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Prostaglandin E2 and SOCS1 have a role in intestinal immune tolerance

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Interleukin 10 (IL-10) and regulatory T cells (Tregs) maintain tolerance to intestinal microorganisms. However, *Il10^{-/-}Rag2^{-/-}* mice, which lack IL-10 and Tregs, remain healthy, suggesting the existence of other mechanisms of tolerance. Here, we identify suppressor of cytokine signalling 1 (SOCS1) as an essential mediator of immune tolerance in the intestine. *Socs1^{-/-}Rag2^{-/-}* mice develop severe colitis, which can be prevented by the reduction of microbiota and the transfer of IL-10-sufficient Tregs. Additionally, we find an essential role for prostaglandin E2 (PGE2) in the maintenance of tolerance within the intestine in the absence of Tregs. *Socs1^{-/-}* dendritic cells are resistant to PGE2-mediated immunosuppression because of dysregulated cytokine signalling. Thus, we propose that SOCS1 and PGE2, potentially interacting together, act as an alternative intestinal tolerance mechanism distinct from IL-10 and Tregs.

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Several mechanisms, including immunoregulation by the innate and adaptive immune systems, allow for millions of bacteria to symbiotically reside in the gastrointestinal tract without disruption of gut homeostasis. Breakdown of tolerance to intestinal microorganisms could have a fundamental role in the pathogenesis of inflammatory bowel disease (IBD). Previous studies of several genetically engineered mouse models have shown that lack of immunosuppression leads to colitis, for example, *Il2*^{-/-} and *Il10*^{-/-}, and dominant-negative transforming growth factor- β (TGF- β) type 2 receptor transgenic (Tg) mice develop spontaneous colitis¹. The major targets of interleukin 2 (IL-2) and TGF- β are T cells: TGF- β suppresses the growth of T cells and their differentiation into Th1 and Th2 cell subsets², and both TGF- β and IL-2 are required for the induction, maintenance or function of Foxp3-expressing regulatory T cells³ (Foxp3⁺Tregs, hereafter referred to as Tregs). IL-10, on the other hand, suppresses the activation of innate immune cells in response to various Toll-like receptor stimuli through a transcription factor signal transducer and activator of transcription 3 (STAT3)⁴.

The Tregs constitute a distinct population of CD4⁺ T cells that have an essential role in the maintenance of systemic tolerance. Alterations in Treg function lead to the activation of T-helper cell subsets known as Th1, Th2 and Th17, which produce the cytokines interferon- γ (IFN γ), IL-4 and IL-17, respectively, as well as to various types of systemic inflammation including inflammation of the gut. However, *Rag1*^{-/-} mice lacking Tregs do not develop colitis spontaneously. Furthermore, *Tgfb1*^{-/-}*Rag2*^{-/-} and *Stat3*^{-/-}*Rag2*^{-/-} mice do not develop colitis despite their lack of Tregs, TGF- β and IL-10 signalling^{5,6}. Therefore, another tolerance mechanism, in addition to TGF- β and IL-10, must exist in the innate immune system.

Many inflammatory cytokines use the janus kinase (JAK)/STAT signalling pathway, which is strictly regulated by negative regulatory mechanisms that limit excessive signalling. Two particularly important players in this negative regulation are the proteins known as suppressor of cytokine signalling 1 (SOCS1) and SOCS3. SOCS1 regulates the signals of various cytokines such as IFN γ , IL-2, IL-4, and IL-7 (ref. 7) *Socs1*^{-/-} mice die as early as 2 to 3 weeks after birth as result of severe inflammation in the liver⁸, though they survived much longer in the absence of IFN γ or in T-cell-deficient conditions.

In this study, we show that *Socs1*^{-/-}*Rag2*^{-/-} mice develop severe spontaneous T-cell-independent colitis. Adoptive transfer of IL-10-sufficient Tregs suppressed the colitis of *Socs1*^{-/-}*Rag2*^{-/-} mice, suggesting that SOCS1 is necessary to maintain the innate immune tolerance in the intestine in the absence of Tregs. IL-10 and IL-10-sufficient Tregs suppressed the production of inflammatory cytokines from lipopolysaccharide (LPS)-stimulated bone marrow-derived dendritic cells (BMDCs), and *Socs1*^{-/-} BMDCs were as sensitive to IL-10-mediated suppression as *Socs1*^{+/+} BMDCs were. Therefore, we hypothesized that there is another immunosuppressive mechanism in the intestinal innate immune system, whose function is impaired in *Socs1*^{-/-} cells, and we sought to identify factors required for this distinct mechanism of immune suppression. Prostaglandin E2 (PGE2) is an IL-10-independent innate immune suppressor, and SOCS1-deficiency abolished its immunosuppressive effect. Furthermore, deprivation of PGE2 by cyclooxygenase (COX) inhibitor provoked severe colitis in *Rag2*^{-/-} but not in *Rag2*^{+/+} mice, which was also prevented by the transfer of Tregs. Thus, we propose that the tolerance in the intestinal innate immune system is regulated by a unique mechanism involving PGE2, the suppressive effect of which could be overridden by inflammatory cytokine signalling, which is strictly regulated by SOCS1.

Results

***Socs1*^{-/-}*Rag2*^{-/-} mice develop spontaneous colitis.** *Socs1*^{-/-} mice die shortly after birth as result of severe systemic inflammation⁸. It has been shown that early lethality can be reverted by crossing with *Ifng*^{-/-} mice, *Tcra*^{-/-} mice or *Rag2*^{-/-} mice, or by restoring SOCS1

expression in T and B cells by crossing with T- and B-cell-specific *Socs1* Tg mice (*Socs1*-KOTg mice)⁸⁻¹¹. Figure 1a shows the course of survival of various *Socs1*^{-/-} mice in our facility. These data suggest that dysregulation of IFN γ signalling and activation of lymphocytes are the main causes of the early lethality in *Socs1*^{-/-} mice. We have previously shown that *Socs1*^{-/-}*Tcra*^{-/-} mice developed more severe colitis than *Tcra*^{-/-} mice¹¹. We suspected that SOCS1 deficiency in T cells might contribute to the development of colitis; however, we later found that *Socs1*-KOTg mice, in which SOCS1 expression is restored in T cells, also developed mild colitis with age¹². Furthermore, we found that *Socs1*^{-/-}*Rag2*^{-/-} mice developed fatal colitis (Fig. 1). These observations suggest that SOCS1 expression in cells other than T cells is important in suppression of colitis.

Socs1^{-/-}*Rag2*^{-/-} mice showed marked growth retardation, but survived for up to 4 or 5 months after birth (Fig. 1a and b). All *Socs1*^{-/-}*Rag2*^{-/-} mice developed severe colitis at 8–9 weeks of age, suffering from diarrhoea and weight loss, whereas littermate *Rag2*^{-/-} mice housed in the same cage never developed colitis (Fig. 1b–d). Through histological examination, we observed many mononuclear and polymorphonuclear cells infiltrating the lamina propria of the large intestines of *Socs1*^{-/-}*Rag2*^{-/-} mice (Fig. 1c). The inflammation extended from the caecum to the rectum, accompanied by the elongation of the crypts, goblet cell depletion and crypt abscesses, all of which are characteristic of chronic colitis.

Flow cytometric analyses showed that CD11b^{high} cells and natural killer cells were predominant in the spleens of both *Rag2*^{-/-} and *Socs1*^{-/-}*Rag2*^{-/-} mice (Fig. 1e). There was an increase in CD11b^{high} cells and Gr-1⁺ cells in *Socs1*^{-/-}*Rag2*^{-/-} mice, probably reflecting granulocytosis caused by active colitis. Immunohistochemical examination showed massive infiltration of CD45⁺ cells in the lamina propria of *Socs1*^{-/-}*Rag2*^{-/-} mice (Fig. 1f). Most of these infiltrating cells were also stained with anti-CD11b or NK1.1, both of which stained significantly more cells in *Socs1*^{-/-}*Rag2*^{-/-} colons than in *Rag2*^{-/-} colons (Fig. 1f). mRNA levels of most of the inflammatory cytokines were upregulated in the colons of *Socs1*^{-/-}*Rag2*^{-/-} mice (Fig. 1g).

To determine the source of these cytokines, we sorted CD11b^{high} NK1.1⁻ and CD11b^{low}NK1.1⁺ cells, and performed quantitative reverse transcription PCR (Fig. 1h). IL-12p35 and IL-6 were produced almost exclusively by the CD11b⁺ cells, whereas tumour necrosis factor- α (TNF α) was produced by both CD11b⁺ and NK1.1⁺ cells (Fig. 1h). The expression levels of these cytokines were upregulated in the *Socs1*^{-/-} group. IFN γ was produced only by NK1.1⁺ cells. NK1.1⁺ cells also expressed small amounts of IL-17A and IL-4. The production of IL-17 from NK1.1⁺ cells has also been reported by Passos *et al.*¹³. Although the expression levels of IFN γ , IL-4 and IL-17A were upregulated in the colons of *Socs1*^{-/-}*Rag2*^{-/-} mice (Fig. 1g), the expression levels of these cytokines per cell in *Socs1*^{-/-} NK1.1⁺ cells were similar to or less than those in *Socs1*^{+/+} NK1.1⁺ cells (Fig. 1h). Therefore, the upregulation of these cytokines in the tissue seemed to be due to the increase in the number of infiltrating cells. Western blot analyses showed that phosphorylation of STAT1 and I κ B α was significantly augmented in the colons of *Socs1*^{-/-}*Rag2*^{-/-} mice, suggesting the activation of JAK/STAT1 and nuclear factor- κ B pathways in these mice (Fig. 1i).

Characterization of the colitis of *Socs1*^{-/-}*Rag2*^{-/-} mice. *Socs1*^{-/-}*Rag2*^{-/-} mice treated with broad-spectrum antibiotics from 6 weeks of age never developed colitis even at 9 weeks of age. Furthermore, colitis was markedly improved by antibiotic treatment even after it had developed (Fig. 2a). These observations suggest that *Socs1*^{-/-} intestinal innate immune cells were highly activated by intestinal commensal bacteria, resulting in the production of inflammatory cytokines, which leads to the development of spontaneous colitis.

Because SOCS1 is a strong negative regulator of the IFN γ signal, we generated *Ifng*^{-/-}*Socs1*^{-/-}*Rag2*^{-/-} mice to examine the contribution of IFN γ to this colitis. As expected, *Ifng* deficiency significantly

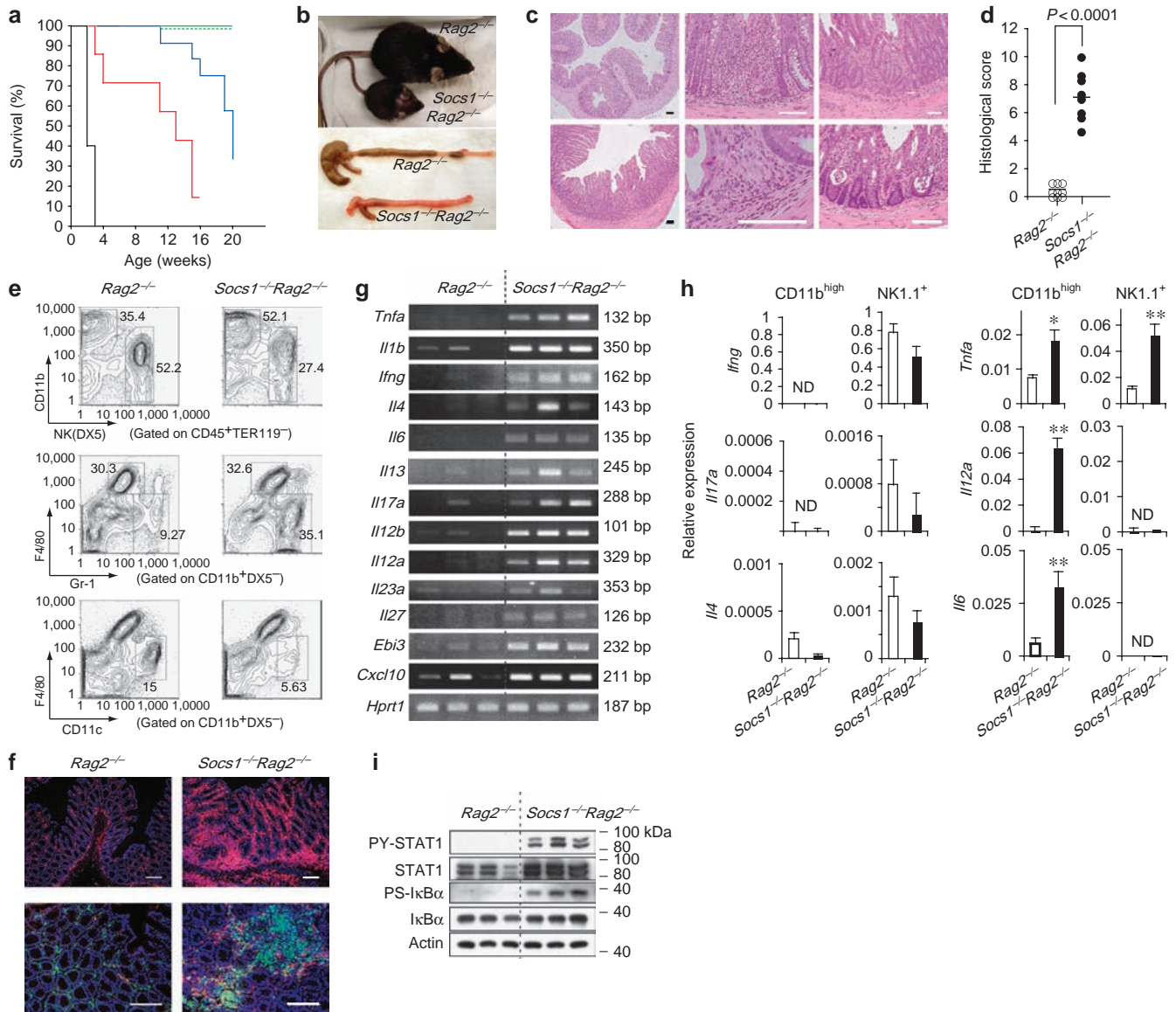


Figure 1 | *Socs1*^{-/-} *Rag2*^{-/-} mice develop spontaneous colitis. (a) The survival curves of *Socs1*^{-/-} (black, n = 5), *Socs1*^{-/-} *Tcrα*^{-/-} (red, n = 14), *Socs1*^{-/-} *Rag2*^{-/-} (blue, n = 12), *Socs1*-KOTg (green, n = 6) and *Ifng*^{-/-} *Socs1*^{-/-} (green, dotted, n = 10) mice. (b) *Rag2*^{-/-} and *Socs1*^{-/-} *Rag2*^{-/-} mice (8-week-old; upper panel). Representative macroscopic observations of the colons of 9-week-old *Rag2*^{-/-} and *Socs1*^{-/-} *Rag2*^{-/-} mice (lower panel). (c) Representative histopathology of the colons of *Rag2*^{-/-} (upper left panel) and *Socs1*^{-/-} *Rag2*^{-/-} mice (other five panels) at 9 weeks of age stained with haematoxylin and eosin. Scale bar, 100 μm. (d) The histological scores of colitis for *Rag2*^{-/-} (n = 9) and *Socs1*^{-/-} *Rag2*^{-/-} (n = 8) mice at 9 weeks of age. The horizontal lines indicate the mean values. (e) Flow cytometric analyses showing the proportions of CD11b^{high} and natural killer (NK) (DX5⁺) cells among CD45⁺ TER119⁻ cells (upper), Gr-1⁺ and F4/80⁺ cells among CD11b⁺ cells (middle) and CD11c⁺ and F4/80⁺ cells among CD11b⁺ cells (lower) in the spleens of *Rag2*^{-/-} and *Socs1*^{-/-} *Rag2*^{-/-} mice. Data are representative of three independent experiments. (f) Fluorescence immunohistochemistry showing the infiltration of CD45⁺ cells (red) into the colonic lamina propria of *Rag2*^{-/-} (upper left panel) and *Socs1*^{-/-} *Rag2*^{-/-} mice (upper right panel). The infiltrating CD45⁺ cells were predominantly CD11b⁺ cells (green), although NK1.1⁺ cells (red) were also significantly increased (lower panels). Nuclei of cells are stained with Hoechst 33342 dye (blue). Data are representative of three independent experiments using samples from different mice. Scale bar, 100 μm. (g) RNAs were extracted from the colons of 9-week-old *Rag2*^{-/-} and *Socs1*^{-/-} *Rag2*^{-/-} mice, and the indicated gene expression was analysed by RT-PCR. Three mice per group are shown. Data are representative of three independent experiments using different samples from one to three mice per group. (h) CD11b^{high} NK1.1⁻ and NK1.1⁺ cells were sorted by FACS Aria from the spleens of *Rag2*^{-/-} and *Socs1*^{-/-} *Rag2*^{-/-} mice, and RNA was purified from the cells. The relative expression levels of the indicated genes were normalized against *Hprt1* using Ct values determined by quantitative real-time RT-PCR. Error bars represent + s.d. of triplicate measurements for each sample. *P < 0.05 and **P < 0.01 compared with *Rag2*^{-/-} cells. Data are representative of two independent experiments. (i) Proteins were extracted from the colons of 9-week-old *Rag2*^{-/-} and *Socs1*^{-/-} *Rag2*^{-/-} mice, and western blot analyses were performed using indicated Abs. Three mice per group are shown. PY, tyrosine phosphorylated; PS, serine phosphorylated. Data are representative of two independent experiments using different samples from two to three mice per group. ND, not determined.

reduced the severity of colitis induced by *Socs1/Rag2* deficiency (Fig. 2b). However, *Ifng* deficiency did not eliminate colitis completely, suggesting that other cytokines that are highly activated in

Socs1^{-/-} *Rag2*^{-/-} mice are also involved in colitis. An obvious candidate was IL-23. IL-23 is regarded as a key inflammatory factor involved in the pathogenesis of colitis¹⁴. The involvement of IL-23

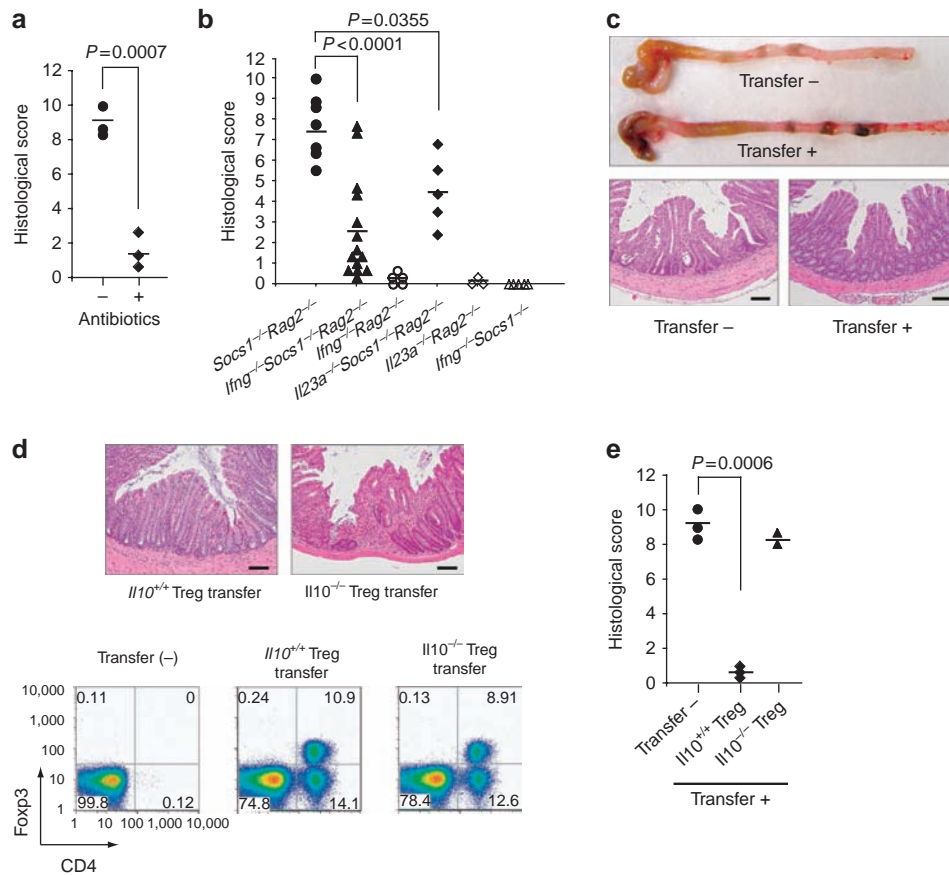


Figure 2 | Characterization of the colitis in *Socs1*^{-/-} *Rag2*^{-/-} mice. (a) Histological scores of the colitis of 16-week-old *Socs1*^{-/-} *Rag2*^{-/-} mice treated with or without antibiotics for 2 weeks from 14 weeks of age ($n=3$). The horizontal lines indicate the mean values. (b) Histological scores of the colitis of *Socs1*^{-/-} *Rag2*^{-/-} mice (black circles, $n=7$), *Ifng*^{-/-} *Socs1*^{-/-} *Rag2*^{-/-} mice (black triangles, $n=14$), *Ifng*^{-/-} *Rag2*^{-/-} mice (white circles, $n=5$), *Il23a*^{-/-} *Socs1*^{-/-} *Rag2*^{-/-} mice (black diamonds, $n=5$), *Il23a*^{-/-} *Rag2*^{-/-} mice (white diamonds, $n=3$) and *Ifng*^{-/-} *Socs1*^{-/-} mice (white triangles, $n=5$) at 9 weeks of age. The horizontal lines indicate the mean values. (c) Macroscopic and microscopic observations of the colons of *Socs1*^{-/-} *Rag2*^{-/-} mice with or without Treg transfer. CD4⁺CD25^{high} T cells (2×10^5 cells) sorted from WT mice were transferred at 10 weeks of age. The mice were killed 6 weeks after the transfer. Representative data are shown. Scale bar, 200 μ m. (d) Representative histopathology of the colons of *Socs1*^{-/-} *Rag2*^{-/-} mice transferred with Tregs sorted from WT (*Il10*^{+/+}) and *Il10*^{-/-} mice. Scale bar, 100 μ m (upper panel). The reconstitution of Tregs confirmed by flow cytometry (lower panel). Spleens were removed from *Socs1*^{-/-} *Rag2*^{-/-} mice without Treg transfer (lower left) or 6 weeks after the transfer with the indicated Tregs (lower middle and right), and the frequencies of CD4⁺ and Foxp3⁺ cells among TER119⁻CD11c⁻ cells gated on particular forward and side scatter were analysed. At 6 weeks after the transfer, nearly 40% of the CD4⁺ cells expressed Foxp3, and the frequency was similar between the *Il10*^{+/+} and *Il10*^{-/-} Treg-transferred group. (e) Histological scores of the colitis of 16-week-old *Socs1*^{-/-} *Rag2*^{-/-} mice without transfer (black circles, $n=3$), transferred with *Il10*^{+/+} Tregs (black diamonds, $n=3$) and transferred with *Il10*^{-/-} Tregs (black triangles, $n=2$) is shown. The horizontal lines indicate the mean values.

in the aggravation of colitis in T-cell-deficient mice has also been reported by previous studies^{15,16}, and the mRNA level of *Il23a* was increased in the colons of *Socs1*^{-/-} *Rag2*^{-/-} mice (Fig. 1g). To test the possibility that IL23 was involved in colitis, we generated *Il23a*^{-/-} *Socs1*^{-/-} *Rag2*^{-/-} mice. Milder colitis was observed in the *Il23a*^{-/-} *Socs1*^{-/-} *Rag2*^{-/-} mice than in the *Socs1*^{-/-} *Rag2*^{-/-} mice (Fig. 2b), which supports the idea that not only IFN γ but also IL-23 is involved in the T-cell-independent development of colitis.

Interestingly, *Ifng*^{-/-} *Socs1*^{-/-} mice never developed colitis, whereas *Ifng*^{-/-} *Socs1*^{-/-} *Rag2*^{-/-} mice did (Fig. 2b). This suggests that the development of colitis caused by SOCS1 deficiency was because of the lack of functional T and B cells. These observations strongly suggest that lymphocytes have a protective role in the suppression of colitis caused by SOCS1 deficiency. Among the lymphocytes, Tregs are the most likely to be involved in this process. To explore this possibility, we transferred CD4⁺CD25^{high} cells from wild-type (WT) mice to *Socs1*^{-/-} *Rag2*^{-/-} mice. As we expected, there was a marked reduction in the severity of colitis in the Treg-transferred group (Fig. 2c). In contrast, when we transferred IL-10-deficient Tregs,

there was no significant improvement, despite the similarity of reconstitution rates (Fig. 2d and e). On the basis of these observations, we concluded that the lack of Treg-derived IL-10 is an important component in the development of colitis in *Socs1*^{-/-} *Rag2*^{-/-} mice.

PGE2 as an immunosuppressor molecule in the colonic mucosa.

Our Treg transfer experiment into *Socs1*^{-/-} *Rag2*^{-/-} mice indicated that suppression by Treg-derived IL-10 could overwhelm the inflammatory effect of SOCS1 deficiency. In accordance with this scenario, IL-10 suppressed the TNF α and IL-12p70 production from *Socs1*^{-/-} BMDCs stimulated with LPS or LPS+IFN γ as efficiently as from *Socs1*^{+/+} BMDCs (Fig. 3a, data for LPS alone shown in Supplementary Fig. S1a). Similarly, Tregs suppressed inflammatory cytokine production from BMDCs in an IL-10-dependent manner and this Treg-mediated suppression was also observed in *Socs1*^{-/-} BMDCs (Fig. 3b). These observations suggested that IL-10/Treg-mediated suppression was not impaired by SOCS1 deficiency; therefore, we hypothesized that another immunosuppressive mechanism was altered in the colons of *Socs1*^{-/-} *Rag2*^{-/-} mice.

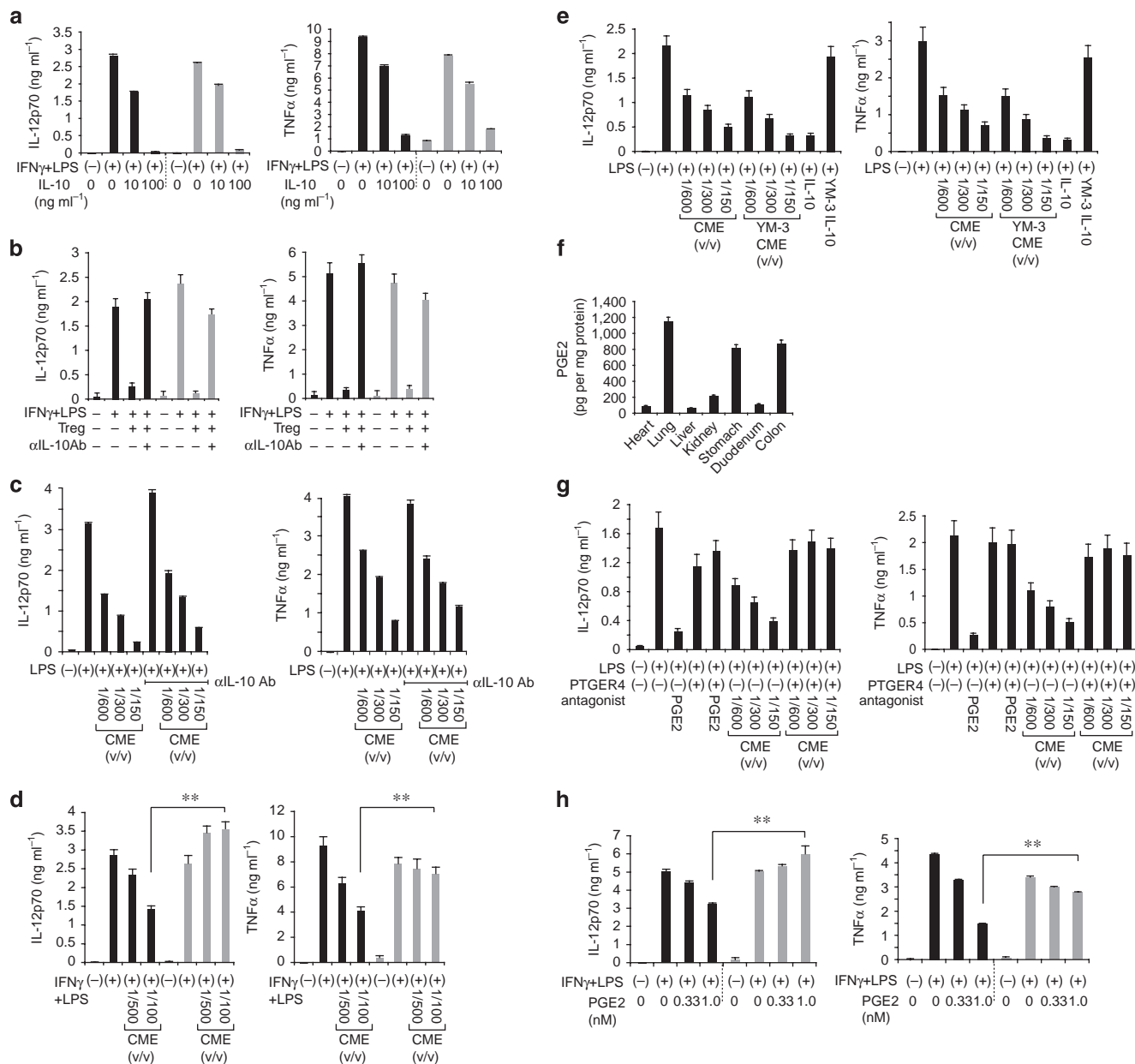


Figure 3 | IFN γ renders *Socs1*^{-/-} BMDCs resistant to PGE2 but not to IL-10. (a–e, g, h) *Socs1*^{+/+} (black bars) and *Socs1*^{-/-} (grey bars) BMDCs (5 × 10⁵ cells per ml) induced from the bone marrow cells of *Rag2*^{-/-} and *Socs1*^{-/-}*Rag2*^{-/-} mice were stimulated for 12 h with indicated reagents, and the levels of IL-12p70 and TNF α in the culture supernatant were measured by ELISA. Error bars represent +s.d. Cells were stimulated in triplicate wells for each condition and s.d. was calculated from the values determined by ELISA. (a) *Socs1*^{+/+} and *Socs1*^{-/-} BMDCs were stimulated with LPS (10 ng ml⁻¹) and murine IFN γ (10 ng ml⁻¹) in the presence of graded concentrations of recombinant murine IL-10. (b) *Socs1*^{+/+} and *Socs1*^{-/-} BMDCs were co-cultured with 2.5 × 10⁵ cells per ml of CD4⁺CD25^{high} T cells in the presence of soluble anti-CD3 Ab (1 μg ml⁻¹). Cells were stimulated with LPS + IFN γ (10 ng ml⁻¹ each) with or without anti-IL-10-neutralizing Ab (αIL-10 Ab, 10 μg ml⁻¹). (c) *Socs1*^{+/+} BMDCs were stimulated with LPS (10 ng ml⁻¹) in the presence or absence of graded concentrations of CME and αIL-10 Ab (10 μg ml⁻¹). (d) *Socs1*^{+/+} and *Socs1*^{-/-} BMDCs were stimulated with LPS + IFN γ (10 ng ml⁻¹ each) in the presence of graded concentrations of CME (v/v, volume of CME/volume of solution (media + CME)). **P < 0.01. (e) *Socs1*^{+/+} BMDCs were stimulated with LPS (10 ng ml⁻¹) in the presence or absence of graded concentrations of CME. The CME filtrated through YM-3 ultrafiltration columns (Millipore), which remove molecules higher than 3,000 Da, retained a suppressive function. IL-10 was used as an experimental control. (f) Tissue extracts from the indicated organs were prepared as described in the Methods, and the levels of PGE2 were determined by ELISA. Error bars indicate +s.d. of triplicate ELISA measurements for each sample. (g) *Socs1*^{+/+} BMDCs were stimulated with LPS (10 ng ml⁻¹) in the presence of PGE2 (1 nM) or graded concentrations of CME with or without PTGER4 antagonist (ONO-AE3-208; 100 nM). (h) *Socs1*^{+/+} and *Socs1*^{-/-} BMDCs were stimulated with LPS + IFN γ (10 ng ml⁻¹ each) in the presence of graded concentrations of PGE2. **P < 0.01. Data are representative of two (f) to three (a–e, g, h) independent experiments.

To define this non-IL-10 tolerance mechanism in the intestine and to specifically identify a soluble factor that could suppress the immune response, we prepared a whole colonic mucosal extract (CME) from the colonic mucosal layer of WT mice and included this CME in the culture media for BMDCs. Interestingly, CME

suppressed LPS-induced production of IL-12p70 and TNF α from BMDCs in a dose-dependent manner (Fig. 3c). Because this suppression was not counteracted by anti-IL-10-neutralizing antibody (Ab), IL-10 could not be involved in the suppression (Fig. 3c). Importantly, the suppressive effect of CME was diminished

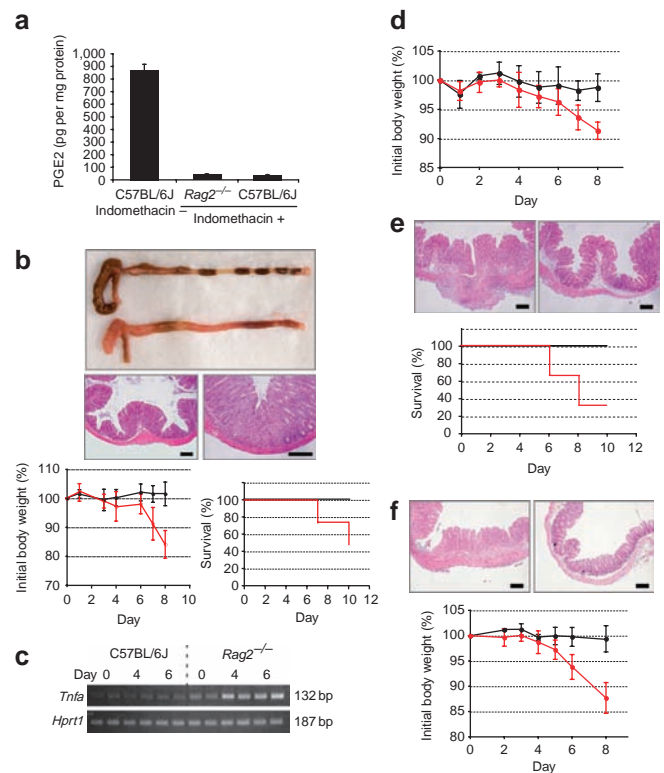
in *Socs1*^{-/-} BMDCs but not in *Socs1*^{+/+} BMDCs when cells were stimulated with LPS + IFN γ (Fig. 3d, data for LPS alone shown in Supplementary Fig. S1b).

In attempting to identify the factor in CME, we noticed that the suppressive factor present in CME was heat resistant (Supplementary Fig. S1c), and its molecular weight was estimated to be <3,000 Da by ultrafiltration (Fig. 3e). Thus, it seemed likely to be a small molecule rather than a protein. We had previously purified a similar factor from mouse embryonic fibroblast culture media and identified it as PGE2 (ref. 13). Therefore, we suspected that the factor in CME was PGE2; in agreement with this hypothesis, the CME contained a large amount of PGE2 (~200 ng ml⁻¹, as determined by enzyme-linked immunosorbent assay (ELISA)). As shown in Figure 3f, extracts from the colon, lung and stomach contained substantial amounts of PGE2, suggesting that these organs have a capacity to produce a relatively large amount of PGE2 compared with other organs under physiological conditions. The suppressive effect of CME as well as PGE2 was completely counteracted by the antagonist of PGE2 receptor 4 (PTGER4, EP4 subtype), a major immunoregulatory receptor for PGE2 (ref. 17) (Fig. 3g). These data suggested that PGE2 is a major suppressor for proinflammatory cytokine production in the intestine, although additional mechanisms of tolerance may also exist. Importantly, just like that of CME, the suppressive effect of PGE2 was diminished in *Socs1*^{-/-} BMDCs when the cells were stimulated with LPS + IFN γ (Fig. 3h). These data indicate that the PGE2-mediated immunosuppressive mechanism was impaired in *Socs1*^{-/-} cells.

PGE2 have an indispensable role in the absence of Tregs. Next, to confirm that PGE2 has a regulatory role in the intestine in the absence of Tregs, we continuously administered the non-selective COX inhibitor indomethacin to WT (*Rag2*^{+/+}) and *Rag2*^{-/-} mice in the drinking water. This treatment effectively suppressed the production of PGE2 in the colons of mice of both genotypes (Fig. 4a). Interestingly, although indomethacin-administered WT mice rarely developed symptoms, *Rag2*^{-/-} mice showed a high susceptibility to indomethacin, and developed diarrhoea and weight loss because of severe inflammation in the caecum and colon (Fig. 4b) accompanied by elevated levels of inflammatory cytokines (Fig. 4c), whereas WT mice developed almost no inflammatory changes in the intestine. When a PTGER4 agonist was administered together with indomethacin, *Rag2*^{-/-} mice did not develop colitis and wasting disease (Fig. 4d), suggesting that the decrease in PGE2/PTGER4 signalling is the direct cause of indomethacin-induced inflammation in *Rag2*^{-/-} mice.

When commensal bacteria were removed by antibiotics, indomethacin did not induce colitis in *Rag2*^{-/-} mice (Fig. 4e), suggesting that a dysregulated immune response to commensal bacteria is involved in this colitis. Furthermore, the transfer of Tregs into *Rag2*^{-/-} mice also prevented colitis of indomethacin-treated *Rag2*^{-/-} mice (Fig. 4f). These findings suggest that PGE2-mediated immunosuppression is indispensable for preventing the hyperactivation of innate immune cells and the resultant intestinal inflammation under Treg-deficient conditions. Interestingly, these features of the colitis of indomethacin-treated *Rag2*^{-/-} mice were almost identical to those of the colitis observed in *Socs1*^{-/-}*Rag2*^{-/-} mice. This was consistent with our *in vitro* observation that the PGE2-mediated immunosuppression was impaired in *Socs1*^{-/-} BMDCs.

SOCS1 maintains the immunosuppressive effect of PGE2. We attempted to ascertain how the PGE2-mediated immunosuppression was impaired in *Socs1*^{-/-} BMDCs. As mentioned in Figure 3h, the suppressive effect of PGE2 on LPS + IFN γ -induced IL-12p70 and TNF α production from BMDCs was counteracted by SOCS1 deficiency, and this antagonism was barely seen in the absence of IFN γ , suggesting that dysregulated IFN γ signal is the cause of the



antagonism. When PGE2 binds to a receptor, especially PTGER4, adenylate cyclase is activated and the level of cyclic AMP (cAMP) is elevated. cAMP then suppresses the production of inflammatory cytokines by several mechanisms^{18,19}. As shown in Figure 5a, *Socs1*^{-/-} BMDCs were also resistant to cAMP-mediated suppression of cytokine production in the presence of LPS + IFN γ . IFN γ

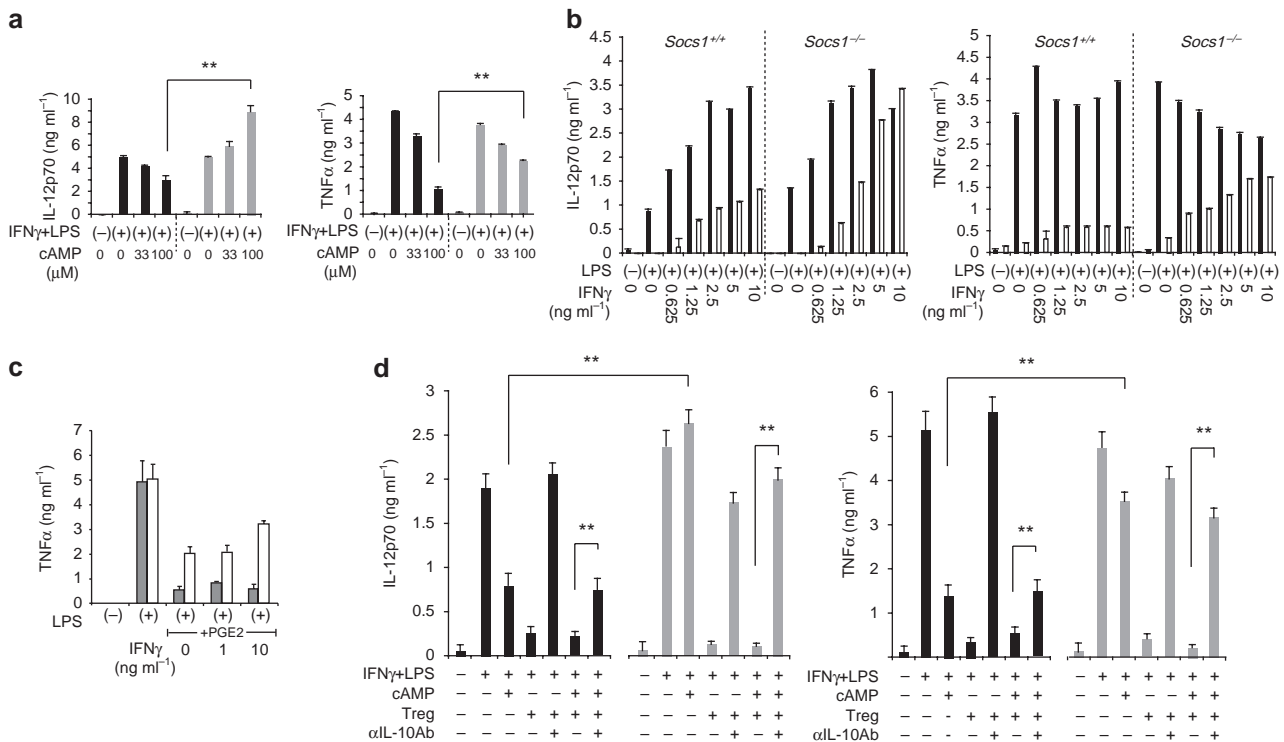


Figure 5 | IFN γ antagonizes cAMP but not Treg-derived IL-10 in *Socs1*^{-/-} BMDCs. (a) *Socs1*^{+/+} (5×10^5 cells per ml; black bars) and *Socs1*^{-/-} BMDCs (5×10^5 cells per ml; grey bars) were stimulated with LPS + IFN γ (10 ng ml^{-1} each) for 12 h in the presence or absence of graded concentrations of membrane-permeable cAMP analogue, 8-Br-cAMP. The levels of IL-12p70 and TNF α in the culture supernatant were measured by ELISA. (b) *Socs1*^{+/+} (5×10^5 cells per ml; left half of each panel) and *Socs1*^{-/-} BMDCs (5×10^5 cells per ml; right half of each panel) were stimulated with LPS (10 ng ml^{-1}) for 12 h with (white bars) or without (black bars) $100 \mu\text{M}$ of 8-Br-cAMP with graded concentrations of murine recombinant IFN γ . The levels of IL-12p70 and TNF α in the culture supernatant were measured by ELISA. IFN γ dose-dependently antagonized cAMP-mediated immunosuppression, which was enhanced by SOCS1 deficiency. (c) *Stat1*^{+/+} (5×10^5 cells per ml; white bars) and *Stat1*^{-/-} (5×10^5 cells per ml; grey bars) BMDCs were stimulated with LPS (10 ng ml^{-1}) for 24 h in the absence or presence of PGE2 and graded concentrations of murine recombinant IFN γ . The level of TNF α in the culture supernatant was measured by ELISA. (d) *Socs1*^{+/+} and *Socs1*^{-/-} BMDCs (5×10^5 cells per ml) were co-cultured with 2.5×10^5 cells per ml of CD4⁺CD25^{high} regulatory T cells in the presence of soluble anti-CD3 Ab ($1 \mu\text{g ml}^{-1}$). Cells were stimulated with 10 ng ml^{-1} of LPS + IFN γ (10 ng ml^{-1} , each) for 12 h with or without anti-IL-10 neutralizing Ab ($10 \mu\text{g ml}^{-1}$) in the presence or absence of membrane-permeable cAMP. The levels of IL-12p70 and TNF α in the culture supernatant were measured by ELISA. Data are representative of two (b–d) to three (a) independent experiments. Error bars represent + s.d. Cells were stimulated in triplicate wells for each condition and s.d. was calculated from the values determined by ELISA. ** $P < 0.01$.

dose-dependently counteracted the suppressive effects of cAMP in *Socs1*^{-/-} BMDCs (Fig. 5b). It is notable that IFN γ partially antagonized the suppressive effect of PGE2 and cAMP in *Socs1*^{+/+} BMDCs (Fig. 5b), although this antagonism was not observed in *Stat1*^{-/-} BMDCs (Fig. 5c). These data suggest that the antagonistic effect of IFN γ /STAT1 pathway against PGE2 and cAMP was amplified in *Socs1*^{-/-} BMDCs.

Lastly, we confirmed the relationship among SOCS1, cAMP and IL-10/Tregs (Fig. 5d). When we co-cultured *Socs1*^{+/+} BMDCs with cAMP and Tregs, the production of IL-12p70 and TNF α was intensively suppressed. The suppressive effect of Treg was diminished by anti-IL-10 Ab, indicating that Treg functioned through IL-10 in this co-culture experiment (Fig. 5d). Although immunosuppressive effect of cAMP was diminished in *Socs1*^{-/-} BMDCs, Tregs potently suppressed cytokine production from *Socs1*^{-/-} BMDCs in an IL-10-dependent manner (Fig. 5d). These phenomena were consistent with our *in vivo* observation that Tregs suppressed colitis in both *Socs1*^{-/-}*Rag2*^{-/-} mice and indomethacin-treated *Rag2*^{-/-} mice. We found that another STAT1-activating cytokine IL-27 and the STAT6-activating cytokines, IL-4 and IL-13, also counteracted the immunosuppressive effect of cAMP in *Socs1*^{-/-} BMDCs, although their effect was not as evident as that of IFN γ (Fig. 6). This may explain why IFN γ deficiency could not completely eliminate the colitis in *Ifng*^{-/-}*Socs1*^{-/-}*Rag2*^{-/-} mice (Fig. 2b). The involvement

of STAT6-activating cytokines, especially IL-13, for pathogenesis of colitis has been shown in previous studies^{20,21}. Although a recent study suggested a role of IL-13 for attenuation of production of IFN γ and IL-17, and the resultant colitis due to IL-10 deficiency²², our study here suggests a synergistic involvement of IFN γ , IL-4 and IL-13 for aggravation of colitis. Collectively, the Treg/IL-10 system seems to be more stable than the PGE2/cAMP system, as the former system was not affected by the strong cytokine signalling, especially IFN γ /STAT1 signalling, resulting from SOCS1 deficiency. The hierarchy among these suppression systems suggested by this study is shown in Figure 7.

Discussion

In this study, we tried to clarify the relationship among Tregs, IL-10, PGE2 and SOCS1 in maintaining the tolerance to commensal microorganisms. Together with a previous report by Berg *et al.*²³, which showed severe indomethacin-induced colitis in *Il10*^{-/-} mice, our study demonstrated the independent contributions of Treg-derived IL-10 and the PGE2/cAMP system for the maintenance of intestinal tolerance. These two systems are regarded as safeguarding each other, thus limiting the aggravation of colitis (Fig. 7). Our study also suggests that inflammatory cytokine signalling would block the immunosuppressive effect of PGE2 and cAMP but not IL-10, and the absence of SOCS1 amplifies the antagonism

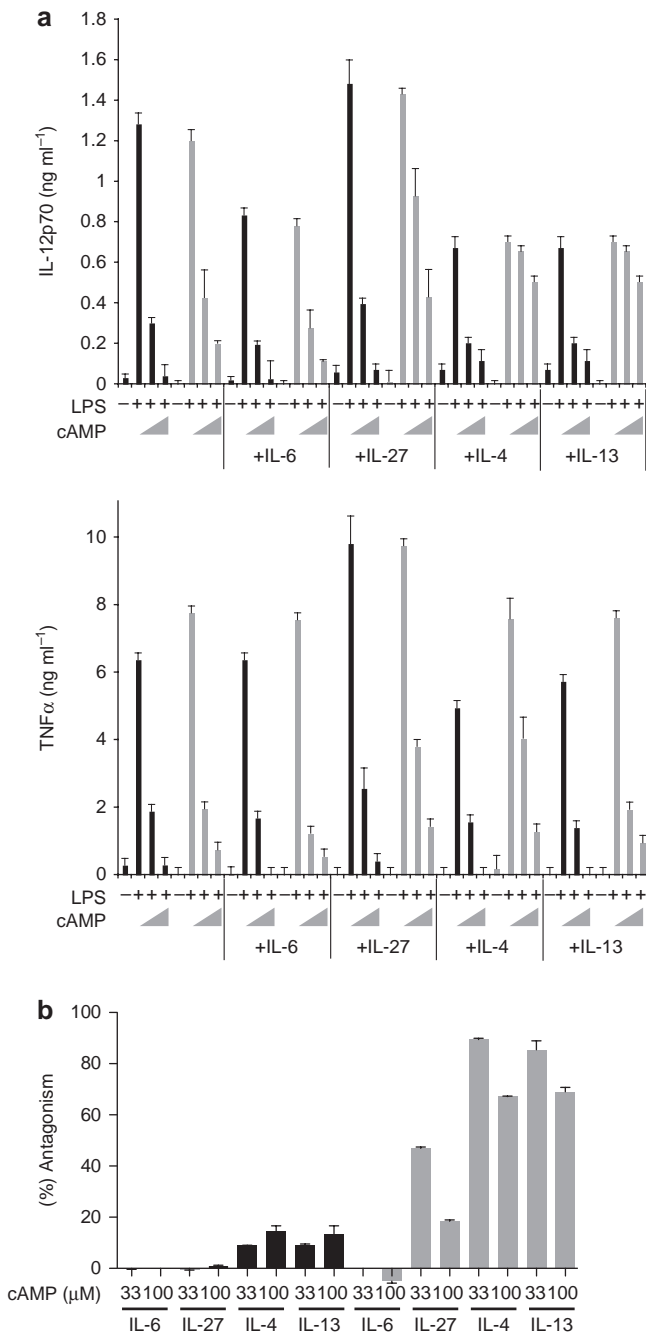


Figure 6 | Select cytokines partially antagonize the cAMP-mediated immunosuppression. *Socs1*^{+/+} (5 × 10⁵ cells per ml; black bars) and *Socs1*^{-/-} (5 × 10⁵ cells per ml; grey bars) BMDCs were stimulated with LPS (10 ng ml⁻¹) and graded concentrations of 8-Br-cAMP (0, 33 and 100 μM) for 12 h with or without recombinant murine IL-6 (20 ng ml⁻¹), IL-27 (100 ng ml⁻¹), IL-4 (20 ng ml⁻¹) and IL-13 (20 ng ml⁻¹). (a) The levels of IL-12p70 and TNFα in the culture supernatant were measured by ELISA. Error bars indicate + s.d. Cells were stimulated in triplicate wells for each condition and s.d. was calculated from the values determined by ELISA. (b) The normalized antagonistic effect (% antagonism) of the indicated cytokines against the suppressive effect of cAMP for the production of IL-12p70. Data were calculated from a data set shown in panel a. Data are representative of two independent experiments (a, b).

to PGE2/cAMP. Although we have shown *ex vivo* that cytokine signalling regulation by SOCS1 is required for efficient PGE2-mediated suppression of activation of innate immune cells, the contribution

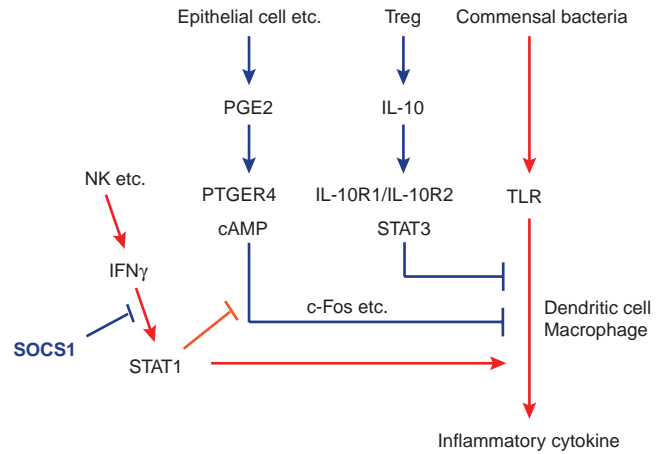


Figure 7 | A model for SOCS1-dependent and -independent mechanism of intestinal tolerance. Tolerance to commensal bacteria is regulated by two independent pathways: PGE2-cAMP and Treg-IL-10. Inflammatory cytokines, especially IFNγ, antagonize PGE2-cAMP-mediated suppression for Toll-like receptor signalling. SOCS1 negatively regulates IFNγ signalling, thus maintaining the PGE2-cAMP-mediated tolerance.

of the resistance to PGE2/cAMP-mediated suppression in *Socs1*^{-/-} cells to the development of colitis of *Socs1*^{-/-} *Rag2*^{-/-} mice is still not clear. Because the molecular mechanisms of PGE2/cAMP-mediated innate immune suppression and STAT-dependent interference to the suppression were not fully resolved, and because we did not have a good molecular target, it was difficult for us to rescue *Socs1*^{-/-} *Rag2*^{-/-} mice from colitis by enhancing the effect of PGE2/cAMP in a manner that could avoid the effect of enhanced STATs signalling. Treatment with a PTGER4 agonist was not suitable for this purpose, because activated STATs would block the effect of the PTGER4 agonist in *Socs1*^{-/-} *Rag2*^{-/-} mice. Therefore, we could not provide direct evidence for the link between the two pathways *in vivo*. Nevertheless, the findings that SOCS1 and PGE2/cAMP system are indispensable for innate immune system to maintain the homeostasis in the colon in the absence of Tregs are interesting. The PGE2/cAMP system has not been well recognized as an important tolerance mechanism *in vivo* probably because this system is vulnerable because of the antagonism by inflammatory cytokines including IFNγ and is also masked by a strong suppressive effect of IL-10. A pivotal role of the PGE2 suppression system appeared only when Tregs or IL-10 are absent. It is well known that the administration of COX inhibitors to human IBD patients may increase the risk of relapse and worsening of the disease²⁰. Similarly, the administration of a COX inhibitor has been shown to exacerbate the course of dextran sulphate sodium salt-induced colitis²¹. The IL-10/Treg system may not be fully functional in these situations.

Although many mechanisms seem to be involved in the gastro-protective effect of PGE2 (ref. 24), suppression of the production of inflammatory cytokines is probably one of the important mechanisms for protection against colitis. The immunosuppressive activity of PGE2 largely depends on the production of cAMP from ATP. Activation of the PGE2 receptors PTGER2 (EP2) and PTGER4 (EP4) usually upregulates intracellular cAMP level, whereas PTGER1 (EP1) upregulates intracellular Ca²⁺ and PTGER3 (EP3) down-regulates cAMP¹⁷; the anti-inflammatory effect of PGE2 is, therefore, context dependent. As shown previously, cAMP suppresses the production of inflammatory cytokines from macrophages and dendritic cells in an IL-10-independent manner^{13,19}. Moreover, cAMP has been shown to be involved in the induction and function of Foxp3⁺ Tregs²⁵⁻²⁷. A recent study has shown that cAMP enhances the induction of human Th17 cells by PGE2/PTGER2/PTGER4 signalling²⁸. Therefore, the effects of PGE2 and cAMP may be complicated

in the presence of T cells. Nevertheless, our study established an essential role of the PGE2/cAMP system to maintain tolerance to commensal bacteria in the intestine.

It is known that not only IL-10 but also TGF- β has an important role in intestinal tolerance and suppression of colitis^{1,28,29}. However, *Tgfb1*^{-/-} mice as well as *Stat3*^{-/-} mice with SCID or *Rag*-deficient backgrounds do not develop colitis^{4-6,30}. We also confirmed that *Il10*^{-/-}*Rag2*^{-/-} mice do not develop colitis during our observation period (2 to 4 months after birth, Supplementary Fig. S2). Although *Il10*^{-/-}*Rag2*^{-/-} mice may still develop colitis with age, it is clear that IL-10 is dispensable for maintenance of intestinal tolerance in young adult lymphopenic mice. Thus, it appears that effector T cells are required for the development of colitis induced by TGF- β or IL-10/STAT3 deficiency. In contrast, *Socs1*^{-/-} mice developed colitis, especially, on a *Rag2*^{-/-} background, and their colitis was suppressed by a transfer of IL-10-producing Tregs. Therefore, the mechanism underlying the colitis induced by SOCS1 deficiency is apparently different from those mechanisms underlying the colitis induced by IL-10 and TGF- β deficiency. The situation of *Socs1*^{-/-} mice was somewhat similar to that of *T-bet*^{-/-} mice³¹. *T-bet*^{-/-} mice also develop colitis only on a *Rag2*^{-/-} background. However, there are also many differences between these two genotypes. In *T-bet*^{-/-}*Rag2*^{-/-} mice, the dysregulation of cytokine production was confined only to TNF α , but in *Socs1*^{-/-}*Rag2*^{-/-} mice, many inflammatory cytokines were upregulated. Rectal prolapse was frequently observed in *T-bet*^{-/-}*Rag2*^{-/-} mice, but was not observed at all in *Socs1*^{-/-}*Rag2*^{-/-} mice. The colitis spread from colitic *T-bet*^{-/-}*Rag2*^{-/-} mice to healthy *T-bet*^{+/+}*Rag2*^{-/-} mice, but not from *Socs1*^{-/-}*Rag2*^{-/-} mice to *Socs1*^{+/+}*Rag2*^{-/-} mice. Because the forms of colitis observed in these two mice are so different, we speculate that T-bet and SOCS1 differentially regulate the intestinal immune response.

We have shown that STAT1 is antagonistic towards the PGE2/cAMP pathway. In addition, STAT6-activating cytokines IL-4 and IL-13 partly counteracted the immunosuppressive effect of cAMP in *Socs1*^{-/-} BMDCs (Fig. 6). However, the molecular mechanisms of PGE2/cAMP-mediated immunosuppression and the antagonisms by STATs remain to be clarified. Recently, we reported that the transcription factor c-Fos is involved in the mechanism of cAMP-mediated immunosuppression in macrophages¹⁹. Another group has recently reported that A kinase-anchoring protein 95 (AKAP95), which interacts with p105 (nuclear factor- κ B1), is responsible for cAMP-mediated suppression³⁰. Further study is necessary to clarify the mechanisms of suppression and the antagonisms by STATs, the balance between which is regulated by SOCS1.

Recent genome-wide association studies demonstrated a significant association of single-nucleotide polymorphisms within the proximity of the *PTGER4* gene region, which regulate *PTGER4* expression, with Crohn's disease³². Polymorphisms in the *STAT3* gene and *JAK2* promoter region also associate with IBD³³⁻³⁶. These studies suggest that alterations in the strength of PGE2 and JAK/STAT signalling can modulate susceptibility to IBD. Our study presented

here also supports this idea, although we should be very careful in translating these results from animal studies to human disease.

We have focused on the role of SOCS1 in the control of intestinal inflammation in this study. The dysregulation of cytokine signalling due to the lack of SOCS1 lead to colitis. Lack of SOCS3, another important negative feedback regulator of cytokine signalling, on the other hand, rendered macrophages and dendritic cells hyporesponsive to various Toll-like receptor stimuli because of the hyperactivation of STAT3 (refs 37, 38), and the macrophage/neutrophil-specific *Socs3*-conditional knockout mice were rather resistant to DSS-induced colitis³⁷. Interestingly, STAT3-hyperactivated innate immune cells are anti-inflammatory in general, whereas STAT3-hyperactivated T cells, including the T cells devoid of *Socs3*, are more prone to produce IL-17 (ref. 39), and thus mediate inflammation in certain conditions. Studies to clarify the cell type-specific roles of SOCS1 and SOCS3 and their relevance in various inflammatory settings *in vivo* are being conducted.

Methods

Mice. *Socs1*^{-/-} mice and *Socs1*-KOTg mice with a C57BL/6J background had been established previously¹². *Il23a*^{-/-} mice have been described previously⁴⁰. *Rag2*^{-/-} mice, *Il10*^{-/-} mice and *Ijng*^{-/-} mice, all with a C57BL/6J background, were obtained from The Jackson Laboratory. Several lines of double knockout and triple knockout mice were generated by intercrossing. All mice were housed in an SPF animal facility at Kyushu University. All experiments using these mice were approved by and performed according to the guidelines of the animal ethics committee of Kyushu University, Fukuoka, Japan.

Flow cytometry and cell sorting. Cells were stained with the indicated fluorescein-conjugated Abs at concentrations of 1:200 to 1:1000, and were analysed on a FACSCalibur or FACSARIA (Becton Dickinson) flow cytometer. All Abs and kits used were purchased from eBioscience. For the isolation of Tregs, CD4⁺ T cells were negatively selected using MACS (Miltenyi Biotec), and CD4⁺CD25^{high}7-AAD⁻ cells were further purified using FACSARIA. The sorted cells, which were used as Tregs, were >95% CD4⁺CD25^{high}, and ~90% of the isolated CD4⁺CD25^{high} cells expressed Foxp3.

Oral indomethacin administration. *Rag2*^{+/+} and *Rag2*^{-/-} mice were continuously fed with 40 μ g ml⁻¹ (corresponding to ~10 mg kg⁻¹ per day) of the non-selective COX inhibitor indomethacin (Nacalai Tesque) in drinking water for 8 to 10 days. To dissolve the indomethacin completely in water, 0.04% dimethyl sulfoxide and 0.2 mM NaOH were also added. The indomethacin solution was replaced every 48 h.

Antibiotic treatment and PTGER4 agonist administration. To remove the intestinal commensal bacteria, imipenem (50 mg kg⁻¹ per day; Banyu Pharmaceutical) and vancomycin (50 mg kg⁻¹ per day; Sigma-Aldrich) were continuously administered in drinking water. Imipenem + vancomycin at this dosage has been reported to effectively reduce both the aerobic and anaerobic bacteria residing in the colon of mice⁴¹. A PTGER4 agonist, ONO-AE1-329 (a generous gift from Ono Pharmaceutical), was injected subcutaneously twice a day at a dosage of 4 μ g kg⁻¹ body weight during the entire period of indomethacin administration.

Histopathological examination. The severity of each colitis case was determined according to the criteria shown in Table 1. In preparation for immunohistochemical examination, frozen cryostat sections were fixed in acetone, blocked with mouse serum and incubated with the fluorescein-conjugated monoclonal Abs (1:100). The slides were examined through a fluorescence digital microscope (Keyence).

Table 1 | Histological scoring system for chronic colitis.

Histological changes	Score			
	0	1	2	3
Loss of goblet cells	None	Mild/focal	Numerous and diffuse	Numerous and diffuse
Crypt abscesses	None	Few (1 in several sections)	A few (1-4 per section)	Many (≥ 5 per section)
Hyperaemia in the mucosa	None/not evident	Mild	Diffuse	Diffuse
Cellular infiltration in the lamina propria	None	A few	Moderate/focal	Numerous and diffuse
Elongation of colonic mucosa	Normal ($\leq 100\%$)	Mild (101-150%)	Evident ($\geq 150\%$)	Evident ($\geq 150\%$)
Epithelial erosion	None	Mild/focal	Evident/diffuse	Evident/diffuse

For the assessment of the severity of colitis, histological scores of the proximal, middle and distal portion of the colons were determined, as indicated, and the average was calculated.

Preparations of CMEs. To extract whole molecules of the substances contained in the colonic mucosa, 0.2 g wet weight of the mucosal and submucosal layers removed from the colons were resuspended in RPMI 1640, frozen at -20°C and then thawed at 4°C . This freezing and thawing cycle was repeated three times. After the third thawing, samples were centrifuged, and the supernatants were filtered through a 0.22- μm filter. The liquid thus obtained was used as CME. For the comparison of PGE2 levels among tissues, several organs were collected from WT mice shortly after the euthanasia, embedded in ice-cold PBS containing 100 μM indomethacin and frozen and thawed twice. Indomethacin was included in the buffer to minimize the *de novo* production of PGE2 during the experimental procedure. After filtration through a 0.22- μm filter, the concentrations of proteins in the supernatants were measured and adjusted to 1 mg ml^{-1} . PGE2 levels of the supernatants were determined using ELISA kit (Cayman Chemical).

Biochemical analysis of bone BMDCs. Granulocyte-macrophage colony stimulating factor-induced BMDCs were stimulated with LPS (10 ng ml^{-1} , *Escherichia coli* serotype 055:B5, Sigma-Aldrich) in the absence or presence of the combinations of cytokines and reagents indicated in the figure legends. The levels of IL-12p70 and TNF α produced were measured with an ELISA kit (eBioscience). 8-Bromoadenosine 3,5'-cyclic monophosphate sodium salt (8-Br-cAMP) was purchased from Sigma-Aldrich. PGE2 was purchased from Nacalai Tesque. Recombinant murine cytokines were purchased from Peprotech. Anti-IL-10 Ab was purchased from eBioscience. A PTGER4 antagonist, ONO-AE3-208 is a gift from Ono Pharmaceutical.

Western blot analysis. Anti-tyrosine-phosphorylated STAT1 and anti-serine-phosphorylated I κ B α monoclonal Abs were purchased from Cell Signaling Technology. Anti-STAT1, anti-I κ B α , anti-c-Fos and anti-p65 Ab were purchased from Santa Cruz Biotechnology. Anti-actin Ab was purchased from Sigma-Aldrich.

Reverse transcription PCR analysis. Total RNAs from cell and tissue specimens were reverse transcribed and analysed by semi-quantitative PCR or quantitative real-time PCR. Quantitative real-time PCR was performed using the Power-SYBR Green Real-time PCR Master Mix (Applied Biosystems) and the ABI 7000 Sequence Detector System (Applied Biosystems). The comparative threshold cycle (Ct) method was used to calculate the relative expression levels of the target genes compared with the internal control (*Hprt1*). The primer sequences are shown in Supplementary Table S1.

Statistical analysis. Statistical analysis was performed using Student's *t*-test. A *P*-value <0.05 was considered to indicate statistical significance.

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Author contributions

T.C. designed and carried out the experiments, analysed data and co-wrote the manuscript with A.Y.; K.K., G.M., R.M., N.I., H.Y., T.S. and R.Y. carried out the experiments with T.C.; K.N. and R.T. supervised the study; and A.Y. designed experiments, analysed data and co-wrote the manuscript with T.C.

Additional information

Supplementary Information accompanies this paper at www.nature.com/naturecommunications

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