IN BRIEF

MACROPHAGES

SOCS2 and SOCS3 in macrophage polarization

M1 macrophages are pro-inflammatory, whereas M2 macrophages are involved in tissue repair. Here, Spence et al. study the role of suppressor of cytokine signalling 2 (SOCS2) and SOCS3 in macrophage polarization. The phenotypes of Socs2^{-/-} or Socs3^{-/-} macrophages were analysed in the absence or presence of polarizing stimuli, and their function was tested in a mouse model of sepsis. Socs2^{-/-} macrophages were biased towards the M1 subset, whereas Socs3-/- macrophages had a stable M2 profile. Moreover, compared with controls, Socs2-/macrophages had higher levels of phosphorylated STAT1 (signal transducer and activator of transcription 1) following stimulation with interferon- γ , and Socs^{3-/-} macrophages had higher levels of active STAT6 in response to interleukin-4. As the blockade of these cytokines partially reversed the polarization bias of Socs2^{-/-} and Socs3^{-/-} macrophages, SOCS2 and SOCS3 may control macrophage polarization by regulating cytokine-STAT signalling.

ORIGINAL RESEARCH PAPER Spence, S. et al. Suppressors of cytokine signaling 2 and 3 diametrically control macrophage polarization. *Immunity* 21 Nov 2012 (doi:10.1016/j.immuni.2012.09.013)

MUCOSAL IMMUNOLOGY

IRF3 maintains gut homeostasis

Here, Negishi *et al.* investigate the role of interferon-regulatory factor 3 (IRF3) in intestinal homeostasis. $Irf3^{-/-}$ mice showed more severe symptoms of dextran sulphate sodium-induced colitis and impaired disease recovery compared with control mice. Thymic stromal lymphopoietin (TSLP) and interleukin-33 (IL-33) have homeostatic functions in the gut, and their expression levels were lower in $Irf3^{-/-}$ mice than in wild-type controls, both in the steady state and during colitis. IRF3-dependent expression of TSLP and IL-33 was induced in response to faecal (possibly microbiota-derived) nucleic acids and involved signalling through one of the adaptor proteins MAVS and STING. Finally, IRF3 cooperated with nuclear factor- κ B to promote *Tslp* transcription. Further molecular links between the intestinal microbiota and IRF3-dependent homeostasis remain to be identified.

ORIGINAL RESEARCH PAPER Negishi, H. et al. Essential contribution of IRF3 to intestinal homeostasis and microbiota-mediated *Tslp* gene induction. *Proc. Natl Acad. Sci. USA* 3 Dec 2012 (doi:10.1073/pnas.1219482110)

GENE REGULATION

STATs control subset-specific enhancer activation

Enhancers are upstream genetic elements that promote transcription, and their activation is marked by several epigenetic changes, including monomethylation of histone H3 lysine 4 and binding of the acetyltransferase p300. Genome-wide analyses of enhancer activation in T helper 1 ($T_{\rm H}$ 1) and $T_{\rm H}$ 2 cells revealed very distinct profiles. Interestingly, analyses in T helper cells deficient for key signal transducer and activation of transcription (STAT) proteins showed that STATs bind to subset-specific enhancers and control the recruitment of p300 and, thereby, enhancer activation. Overexpression of the master transcriptional regulators T-bet and GATA3 in STAT-deficient $T_{\rm H}$ 1 and $T_{\rm H}$ 2 cells, respectively, was not sufficient to restore the profile of enhancer activation. Thus, STATs seem to link cytokine signals to transcriptional regulation.

ORIGINAL RESEARCH PAPER Vahedi, G. et al. STATs shape the active enhancer landscape of T cell populations. *Cell* **151**, 981–993 (2012)