fukuoka, Japan

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2 H.T. and H.N. contributed equally to this work.

3 Address correspondence and reprint requests to Dr. Hiroshi Nakajima, Department of Allergy and Clinical Immunology, Chiba University School of Medicine, 1-8-1 Inohana, Chiba City, Chiba 260-8670, Japan. E-mail address: nakajimh@faculty.chiba-u.jp

4 Abbreviations used in this paper: SOCS, suppressor of cytokine signaling; WT, wild type; ChIP, chromatin immunoprecipitation; LUC, luciferase; MGF, mammary gland factor.
Materials and Methods

Mice

Stat5α-deficient (Stat5α−/−) mice (15) were backcrossed to BALB/c mice (Charles River Breeding Laboratories) for eight generations and littermate wild-type (WT) mice were used as controls. All experiments were performed according to the guidelines of Chiba University (Chiba, Japan).

Cell culture

Splenocytes (2 × 10^6 cells/ml) from WT mice or Stat5α−/− mice were stimulated with plate-bound anti-CD3 mAb (5 µg/ml, clone 145-2C11; BD Pharmedics) in RPMI 1640 medium supplemented with 10% heat-inactivated FCS, 50 µM 2-ME, 2 mM L-glutamine, and antibiotics in a 24-well microtiter plate at 37°C for 48 h. As indicated, IL-12 (15 ng/ml; Peprotech) was added to polarize toward Th1 cells (Th1 condition), and IL-4 (15 ng/ml; PeproTech) and anti-IFN-γ mAb (15 ng/ml, clone XM12.2; BD Pharmedics) were added to polarize toward Th2 cells (Th2 condition) (11). Cells were washed with PBS and then cultured for another 72 h in Th1 or Th2 condition in the presence of IL-2 (10 ng/ml; PeproTech). In some experiments, either IL-4 (15 ng/ml) or anti-IFN-γ mAb (15 ng/ml) was added to the culture. In separate experiments, the indicated amounts of IL-12 (1–8 ng/ml) were added to the culture.

Flow cytometric analysis

Cells were stained and analyzed on a FACSCalibur (BD Biosciences) using CellQuest software. The following Abs were purchased from BD Pharmingen: anti-CD4 FITC, anti-CD4 PE, anti-CD4 PerCP (H129.19; BD Pharmingen) and anti-hamster IgG PE (BD Pharmingen), respectively. Before staining, FcRs were blocked with anti-CD16/CD32 Ab (2.4G2; BD Pharmingen). Negative controls consisted of isotype-matched, nonspecific Abs (BD Pharmingen).

Intracellular cytokine analysis

Intracellular cytokine staining for IFN-γ was performed as previously described (11). In brief, cultured splenocytes were restimulated with plate-bound anti-CD3 mAb at 37°C for 6 h, with monensin (2 µM; Sigma-Aldrich) added for the final 4 h. After FcRs were blocked with anti-CD16/CD32 Ab (2.4G2; BD Pharmingen), cells were stained with anti-CD4 PerCP (H129.19; BD Pharmingen), fixed with IC FIX (BioSource International), permeabilized with IC PERM (BioSource International), and stained with anti-CD4 PE (BV4D-1D11; BD Pharmingen) and anti-IFN-γ allophycocyanin (XM12.2; BD Pharmingen) at 4°C for 30 min. Cytokine profile (IL-4 vs IFN-γ) on CD4+ cells or CD4+ GFP+ cells (in the case of retrovirus experiments) was analyzed on a FACSCalibur.

Intracellular staining for phosphorylated Stat4

Intracellular staining for tyrosine-phosphorylated Stat4 was performed as described elsewhere (16) with a minor modification. In brief, spleenocytes from WT mice and Stat5α−/− mice were stimulated with plate-bound anti-CD3 mAb at 37°C for 48 h in Th1 or Th2 condition. Cells were washed and then cultured for another 72 h with fresh medium in Th1 or Th2 condition in the presence of IL-2 (10 ng/ml). Cells were stained from cytokines for 8 h and then stimulated with or without IL-12 (15 ng/ml) at 37°C for 20 min. Cells were stained with anti-CD4 FITC at 4°C, fixed with IC FIX, and permeabilized with 90% methanol and subsequently with IC PERM. Cells were then incubated with rabbit polyclonal anti-phospho-Stat4 Ab (Zymed Laboratories) or normal rabbit serum (as a control) for 30 min and visualized by anti-mouse IgG2a FITC (BD Pharmingen) and anti-hamster IgG PE (BD Pharmingen), respectively. Before staining, FcRs were blocked with anti-CD16/CD32 Ab (2.4G2; BD Pharmingen). Negative controls consisted of isotype-matched, nonspecific Abs (BD Pharmingen).

Western blot analysis

Splenocytes from WT mice or Stat5α−/− mice were stimulated with plate-bound anti-CD3 mAb for 48 h in Th2 condition. Cells were cultured for another 72 h in Th2 condition in the presence of IL-2. After CD4+ T cells were purified using anti-CD4 FITC and anti-FITC microbeads (BD Pharmingen) (>90% pure by flow cytometry) and rested for 8 h in the fresh medium, cells were stimulated with IL-12 (15 ng/ml) for 20 min and whole cell lysates were subjected to immunoblotting as previously described (17). Anti-phospho-Stat4 Ab and anti-mouse Stat4 Ab were purchased from Zymed Laboratories.

TagMan PCR analysis

Total cellular RNA was prepared as previously described (18). The expression levels of SOCS3 mRNA were determined by real-time PCR using a standard protocol on ABI PRISM 7000 instrument (Applied Biosystems). PCR primers and a fluorographic probe for SOCS3 were previously described (13). The levels of SOCS3 mRNA were normalized to the levels of GAPDH mRNA (Applied Biosystems).

Chromatin immunoprecipitation (ChIP) assay

ChIP assays were conducted using the ChIP Assay kit (Upstate Biotechnology) according to the manufacturer’s instruction with some modifications. Briefly, splenocytes from WT mice or Stat5α−/− mice were stimulated with plate-bound anti-CD3 mAb at 37°C for 48 h. CD4+ T cells were purified (>90% pure by flow cytometry) using a CD4+ T cell enrichment column (R&D Systems), starved in cytokines in fresh RPMI 1640 medium for 3 h, and then stimulated with IL-2 (20 ng/ml) at 37°C for 30 min. Cells were fixed with 1% formaldehyde to cross-link chromatin at room temperature for 15 min. Chromatin was then sonicated with SDS lysis buffer and then sonicated on ice to shear DNA to lengths between 500 and 700 bp. Sonicated lysates were centrifuged at 13,000 rpm for 4°C for 10 min, and the sonicated cell pellet was suspended in ChIP dilution buffer. The sonicated chromatin was then immunoprecipitated with anti-Stat5a antisera (R&D Systems) or control rabbit serum at 4°C for 12 h. The anti-Stat5a immunoepitopes were purified with protein A-agarose. After deproteinization and reverse-protonation and ligation of MmeI, the amounts of selected DNA sequences in the immunoprecipitates were assessed by PCR. The sequences of the primers for SOCS3 promoter were 5′-TTTTGTCCT CTTCTCGGTTA-3′ and 5′-GTTGAGTACGATGAGGAGG-3′. The sequences of the primers for β-globin promoter (as a control) were described elsewhere (19).

Luciferase assay

Stat6-dependent reporter plasmid, TPU474 (20), was a kind gift from Dr. U. Schindler (Tularik, San Francisco, CA). Stat6-dependent reporter plasmid, mammalian gland factor luciferase (MGF-Luc) (21), was a kind gift from Dr. H. Wakao (RIKEN Research Center for Allergy and Immunology, Kanagawa, Japan). Murine SOCS3 promoter (273 to +160) (22) was amplified by PCR using murine genomic construct as a template and inserted into pCMV/Bag/II site of pGL3-basic vector (Promega) to generate murine SOCS3-Luc. The reporter plasmid with Stat5-binding sites (PuG5) was then transfected into T cells (Promega) using FuGENE6 transfection reagents (Roche Diagnostics). Twenty-four hours after transfection, the luciferase activity was measured by the dual luciferase assay system (Promega). Firefly luciferase activity of TPU474, MGF Luc, SOCS3 WT Luc, SOCS3 mt1 Luc, SOCS3 mt2 Luc, or SOCS3 mt3 Luc with pRL-TK in the presence or absence of the expression vectors of a constitutively active form of Stat5a (16 Stat5a) (23) (pCDNA 16 Stat5a) or a constitutively active form of Stat6 (Stat6VT) (24) (pCDNA Stat6VT) using FuGENE6 transfection reagents (Roche Diagnostics). Twenty-four hours after transfection, the luciferase activity was measured by the dual luciferase assay system (Promega). Firefly luciferase activity of TPU474, MGF Luc, SOCS3 WT Luc, SOCS3 mt1 Luc, SOCS3 mt2 Luc, or SOCS3 mt3 Luc was normalized by Renilla luciferase activity of pRL-TK. All values were obtained from experiments conducted in triplicate and repeated at least four times.

Retrovirus-mediated gene expression

Bicistronic retrovirus vector pMX-IRES-GFP (23) was a kind gift from Dr. T. Kitamura (Tokyo University, Tokyo, Japan). pMX-SOCS3-IRES-GFP was previously described (14). Retroviruses were produced with a transient retrovirus packaging cell line, Plat-E (25), and stored at −80°C until use. For infection to T cells, putative retrovirus vector supernatants from WT mice or Stat5α−/− mice were stimulated with plate-bound anti-CD3 mAb for 40 h in the presence or absence of IL-4 (15 ng/ml), cells were incubated with 500 µl of the retrovirus in the presence of IL-2 (20 ng/ml) in a 24-well microtiter plate
that was coated with RetroNectin (27 μg/ml; Takara Shuzo). After 4 h of infection, 500 μl of fresh medium was added to the culture well and cells were allowed to grow for another 72 h in the presence or absence of IL-4 before being subjected to intracellular cytokine analysis. Under these conditions, the efficiency of infection to CD4+ T cells was 15–30% as assessed by GFP+ cells by FACS.

**Data analysis**

Data are summarized as mean ± SD. The statistical analysis of the results was performed by the unpaired t test. Values for p < 0.05 were considered significant.

**Results**

**IFN-γ plays a dominant inhibitory role in the down-regulation of Th2 cell differentiation in Stat5a−/− CD4+ T cells**

We have previously shown that Th2 cell differentiation is impaired but Th1 cell differentiation is increased in Stat5a−/− CD4+ T cells (11). Because Th1 cells suppress Th2 cell differentiation by producing cytokines such as IFN-γ (1–3), it is possible that the enhanced Th1 cell differentiation is a principal reason for the decreased Th2 cell differentiation of Stat5a−/− CD4+ T cells. To determine whether Th1 cytokines participate in the decreased Th2 cell differentiation in Stat5a−/− CD4+ T cells, we first examined the effect of neutralizing anti-IFN-γ Ab on Th2 cell differentiation in Stat5a−/− CD4+ T cells. After a 5-day culture of anti-CD3-stimulated splenocytes, Th2 cell differentiation was significantly decreased but Th1 cell differentiation was increased in Stat5a−/− CD4+ T cells as compared with those in WT CD4+ T cells (Fig. 1, a and b). IL-4 alone could not significantly induce Th2 cell differentiation of Stat5a−/− CD4+ T cells (Fig. 1, a vs b). In contrast, in the presence of anti-IFN-γ Ab, IL-4 strongly induced Th2 cell differentiation in Stat5a−/− CD4+ T cells (Fig. 1a). Also, IL-4 itself significantly induced Th2 cell differentiation of WT CD4+ T cells (Fig. 1, a vs e) and the Th2 cell differentiation was further increased in the presence of anti-IFN-γ Ab (Fig. 1g). These results indicate that enhanced Th1 cell differentiation and the resultant IFN-γ production play a dominant inhibitory role in the down-regulation of IL-4-induced Th2 cell differentiation of Stat5a−/− CD4+ T cells. In addition, even in the presence of anti-IFN-γ Ab, the frequency of Th2 cells was still lower in Stat5a−/− CD4+ T cells than that in WT CD4+ T cells (Fig. 1, g vs h), suggesting that other mechanisms are also involved in Stat5a-induced Th2 cell differentiation.

**IL-12-induced Th1 cell differentiation is enhanced in Stat5a−/− CD4+ T cells**

IL-12 plays a critical role in the induction of Th1 cell differentiation (26). To determine whether IL-12/Stat4 signaling plays a causative role in the enhanced Th1 cell differentiation in Stat5a−/− CD4+ T cells, we first examined the sensitivity of Stat5a−/− CD4+ T cells to IL-12-induced Th1 cell differentiation. As shown in Fig. 2, even in the absence of exogenous IL-12, the frequency of Th1 cells after a 5-day culture of anti-CD3-stimulated splenocytes was significantly higher in Stat5a−/− CD4+ T cells than that in WT CD4+ T cells (n = 4, p < 0.05). Importantly, a low concentration (1 ng/ml) of IL-12 significantly increased the number of Th1 cells in Stat5a−/− CD4+ T cells but not in WT CD4+ T cells (n = 4, p < 0.01), whereas a high concentration of IL-12 (4 or 8 ng/ml) increased the number of Th1 cells not only in Stat5a−/− CD4+ T cells but also in WT CD4+ T cells (Fig. 2). In contrast, the levels of IL-12 production in CpG ODN-stimulated CD11c+ dendritic cells were similar in Stat5a−/− mice and WT mice data (not shown). These results indicate that enhanced IL-12 responsiveness of CD4+ T cells but not the capacity of IL-12 production from APCs is responsible for the increased Th1 cell differentiation in Stat5a−/− CD4+ T cells.

**IL-12-induced Stat4 phosphorylation is enhanced in Stat5a−/− CD4+ T cells**

It is well recognized that Stat4 activation is essential for IL-12-induced Th1 cell differentiation (26). Therefore, we next examined IL-12-induced Stat4 phosphorylation in Stat5a−/− CD4+ T cells. Interestingly, Stat5a−/− CD4+ T cells that were cultured in Th2-polarizing condition exhibited the enhanced IL-12-induced Stat4 phosphorylation as compared with WT CD4+ T cells (n = 4, p < 0.01) (Fig. 3a). In contrast, IL-12-induced Stat4 phosphorylation was not significantly enhanced in Stat5a−/− CD4+ T cells that were cultured in Th1-polarizing condition (Fig. 3a). Enhanced IL-12-induced Stat4 phosphorylation of Stat5a−/− CD4+ T cells in
expression of IL-12R and for another 72 h in Th1 or Th2 condition in the presence of IL-2. The ululated with plate-bound anti-CD3 mAb for 48 h in Th1 or Th2 condition. After CD4 T cells were stimulated with plate-bound anti-CD3 mAb in Th1 or Th2 condition. After CD4 T cells were washed and rested for 8 h in the presence of IL-2. After cells were washed and rested for 8 h in the fresh medium, cells were then stimulated with IL-12 (15 ng/ml) for 20 min, and intracellular staining for the phosphorylated form of Stat4 was performed. Representative anti-phospho-Stat4 staining gated on CD4 T cells (left) and the mean fluorescence intensity (MFI) of anti-phospho-Stat4 staining (right) are shown. Dashed lines are FACS profiles of anti-phospho-Stat4 staining without IL-12 stimulation. Data are mean ± SD from four experiments.

Expression of SOCS3 at mRNA level was examined in WT and Stat5a−/− T cells. Total RNA was prepared from freshly isolated WT CD4+ T cells or Stat5a−/− CD4+ T cells. TaqMan PCR analysis for SOCS3 and GAPDH (as a control) mRNA was performed and the levels of SOCS3 mRNA were normalized to the levels of GAPDH mRNA. Data are means ± SD from four experiments.

Increasing evidence indicates that SOCS family proteins negatively regulate JAK/STAT signaling pathways (28–30). Recently, it has been demonstrated that one of SOCS family proteins, SOCS3, is preferentially expressed in Th2 cells and inhibits IL-12-induced Stat4 phosphorylation (13, 14). To determine whether SOCS3 is involved in the enhanced IL-12 responsiveness in Stat5a−/− CD4+ T cells, we first examined the expression levels of SOCS3 mRNA in Stat5a−/− CD4+ T cells. Consistent with previous reports (13, 14), SOCS3 mRNA was detectable in freshly isolated WT CD4+ T cells by real-time PCR analysis (Fig. 4a) and the expression levels were enhanced when

**FIGURE 3.** IL-12-induced Stat4 phosphorylation is enhanced in Stat5a−/− CD4+ T cells. a, IL-12-induced Stat4 phosphorylation in Stat5a−/− CD4+ T cells. Splenocytes from WT mice or Stat5a−/− mice were stimulated with plate-bound anti-CD3 mAb for 48 h in Th1-polarizing condition (15 ng/ml, in the presence of IL-12) or Th2-polarizing condition (15 ng/ml, in the presence of IL-4 and 15 μg/ml, anti-IFN-γ Ab). Cells were cultured for another 72 h in Th1 or Th2 condition in the presence of IL-2. b, Intracellular staining for the phosphorylated form of Stat4 was performed. Representative anti-phospho-Stat4 staining gated on CD4+ T cells (top) and the mean fluorescence intensity (MFI) of anti-phospho-Stat4 staining (bottom). Data are mean ± SD from four experiments.

**FIGURE 4.** SOCS3 is diminished in Stat5a−/− CD4+ T cells. a, Expression levels of SOCS3 mRNA in freshly isolated CD4+ T cells. Total RNA was prepared from freshly isolated splenic WT CD4+ T cells or Stat5a−/− CD4+ T cells. TaqMan PCR analysis for SOCS3 and GAPDH (as a control) mRNA was performed and the levels of SOCS3 mRNA were normalized to the levels of GAPDH mRNA. Data are means ± SD from four experiments.

**Th2-polarizing condition was confirmed by immunoblotting (Fig. 3b). Yet, IFN-γ-induced Stat1 phosphorylation was similarly observed in WT CD4+ T cells and Stat5a−/− CD4+ T cells in both Th1- and Th2-polarizing conditions (data not shown).** These results suggest that the inhibitory machinery that prevents IL-12/Stat4 signaling in developing Th2 cells is impaired in Stat5a−/− CD4+ T cells. Because IL-12 responsiveness is regulated in part by the expression levels of its receptor (27), we next examined the expression of IL-12Rβ1 and IL-12Rβ2 on Stat5a−/− CD4+ T cells. However, FACS analysis revealed that both IL-12Rβ1 and IL-12Rβ2 were normally expressed in Stat5a−/− CD4+ T cells even in Th2-polarizing condition (Fig. 3c), suggesting that the expression levels of IL-12Rs are not likely to be responsible for the enhanced IL-12-induced Stat4 phosphorylation in developing Th2 cells by the absence of Stat5a.

**SOCS3 expression is decreased in Stat5a−/− CD4+ T cells**

Because IL-12 responsiveness is regulated in part by the expression levels of its receptor (27), we next examined the expression of SOCS3 in Stat5a−/− CD4+ Th cells. SOCS3 is a component of the inhibitory machinery that prevents IL-12-induced Stat4 phosphorylation in developing Th2 cells by the absence of Stat5a.

It has been demonstrated that one of SOCS family proteins, SOCS3, is preferentially expressed in Th2 cells and inhibits IL-12-induced Stat4 phosphorylation (13, 14). To determine whether SOCS3 is involved in the enhanced IL-12 responsiveness in Stat5a−/− CD4+ T cells, we first examined the expression levels of SOCS3 mRNA in Stat5a−/− CD4+ T cells. Consistent with previous reports (13, 14), SOCS3 mRNA was detectable in freshly isolated WT CD4+ T cells by real-time PCR analysis (Fig. 4a) and the expression levels were enhanced when
cells were cultured in Th2-polarizing condition (Fig. 4b). As shown in Fig. 4a, the expression levels of SOCS3 mRNA were significantly decreased in freshly isolated Stat5a-/- CD4+ T cells (n = 4, p < 0.01). The expression of SOCS3 mRNA was up-regulated in Th2-polarizing condition even in Stat5a-/- CD4+ T cells, but the expression levels were still lower than those in WT CD4+ T cells (n = 4, p < 0.01) (Fig. 4b). Decreased expression of SOCS3 of Stat5a-/- CD4+ T cells in Th2-polarizing condition was confirmed at protein levels by intracellular FACS analysis (n = 4, p < 0.01) (Fig. 4c). These results suggest that Stat5a regulates the expression levels of SOCS3 in CD4+ T cells.

Stat5a activates SOCS3 promoter

It has been shown that the SOCS3 promoter contains putative tandem STAT-binding sequences (Fig. 5a) (22). We therefore investigated whether Stat5a activated SOCS3 promoter. Either a constitutively active form of Stat5a (1*6 Stat5a) or of Stat6 (Stat6VT) was expressed in COS7 cells, and SOCS3 WT Luc was determined by a reporter assay (Fig. 5b). The activity of SOCS3 WT Luc was significantly enhanced by the expression of 1*6 Stat5a but not of Stat6VT (Fig. 5b). When one of the putative STAT-binding sequences located in the SOCS3 promoter was mutated (SOCS3 mt1 Luc or SOCS3 mt2 Luc) (Fig. 5a), 1*6 Stat5a-induced activation was largely abolished (Fig. 5b). 1*6 Stat5a-induced activation of the SOCS3 promoter was more severely decreased when both STAT-binding sequences were simultaneously mutated (SOCS3 mt3 Luc) (Fig. 5b). As positive controls, we confirmed that the expression of 1*6 Stat5a preferentially activated a Stat5-dependent reporter construct (MGF-Luc) (21), whereas Stat6VT preferentially activated a Stat6-dependent reporter construct TPU474 (Fig. 5b) (20). These results indicate that Stat5a but not Stat6 activates the SOCS3 promoter.

Stat5a binds to the SOCS3 promoter in CD4+ T cells

To determine whether Stat5a binds to the SOCS3 promoter in CD4+ T cells, we next examined Stat5a binding to the SOCS3 promoter by a ChIP. Activated CD4+ T cells from WT mice or Stat5a-/- mice were stimulated with IL-2 for 30 min, fixed with formaldehyde, and sonicated to reduce the DNA length between 500 and 700 bp. After the sonicated chromatin from these cells were immunoprecipitated with anti-Stat5a Ab, the amount of DNA sequences derived from the SOCS3 promoter (from -214 to +13) or the β-globin promoter (as a control) in the immunoprecipitates was assessed by PCR. As shown in Fig. 6, anti-Stat5a Ab precipitated DNA derived from the SOCS3 promoter but not from the β-globin promoter in IL-2-stimulated WT CD4+ T cells. As anticipated, anti-Stat5a Ab did not precipitate DNA derived from the SOCS3 promoter in Stat5a-/- CD4+ T cells even when stimulated with IL-2 (Fig. 6). These results indicate that Stat5a binds to the SOCS3 promoter in CD4+ T cells.

Enforced expression of SOCS3 restores Th cell differentiation in Stat5a-/- CD4+ T cells

We finally examined the effect of SOCS3 expression on the impaired Th cell differentiation of Stat5a-/- CD4+ T cells. We used bicistronic retrovirus-mediated gene expression system, in which infected cells were identified by coexpressed GFP. Splenocytes from Stat5a-/- mice were stimulated with anti-CD3 mAb and infected with pMX-SOCS3-IRES GFP retrovirus or pMX-IRESGFP retrovirus (as a control) in the presence or absence of IL-4. As a control, splenocytes from WT mice were stimulated with anti-CD3 mAb and infected with pMX-β-globin promoter (as a control) in the immunoprecipitates from Stat5a-/- CD4+ T cells (Fig. 7). Taken together, these results suggest that the diminished SOCS3 expression is involved in the impaired Th1/Th2 balance in Stat5a-/- CD4+ T cells.

FIGURE 5. Stat5a directly activates the SOCS3 promoter. a. The murine SOCS3 promoter (WT) and the mutants (mt1, mt2, and mt3) of putative STAT-binding sequences. b. Stat5a preferentially activates the SOCS3 promoter. COS7 cells were transfected with TPU474, MGF-Luc, SOCS3 WT Luc, SOCS3 mt1 Luc, SOCS3 mt2 Luc, or SOCS3 mt3 Luc in the presence or absence of the expression vectors for the constitutively active form of Stat6 (pcDNA3 Stat6VT) or the constitutively active form of Stat5a (pcDNA3 1*6 Stat5a). Twenty-four hours after transfection, the luciferase activity of TPU474, MGF-Luc, SOCS3 WT Luc, SOCS3 mt1 Luc, SOCS3 mt2 Luc, or SOCS3 mt3 Luc was evaluated by the dual luciferase reporter system. Data are mean ± SD from four experiments.

FIGURE 6. Stat5a binds to the SOCS3 promoter in CD4+ T cells. Splenocytes from WT mice or Stat5a-/- mice were stimulated with plate-bound anti-CD3 mAb for 48 h. CD4+ T cells were purified (>90% pure by flow cytometry) using a CD4+ T cell enrichment column, starved from cytokines in fresh medium for 3 h, and then stimulated with IL-2 (20 ng/ml) for 30 min. Cells were fixed with formaldehyde, lysed, and sonicated to reduce the DNA length between 500 and 700 bp. The sonicated chromatin was immunoprecipitated with anti-Stat5a antisera. After deproteinization and reversal of cross-links, the amounts of DNA sequence for the SOCS3 promoter and the β-globin promoter (as a control) in the immunoprecipitates were assessed by PCR. The input DNA and genomic DNA (as a positive control (PC)) were also subjected to PCR analysis. Shown are representative data from four independent experiments.
We also show that Stat5a regulates the expression of SOCS3 in CD4+ T cells. Increasing evidence has revealed that SOCS family proteins are involved in a negative feedback loop of JAK/STAT signaling (28–30). Among SOCS family proteins, SOCS3 has been shown to be preferentially expressed in Th2-polarized cells and to prevent IL-12-induced Th1 cell differentiation (13, 14). In this study, we found that the expression of SOCS3 was decreased not only in freshly isolated CD4+ T cells but also in Th2-polarized CD4+ T cells in Stat5a−/− mice (Fig. 4). We also found that Stat5a bound to the SOCS3 promoter in CD4+ T cells upon IL-2 stimulation (Fig. 6). In addition, we found that a constitutively active form of Stat5a but not a constitutively active form of Stat6 could activate the SOCS3 promoter in a STAT-binding sequence-dependent fashion (Fig. 5b), which is in agreement with a previous finding that Stat5a preferentially recognizes TTC-N3-GAA STAT-binding sequence, whereas Stat6 preferentially recognizes TTC-N4-GAA STAT-binding sequence (31). Taken together, these results suggest that Stat5a but not Stat6 induces SOCS3 expression in the developing Th2 cells. Moreover, because the retrovirus-mediated expression of SOCS3 restored the altered Th cell differentiation of Stat5a−/− CD4+ T cells (Fig. 7), the reduced expression of SOCS3 is likely to be involved in the disregulated Th1/Th2 balance in Stat5a−/− CD4+ T cells.

Accumulating evidence suggests that the Stat5a-induced SOCS3 expression is also involved in the regulation of Th2 cell-mediated allergic inflammation in vivo. First, we have previously shown that Th2 cell-mediated allergic airway inflammation is decreased in Stat5a−/− mice (10), indicating that Stat5a is involved in the induction of in vivo Th2 cell-mediated immune responses. Second, a recent study has shown that SOCS3 expression is increased in peripheral T cells in asthma patients and that the constitutive expression of SOCS3 within T cells results in the enhanced airway hyperreactivity in a mouse model of asthma (14), suggesting that SOCS3 also plays an important role in the induction of Th2 cell-mediated allergic airway inflammation. Third, in the present study, we show that Stat5a is essential for the appropriate expression of SOCS3 in CD4+ T cells, especially in developing Th2 cells (Figs. 4–6). Therefore, although further studies are required, it is suggested that the Stat5a-mediated SOCS3 induction participates in the induction of Th2 cell-mediated allergic airway inflammation.

Because Stat5 has been shown to up-regulate a number of SOCS family proteins (28) and because it is suggested that, in addition to SOCS3, some of SOCS family proteins may regulate Th1 cell and Th2 cell differentiation (30), it is possible that other SOCS family proteins are also involved in Stat5a-mediated Th cell differentiation. For example, SOCS1, an important negative regulator of IFN-γ signaling (32, 33), has been shown to be induced by Stat5 activation (23). However, we found that IFN-γ-induced Stat1 phosphorylation was not enhanced in Stat5a−/− CD4+ T cells (data not shown), suggesting that SOCS1 may not be involved in Stat5a-mediated suppression of Th1 cell differentiation. The possible involvement of other SOCS family proteins in Stat5a-induced Th cell differentiation needs to be determined in future.

Recently, progress has been made on an upstream cytokine for Stat5a activation during Th cell differentiation. Among a number of cytokines that activate Stat5a, it has been demonstrated that blocking of IL-2, either by the neutralization of IL-2 itself or the blocking of IL-2R, decreases Th2 cell differentiation (12, 34, 35). It has also been shown that the developing Th2 cells express higher levels of the IL-2R σ-chain and exhibit stronger Stat5 activation than the developing Th1 cells (34), consistent with a previous finding that Stat5a functions as an amplifier of IL-2 signaling by inducing the expression of the IL-2R σ-chain (36). Moreover, it has recently been demonstrated that IL-2 but not IL-9, IL-15, or IL-21...
induces Stat5 phosphorylation and IL-4 production in activated CD4+ T cells (37). Therefore, it is suggested that IL-2 is most likely to be a cytokine responsible for Stat5a activation during Th cell differentiation.

It is well recognized that Stat5a regulates the expression of CD25 by directly binding to the 5′ regulatory region of the CD25 gene (38, 39). Consistent with this observation, we have previously shown that the number of CD4+ T cells that express CD25 (CD25−/CD4+ T cells) is decreased in Stat5a−/− mice and we have suggested that the decreased number of CD25+ CD4+ T cells may account for the altered Th cell differentiation of Stat5a−/− CD4+ T cells to some extent (11). In addition, it has been demonstrated that Stat5a directly induces IL-4 production by regulating the accessibility of the IL-4 gene (12). Moreover, we show in this study that the induction of SOCS3 expression by Stat5a in conventional CD4+ T cells is important for Stat5a-mediated Th cell differentiation. Therefore, it is suggested that Stat5a regulates Th cell differentiation in multiple pathways. Further studies are required for the understanding of the relative importance of these pathways in Stat5a-mediated Th cell differentiation.

Because Stat5b is highly homologous to Stat5a (40) and because the mice lacking both Stat5a and Stat5b exhibit a severe defect in T cell responses as compared with Stat5a−/− mice or Stat5b−/− mice (41), it is suggested that the function of Stat5a and Stat5b is somehow overlapped. However, the different phenotypes of Stat5a−/− mice and Stat5b−/− mice underscore the distinctive roles of Stat5a and Stat5b. For example, it has been demonstrated that although Stat5a−/− T cells exhibit no detectable defects in anti-CD3-induced proliferation, Stat5b−/− T cells are defective in anti-CD3-induced proliferation (10, 36, 42), suggesting that Stat5b is likely to play a role in the proliferation and/or survival of activated T cells and that this function of Stat5b may not be shared with Stat5a.

Regarding Th cell differentiation, we have recently found that Stat5a is essential for the residual Th2 cell differentiation in Stat6−/− CD4+ T cells by comparing Stat6−/− mice to Stat5a− and Stat6-deficient double-deficient mice in the same genetic background (44). Because Stat5b is normally expressed and activated in response to IL-2 even in the absence of Stat5a (36, 42), the results suggest that Stat5b cannot compensate the role of Stat5a in Stat6-independent Th2 cell differentiation. In contrast, we have previously shown that in addition to Th2 cell differentiation, Th1 cell differentiation is also decreased in Stat5b−/− mice (10). Because it has recently been shown that Stat5 activates the distal region of the human IFN-γ promoter (43), Stat5b may be involved in the induction of IFN-γ production during Th1 cell differentiation.

In conclusion, we show that Stat5a forces the Th1/Th2 balance toward a Th2-type by preventing IL-12-induced Th1 cell differentiation through the induction of SOCS3. Because it has been demonstrated that SOCS3 regulates the onset and maintenance of Th2 cell-mediated allergic diseases such as asthma and atopic dermatitis (14), it is suggested that Stat5a-mediated SOCS3 induction could be a target for the treatment of Th2 cell-mediated allergic diseases.

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Disclosures
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