

From the Department of Microbiology, Tumor and Cell Biology  
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# **ROLE OF SOCS PROTEINS DURING MYCOBACTERIAL INFECTIONS**

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*Think outside the (SOCS) box.*



## ABSTRACT

*Mycobacterium tuberculosis* is the world's most successful bacterial killer. During infection, mycobacteria reside inside host cells encapsulated within a granuloma structure in the latent, asymptomatic phase of infection. Only 10% of latently infected develop active, infectious tuberculosis months or years after the initial infection. The mechanisms that determine latency and bacterial control as well as protection during active tuberculosis are still not fully understood. Members of the SOCS protein family are regulators of cytokine signaling via inhibition of JAK-STAT activation and their expression is increased during different kinds of infections. Therefore, the purpose of this thesis was to study the role of SOCS1, SOCS2 and SOCS3 during mycobacterial infections.

We demonstrated that infection with *M. tuberculosis in vitro* and in mice strongly upregulated SOCS1 expression. Interestingly, SOCS1 reduced IFN- $\gamma$  secretion by macrophages in response to IL-12 rather than responses to IFN- $\gamma$  itself. In line, SOCS1-deficient macrophages showed improved growth control of mycobacteria *in vitro*. Furthermore, in a mouse model of infection, we demonstrated that SOCS1 expression by macrophages impaired bacterial clearance before the onset of protective adaptive immune responses. However, SOCS1 did not hamper adaptive immune-controlled bactericidal mechanisms at later time points. At this stage of infection, SOCS1 expression by non-macrophage cells protected mice from severe immunopathology.

Additionally, we showed that SOCS2 expression was induced in an IRF3-dependent manner after infection with *M. bovis* BCG or incubation with LPS *in vitro*. However, SOCS2-deficient and control mice infected with *M. tuberculosis* displayed similar bacterial burdens in the lungs.

In studying the role of SOCS3 in different mouse models, we found that the lack of SOCS3 in either myeloid or T cells dramatically increased susceptibility to *M. tuberculosis* infection. During infection, SOCS3 expression in macrophages and dendritic cells was required to prevent an inhibitory effect of IL-6 on TNF and IL-12 secretion and elevated IFN- $\gamma$  expression by CD4<sup>+</sup> T cells. More detailed studies revealed that the lack of SOCS3 in myeloid cells could be mimicked by mutating the SOCS3 binding site of the gp130 receptor. This indicates that among the receptors, which can be regulated by SOCS3, the control over the IL-6 family gp130 receptor is fundamental for proper immune responses. Surprisingly, mice bearing SOCS3-deficient T cells were not susceptible to BCG infection. Moreover, a proper defense against challenge with *M. tuberculosis* infection was restored if mice deficient for SOCS3 in T cells had been BCG-vaccinated.

In conclusion, we demonstrated a pivotal role of SOCS1 and SOCS3 on the outcome of infection with *M. tuberculosis*. SOCS1 expression allows fast bacterial growth during the early phase of infection and protects from severe inflammation during later stages. SOCS3 expression in myeloid and T cells independently mediates resistance to *M. tuberculosis* infection by modulating T cell functions. Based on the obtained data, we suggest that SOCS3-regulated pathways are promising targets for future therapies as well as vaccination strategies.

## POPULAR SCIENCE SUMMARY

Tuberculosis is a long known infectious disease that continues to cause 1.5 million deaths every year. Although treatment is available, it is long lasting and the numbers of antibiotic-resistant cases are increasing. Moreover, the available vaccine is not efficient in adults. To combat tuberculosis, more knowledge about how the immune system can protect against *M. tuberculosis* is required. Members of the SOCS protein family have important functions in the inhibition of immune signaling pathways. Since proper regulation and balance of immune responses during mycobacterial infections is crucial for infection control, we investigated the role of SOCS1, SOCS2 and SOCS3 during *M. tuberculosis* infection.

We chose mice as the animal model of infection for our studies. In order to investigate the role of the different SOCS molecules during mycobacterial infections, we took advantage of mice lacking these specific proteins and compared their outcome of infection with infected control mice.

Regarding the function of SOCS1 in *M. tuberculosis*-infected mice, we found that SOCS1 impaired early immune activation leading to increased bacterial numbers in the lungs. However, presence of SOCS1 was beneficial for the infected host restricting infection-induced inflammation at later time points.

For SOCS2, we demonstrated that SOCS2 expression was upregulated in mycobacteria-infected macrophages, but SOCS2-deficient mice did not show increased susceptibility to *M. tuberculosis* infection.

Furthermore, we studied the role of SOCS3 in different immune cells, which are known to be important during *M. tuberculosis* infection. We found that the lack of SOCS3 in myeloid cells, which belong to the first line of defense, as well as in T cells, which belong to the adaptive immunity as second line of defense, dramatically increased susceptibility to *M. tuberculosis* infections. Interestingly, if mice deficient for SOCS3 in T cells were vaccinated with the attenuated *M. bovis* strain BCG before *M. tuberculosis* infection, they could control *M. tuberculosis* infection equally well as mice expressing SOCS3.

Summarizing, we illustrated a role for SOCS1 and SOCS3 during mycobacterial infections, whereas the function of SOCS2 was shown to be minor or redundant. SOCS1 improves bacterial growth during the early phase of infection and protects from severe inflammation during later stages. SOCS3 expression in myeloid and T cells is required to mediate resistance to *M. tuberculosis* infection. This makes SOCS3-regulated pathways promising targets for future therapies and suggests that their modulation may improve vaccination as well as the control of mycobacterial infections.

## LIST OF PUBLICATIONS

- I. \***Carow B**, \*Ye X, Gavier-Widén D, Bhujju S, Oehlmann W, Singh M, Sköld M, Ignatowicz L, Yoshimura A, Wigzell H, Rottenberg ME.  
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- II. Hu J, Lou DH, **Carow B**, Winerdal ME, Rottenberg ME, Wikström AC, Norstedt G, and Winqvist O.  
LPS regulates SOCS2 transcription in a Type I interferon dependent autocrine-paracrine loop.  
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- III. **Carow B**, Reuschl AK, Gavier-Widén D, Yoshimura A, Chambers B, Rottenberg ME.  
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## LIST OF ABBREVIATIONS

BMDC	Bone marrow-derived dendritic cells
BMM	Bone marrow-derived macrophages
CpG	Cytidine-phosphatate-guanosin
<i>C. pneumoniae</i>	<i>Chlamydia pneumoniae</i>
DC	Dendritic cell
EAE	Experimental autoimmune encephalitis
GH	Growth hormone
G-CSF	Granulocyte colony-stimulating factor
HIV	Human immunodeficiency virus
IFN	Interferon
IFNar1	Type I IFN receptor
IL	Interleukin
iNOS	Inducible nitric oxide synthase
IRAK	Interleukin-1-associated kinase
IRF	Interferon regulatory factor
JAK	Janus kinase
JNK	c-Jun N-terminal kinase
KIR	Kinase inhibitory region
LCMV	Lymphocytic choriomeningitis virus
LIF	Leukemia inhibitory factor
<i>L. major</i>	<i>Leishmania major</i>
LPS	Lipopolysaccharide
MAL	MyD88-adaptor-like protein
<i>M. bovis</i> BCG	<i>Mycobacterium bovis</i> Bacillus Calmette-Guérin
MSMD	Mendelian susceptibility to mycobacterial diseases
<i>M. tuberculosis</i>	<i>Mycobacterium tuberculosis</i>
MyD88	Myeloid differentiation factor 88
NF- $\kappa$ B	Nuclear factor- $\kappa$ B
NK	Natural killer
NO	Nitric oxide
PIAS	Protein inhibitors of activated STAT
R	Receptor

RAG	Recombination activating gene
SH2	Src homology 2
SHP2	Src homology tyrosine phosphatase 2
siRNA	Small interfering RNA
SOCS	Suppressor of cytokine signaling
STAT	Signal transducer and activator of transcription
TGF- $\beta$	Transforming growth factor- $\beta$
TLR	Toll-like receptor
TNF	Tumor necrosis factor
<i>T. gondii</i>	<i>Toxoplasma gondii</i>
Treg	Regulatory T cell
TRIF	Toll-IL-1 receptor domain-containing adaptor-inducing IFN- $\beta$
WHO	World Health Organization
WT	Wild type

# 1 INTRODUCTION

## 1.1 MYCOBACTERIUM TUBERCULOSIS

*Mycobacterium tuberculosis* is the causative agent of the infectious disease tuberculosis<sup>1</sup>. It is estimated that around one third of the global population is currently infected with *M. tuberculosis*, but the probability of those infected to develop active tuberculosis during their lifetime is only around 10%<sup>2</sup>. Prediction for onset of clinical disease is difficult as critical biomarkers in infected but healthy individuals are not available<sup>3</sup>. However, the risk of developing active tuberculosis increases to 10% per year if a co-infection with the human immunodeficiency virus (HIV) and *M. tuberculosis* occurs<sup>4</sup>, suggesting that immune responses are responsible for the asymptomatic infection. In 2010, the World Health Organization (WHO) reported for tuberculosis an incidence of over 8 million people and 1.2-1.5 million deaths (including deaths from tuberculosis among HIV-positive patients) making tuberculosis the second leading cause of death from an infectious disease worldwide, after HIV<sup>5</sup>. Other immune suppressive conditions such as diabetes, alcoholism, smoking, advanced age and malnutrition are known to promote disease<sup>6,7</sup>. However, a detailed understanding of why reactivation of *M. tuberculosis* occurs in some individuals has yet to be determined.

The treatment regiment for tuberculosis with combined antibiotics is long, but efficient when followed properly. Due to insufficient drug treatment, drug-resistant *M. tuberculosis* strains have developed over the last decades, including strains resistant to isoniazid and rifampicin, the standard treatment antibiotics, as well as extensively drug-resistant strains that additionally resist treatment with one fluoroquinolone or injectable second-line drug<sup>5,8</sup>.

The only available vaccine against tuberculosis is the attenuated *Mycobacterium bovis* Bacille Calmette Guérin (BCG) that was first introduced in 1921. BCG is one of the most used vaccines worldwide, but protection to tuberculosis, especially in adults, varies from 0% to 80%<sup>9-11</sup>. Moreover, BCG vaccination does not provide sterilizing immunity that is needed to prevent the establishment of a latent persistent *M. tuberculosis* infection or active pulmonary tuberculosis. However, over the last years several new tuberculosis vaccines have reached phase I clinical trials<sup>12</sup>.

Mycobacteria are non-motile rods of 2-4 micrometers length. Due to the high lipid content of its cell wall, *M. tuberculosis* cannot be classified as Gram-positive or

Gram-negative, but it can be identified via acid fast staining (Ziehl-Neelsen stain). The cell wall consists of peptidoglycans and complex lipids, where the major lipid components are mycolic acid, cord factor and wax-D. Different compositions of the cell wall can contribute to the virulence of different mycobacteria strains<sup>13</sup>. *M. tuberculosis* is an obligate aerobe and a facultative intracellular pathogen, and it usually infects myeloid cells. *M. tuberculosis* has a slow growing rate with a doubling time of 18 to 24 hours, so that cultures of clinical specimens must be held for 6-8 weeks, which makes diagnostics difficult.

### 1.1.1 Experimental models of *M. tuberculosis*

As *M. tuberculosis* has a very limited host range with no known natural hosts beyond humans<sup>14</sup>, alternative models to study the disease have been tested. Based on the infection of phagocytes by *M. tuberculosis*, cell cultures of primary murine macrophages, human monocyte-derived macrophages and dendritic cells, have been applied to investigate pathogen-host interactions on intracellular and cellular levels<sup>15</sup>.

The first animal model, the guinea pig, was used by Robert Koch when he identified *M. tuberculosis* as the causative agent of tuberculosis in 1882<sup>1</sup>. Guinea pigs show clinical symptoms around 30 days after infection and survive for 100-140 days<sup>16,17</sup>. Importantly, lung responses to mycobacterial stimuli in guinea pigs are similar to human responses, even leading to the development of primary pulmonary granulomas, which closely resemble human granulomas<sup>17-19</sup>. The hallmark of human granulomas, necrotic centers, are found in infected guinea pigs making them a valuable model to test efficacy of tuberculosis vaccines and anti-mycobacterial drugs<sup>20,21</sup>.

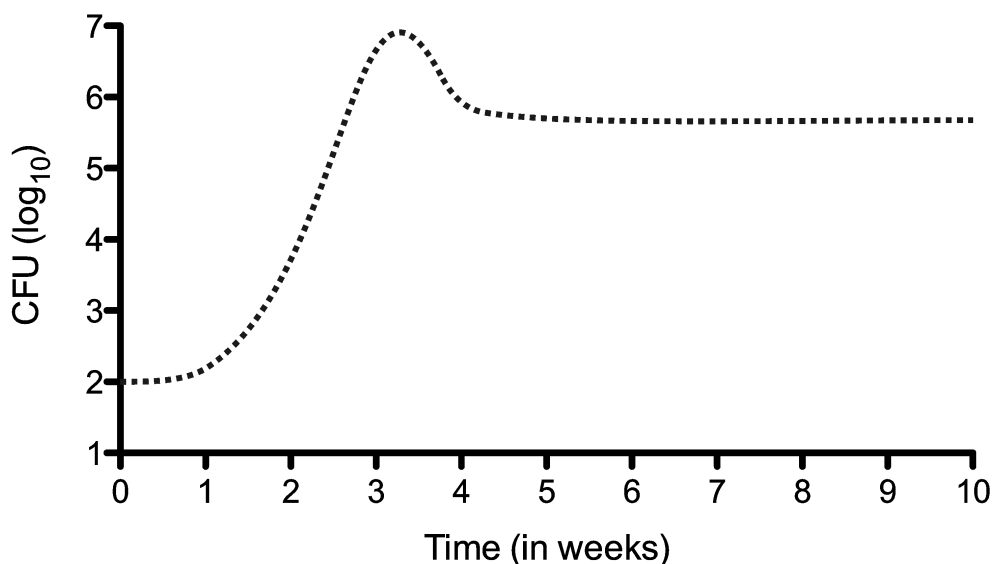
Mice are the most widely used experimental animal model today due to their lower overhead costs as compared to monkeys or guinea pigs, as well as their high reproductive rate, the availability of reagents to characterize mouse responses and established methods for manipulation of gene expression with knockout, conditional knockdown and transgenic mice. The infection of mice is performed either intravenously, intranasally or via aerosol, the latter mimics the natural infection route in humans. A major drawback of the mouse model is the relatively small protective effect of BCG vaccination that reduces bacteria loads after *M. tuberculosis* challenge around 1 log, in comparison to the 2-3 logs reduction seen in guinea pigs<sup>21</sup>. Furthermore, the formed granulomas in infected mice are not highly structured;

consisting of cellular infiltrations with irregular borders, whereas a central necrosis and caseation, as found in human tuberculosis, is absent<sup>22</sup>.

Mycobacteria in aerosol-infected mice replicate progressively in the lungs during the first 3 weeks after infection (Fig. 1)<sup>23</sup>. The onset of adaptive immune responses slows down the bacterial growth, reduces bacteria levels approximately 1 log, and after 4-5 weeks of infection, bacterial levels reach a level that will remain constant throughout the infection (Fig. 1)<sup>24</sup>.

Tuberculosis is typically manifested in the lungs but can also affect other sites, described as extra-pulmonary tuberculosis. In mice, bacterial dissemination to other organs such as the draining lymph node, spleen and liver can be detected after 10 days of aerosol infection<sup>25</sup>. Depending on the initial dose and genetic background of infected mice, they may survive for longer than 200 days, but since all mice develop active disease, they do not provide a model to study latency<sup>26</sup>.

One experimental option to mimic latency in mice is the treatment of *M. tuberculosis*-infected mice with anti-mycobacterial drugs to reduce the bacterial load to undetectable levels followed by either spontaneous or induced reactivation of infection<sup>27,28</sup>. However, it is difficult to evaluate how well this model reflects *M. tuberculosis* latency in humans.



**Figure 1: Kinetic of colony forming units (CFU) of *M. tuberculosis* in lungs of aerosol-infected mice (adapted from<sup>29</sup>).**

Mycobacteria replicate progressively in the lungs during the first 3 weeks of infection, bacterial numbers peak after around 4 weeks and stay at a plateau for more than 10 weeks.

The outcome of infection in cynomolgus macaques displays strong similarities with human disease progression. After a low dose infection some primates generate latent infections, an important stage of *M. tuberculosis* infection that cannot be investigated in the commonly used mouse model<sup>30</sup>.

### 1.1.2 Transmission of *M. tuberculosis*

When infected humans with active pulmonary tuberculosis cough or sneeze, they can spread infectious droplets that are required for transmission of *M. tuberculosis*. Infection occurs if inhaled droplets of bacilli reach the alveoli of the lungs. Exhaled droplets or nuclei are known to remain in the atmosphere for several hours and the infectious dose is low, probably under 10 bacteria<sup>31</sup>.

### 1.1.3 Innate immune responses

The first contact between inhaled mycobacteria and the host is the uptake of *M. tuberculosis* by resident alveolar macrophages within the airways. The ingested mycobacteria are often destroyed directly depending on the mycobactericidal capacities of the host phagocyte and the virulence of the mycobacterium. Pathogenic mycobacteria replicate in the phagosome of the primary host cell and successfully prevent fusion of the phagosome with late endosome/lysosome organelles<sup>32,33</sup>. Moreover, virulent mycobacteria were shown to escape from the lysosomes into the cytosol<sup>34</sup>. Phagocytosis of mycobacteria can be mediated by several receptors such as complement, mannose and scavenger receptors<sup>35-37</sup>. The binding of *M. tuberculosis* molecules to various pattern recognition receptors stimulates the activation of innate immune responses in the host. For *M. tuberculosis*, recognition mainly by toll-like receptor (TLR) 2, 4 and 9 and subsequent activation of MyD88 and TRIF-dependent pathways has been described<sup>38-42</sup>. This innate immune activation leads to up-regulation of co-stimulatory molecules and production of cytokines by infected phagocytes. In addition to macrophages, dendritic cells and neutrophils can also be infected with *M. tuberculosis*<sup>43</sup>. However, innate immune mechanisms are not sufficient to control mycobacterial growth, so that adaptive immune responses need to be activated as a support. Infected dendritic cells migrate to the draining lymph nodes to activate naïve T cells, whereas macrophages recruit immune cells to the site of infection by secretion of attractant chemokines<sup>24</sup>. The following steps are required for a successful protective immune response to *M. tuberculosis*: (1) activation of

antigen-specific effector T cells that migrate back to the lung to activate infected macrophages and (2) the formation of granulomas.

#### 1.1.4 Adaptive immune responses

Antigen-specific CD4<sup>+</sup> and CD8<sup>+</sup> T cells were shown to be required for the control of *M. tuberculosis* infection<sup>44-46</sup>. This is especially apparent in HIV patients, in which a decline of CD4<sup>+</sup> T cells coincides with tuberculosis disease progression<sup>47</sup>. Moreover, deficiency of these cell populations increased susceptibility in murine models of the acute and chronic infection<sup>48,49</sup>. Dissecting the role of T cell subsets in mice demonstrated that the control of *M. tuberculosis* infection was to a greater extent dependent on CD4<sup>+</sup> than on CD8<sup>+</sup> T cells<sup>46</sup>.

IL-12 produced by dendritic cells is crucial for the initiation of the adaptive immune response and stimulates the development of the CD4<sup>+</sup> T cell subpopulation Th1 in response to *M. tuberculosis* infection. The importance of IFN- $\gamma$ -secreting Th1 cells during *M. tuberculosis* infection is well established in mice and humans<sup>50-53</sup>. Moreover, lack of IFN- $\gamma$  as well as the Th1 specific transcription factor Tbet, increases susceptibility to *M. tuberculosis* infection in mice<sup>50,54</sup>.

IFN- $\gamma$  produced by Th1 cells during *M. tuberculosis* infection counteracts the generation of Th17 cells and thereby regulates the balance between Th1 and Th17 cells<sup>55</sup>. Th17 cells are characterized by the secretion of IL-17 and the expression of ROR $\gamma$  transcription factor,<sup>56</sup> and have been shown to be induced during *M. tuberculosis* infection<sup>57</sup>.

Interestingly, the onset of adaptive immune responses after *M. tuberculosis* infection via aerosol is delayed in comparison to other pathogens. In the mouse model, the arrival of antigen-specific T cells into the lung associated with mycobacterial growth restriction, typically occurs 3 to 4 weeks after infection. Studies using transgenic mice that express T cell receptors specific for mycobacterial antigens suggested two time limiting steps: (1) the arrival of infected dendritic cells in the draining lymph node and (2) a threshold of around 1500 mycobacteria in the lymph node that are required for naïve antigen-specific T cells activation<sup>24</sup>. Studies comparing relatively resistant to more susceptible mouse strains showed that bacteria in resistant mouse strains appeared earlier in draining lymph node and could therefore sooner mount an antigen-specific T cell response<sup>25</sup>.

The induction of pathogen-specific regulatory T cells (Tregs) may contribute to the delay of T cell responses during *M. tuberculosis* infection. In humans and mice, regulatory T cells were found to expand and accumulate at the site of infection<sup>58-61</sup>. Regulatory T cells, characterized by expression of transcription factor foxp3 and surface markers CD4 and CD25, are important to down-regulate immune activation to prevent autoimmune diseases<sup>62</sup>, but their role during tuberculosis infection is under debate. On the one hand, transfer of *M. tuberculosis*-specific T regs in mice resulted in increased bacterial burdens in the lungs, whereas others reported an only transient reduction of bacterial loads after depletion of CD25+ T cells<sup>63,64</sup>.

#### 1.1.5 Granuloma formation

The hallmark of tuberculosis and other diseases, the granuloma, typically consists of a core of infected macrophages or epithelioid cells, and giant cells surrounded by foamy macrophages and other mononuclear phagocytes encircled by a mantle of lymphocytes. Mature granulomas are further encapsulated by collagen fibers. The initiation of a stable granuloma during a subclinical stage of infection leads to containment of the mycobacteria and thus is considered to be a host-protective structure<sup>65,66</sup>. This exclusive perspective is questioned by recent findings suggesting granulomas as a niche, in which the mycobacterium can grow and persist<sup>67</sup>. The environment of the granuloma is believed to respond to mycobacteria, and mycobacterial antigens were shown to be required for the granuloma formation<sup>68</sup>. Mycobacteria-infected macrophages undergo necrosis, resulting in a caseous center of the lesion that is characterized by low oxygen, pH and nutrient levels<sup>69</sup>. The replication of mycobacteria is reduced, and bacteria differentiate into a latent stage, but are not eradicated<sup>69,70</sup>. Yet, in less immune-competent or immune-compromised individuals, the fibrous walls lose their integrity, allowing the bacteria to reach the bronchi (transmission of infection) or the blood vessels (dissemination of infection). Tuberculosis can be manifested in almost any organ, but it occurs most frequently in the lungs and lymph nodes, or as generalized lesions resulting from haematogenous dissemination. Cases of genital, skin, meningeal, bone and hepatic tuberculosis are frequent in highly endemic countries<sup>71</sup>.



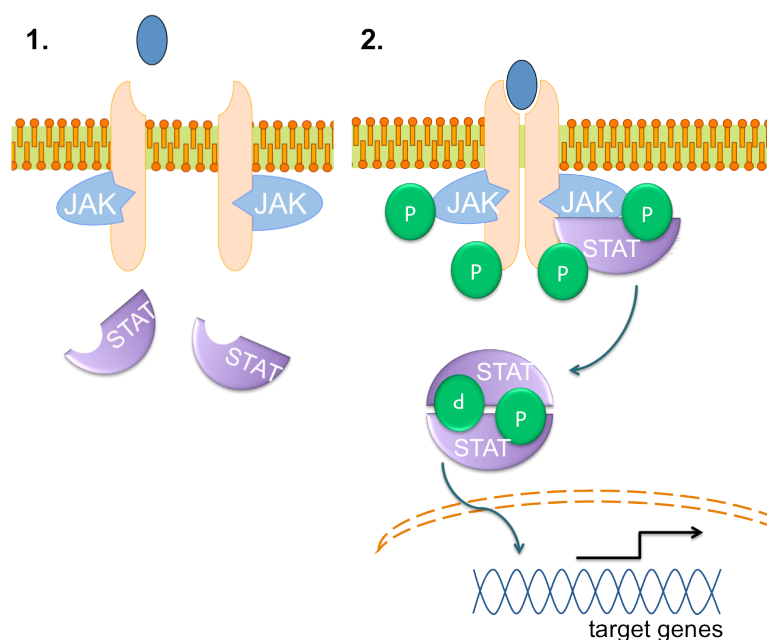
## 1.2 CYTOKINES

Cytokines are usually soluble protein messengers that coordinate the development and function of immune cells, and therefore are indispensable for the initiation, maintenance and termination of immune responses<sup>72</sup>. The group of cytokines includes interleukins and other cell signal molecules, such as tumor necrosis factor and interferons, which are secreted by activated cells to trigger inflammation and respond to infections<sup>73</sup>. Cytokines have a short half-life; acting in either an autocrine or paracrine fashion. Their mode of action is regarded as pro- or anti-inflammatory, depending on the change of activation status of the target cell. The balance of pro- and anti-inflammatory cytokines is an important mechanism to modulate the effect of immune responses: excess of anti-inflammatory cytokines can suppress immune activation, whereas excess of pro-inflammatory cytokines can create harmful immunopathology.

Cytokines exert their function by binding to specific cytokine receptors that can be either membrane bound or soluble, which need to bind to co-receptors on the target cell<sup>73</sup>.

Cytokine receptors can be divided in different groups regarding structural homologies: type I and II cytokine receptors, TNF receptors, IL-1 family receptors, seven transmembrane receptors and G-protein-coupled receptors<sup>73</sup>. Type I and type II cytokine receptors do not possess an intrinsic cytoplasmic kinase activity and therefore are dependent on associated Janus kinases (JAKs) to mediate intracellular signaling after cytokine binding<sup>74</sup>. Binding of cytokines to their type I or II receptor chain leads to the formation of active receptor heterodimers (or cluster of heterodimers) and activation of associated JAKs, which cross-phosphorylate themselves and phosphorylate the cytokine receptor creating binding sites for the SH2 domain of the transcription factors "signal transducers and activators of transcription" (STATs)<sup>74</sup>. Recruited STATs are in turn phosphorylated by the adjacent JAKs and act as binding sites for the SH2 domain of another STAT, which also will be phosphorylated. The phosphorylated STAT dimer is released from the cytokine receptor and translocates into the nucleus where it acts as a transcription activator of specific target genes (Fig. 2). The specificity of the intracellular pathways is due to the presence of 7 different STAT molecules, activated by different receptors<sup>75</sup>. Moreover, STATs can form homo- or heterodimers depending on the activated

receptor, which increases the diversity of target promoters and thereby of gene patterns that can be activated.



**Figure 2: JAK-STAT pathway mediates responses to cytokines.**

Binding of cytokines to their receptors leads to activation of the associated JAKs. Thereafter, STAT molecules get recruited to the receptor and are phosphorylated resulting in activated STAT dimers that act as transcription factors in the nucleus.

A tight regulation of cytokine functions is crucial for both the control of infections and the prevention of infection-associated immunopathology. Different intracellular mechanisms of cytokine signal inhibition are involved in the regulation of innate and adaptive immune responses. The JAK-STAT pathway can be negatively regulated at different stages: protein tyrosine phosphatases remove phosphates from cytokine receptors and activated STATs, SOCS proteins inhibit STAT activation and label cytokine receptors for degradation, whereas PIAS (protein inhibitors of activated STATs) act in the nucleus<sup>76-79</sup>. The role of several of these molecules in the control of infection and disease is not completely understood.

### 1.2.1 Tumor necrosis factor

Tumor necrosis factor (TNF, formerly TNF- $\alpha$ ) is a pro-inflammatory cytokine, mainly produced by activated mononuclear phagocytes, but also by antigen-stimulated T cells and mast cells<sup>80</sup>. TNF is a typical mediator of acute inflammatory responses, sepsis, cachexia and can cause fever but also systemic complications leading to septic shock<sup>81</sup>. During infections, TNF is an important early cytokine that

recruits neutrophils and granulocytes to the site of infection and evokes microbicidal activities in macrophages and neutrophils<sup>80</sup>.

TNF plays a diversified role during *M. tuberculosis* infection contributing to the control of mycobacterial growth as well as to the immunopathology as it participates in granuloma formation. Genetic deletion and neutralization of TNF in mice led to increased bacterial burden and lethality after *M. tuberculosis* infection<sup>82,83</sup>. Furthermore, patients receiving anti-TNF therapy (used for treatment of immune disorders as rheumatoid arthritis) showed an increased rate of reactivation of latent tuberculosis<sup>84</sup>. Similarly in mice, TNF was found to be essential to prevent reactivation in a model of latent *M. tuberculosis* infection<sup>85-87</sup>.

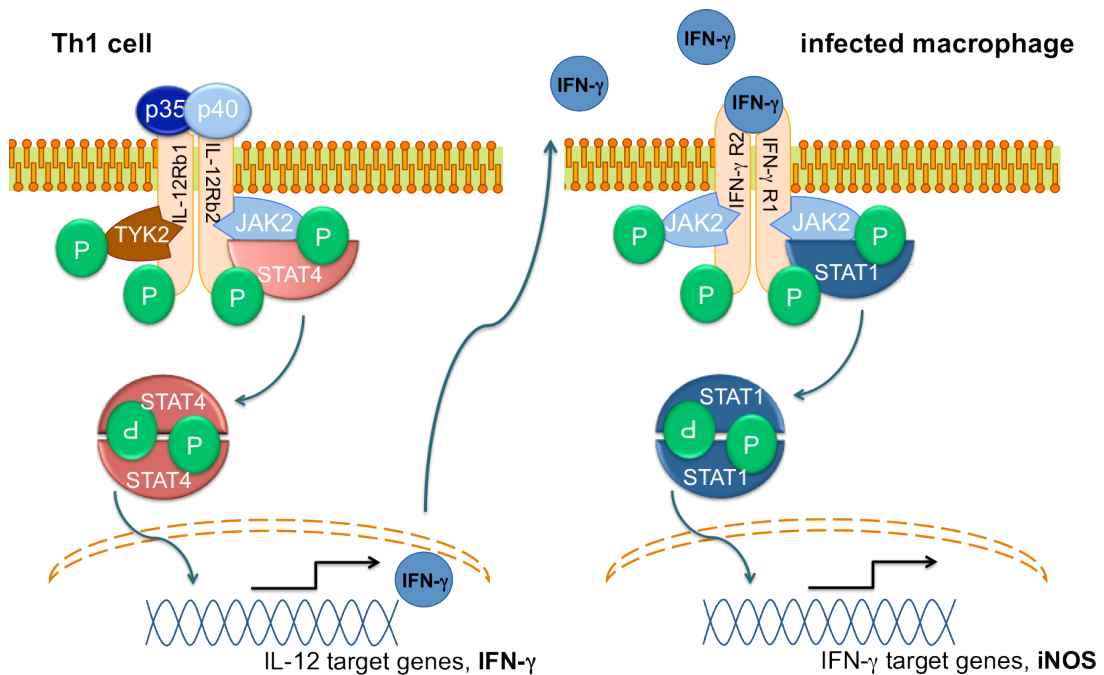
As TNF-deficient mice displayed decreased or disorganized granulomas after mycobacteria infection, part of its protective role could be assigned to its influence on granuloma formation and maintenance<sup>82,86,88-91</sup>. Accordingly, TNF modulates the expression of the chemokines by *M. tuberculosis*-infected macrophages that recruit effector T cells, monocytes and macrophages to the site of infection<sup>90,92,93</sup>. Neutralization of TNF in a chronic infection, when granulomas were already formed, disorganized existing granulomas and aberrant pathology was observed<sup>87</sup>. However, the exact mechanisms for how TNF contributes to the control of mycobacterial growth are poorly understood and recent data in the zebrafish model showed that granuloma formation in *M. marinum* infection may be independent of TNF<sup>94</sup>.

### 1.2.2 Interleukin-12

Interleukin-12 (IL-12) is an important initiator of adaptive immune responses promoting differentiation of naïve CD4<sup>+</sup> T cells to Th1 cells, which produce IFN- $\gamma$  and augment cell-mediated immune responses. Bioactive IL-12 (IL-12p70) is composed of two covalently linked subunits, p35 and p40, that bind to the receptor subunits IL-12R $\beta$ 2 and  $\beta$ 1<sup>95</sup>. Binding of IL-12 to the receptor leads to activation of the Janus kinases JAK2 and TYK2, mediating phosphorylation and activation of the transcription factor STAT4 (Fig. 3)<sup>95</sup>.

The role for IL-12 during *M. tuberculosis* infection was highlighted by the discovery that deficiencies for IL-12p40 and IL-12R $\beta$ 1 can lead to “mendelian susceptibility to mycobacterial disease” (MSMD) due to non virulent mycobacteria<sup>96</sup>. Moreover, mice lacking IL-12p40 or IL-12p35 showed highly increased susceptibility to *M. tuberculosis* infection<sup>97</sup>. In turn, exogenous administration of IL-12p70 increased resistance of BALB/c mice to *M. tuberculosis* infection<sup>98</sup>.

IL-12p40 can be induced in response to *M. tuberculosis* in mononuclear cells by ligation of TLR2 and TLR9 *in vitro* while TLR9 is required for IL-12p40 induction *in vivo*<sup>38</sup>. Early induction of IL-12p40 during *M. tuberculosis* infection is crucial to enable the migration of infected dendritic cells to the draining lymph nodes followed by the activation of naïve T cells<sup>99</sup>. Although IL-12p40 homodimers IL-12(p40)<sub>2</sub> can restore the delayed hypersensitivity response in IL-12p35/IL-12p40-deficient mice, optimal expression of IFN- $\gamma$  by antigen-specific T cells can only be achieved in the presence of IL-12p70<sup>97,100</sup>. Moreover, IL-12p70 is needed to maintain Th1 effector functions during *M. tuberculosis* infection, since exogenous IL-12p70 can reconstitute protection in *p40*<sup>-/-</sup> mice that is again lost after IL-12p70 removal<sup>101,102</sup>. This observation could be confirmed in humans with IL-12R $\beta$ 1-deficiency that fail to maintain a Th1 effector memory population showing that IL-12 plays indispensable roles not only in initiation but also maintenance of protection against mycobacteria<sup>102</sup>.



**Figure 3: Th1 cells activate infected macrophages.**

IL-12 induces IFN- $\gamma$  production in Th1 cells via STAT4 activation that in turn activates infected macrophages. These macrophages induce the transcription of IFN- $\gamma$ -responsive genes as *iNos*.

### 1.2.3 Interferon- $\gamma$

Interferon- $\gamma$  (IFN- $\gamma$ ) is critical during intracellular bacterial and protozoal infections as well as tumor growth control, but aberrant IFN- $\gamma$  production can induce autoimmune diseases<sup>103-107</sup>. Responses to IFN- $\gamma$  are mediated through its interaction with the IFN- $\gamma$  receptor that initiates the activation of STAT1 (Fig. 3)<sup>108</sup>.

The importance of IFN- $\gamma$  for the defense against *M. tuberculosis* has been demonstrated in experimental studies using knockout mice and was confirmed in patients with defects in signaling or production of IFN- $\gamma$  as both show dramatically increased susceptibility to mycobacterial infections<sup>50,109-112</sup>. As previously discussed, CD4+ T cells play an important role during *M. tuberculosis* infection and are considered to be the main source of IFN- $\gamma$ , whereas IFN- $\gamma$  secreted by natural killer (NK) cells appears to be redundant<sup>113,114</sup>.

Early appearance of antigen-specific IFN- $\gamma$ -producing CD4+ T cells in the lungs coincided with earlier restriction of growth after aerosol *M. tuberculosis* infection<sup>115</sup>. Furthermore, a delay in priming of effector T cells led to increased bacterial burden in the lungs<sup>63</sup>.

T cells elicit the bactericidal effects of infected macrophages. IFN- $\gamma$ -activated macrophages were found to overcome bacteria-induced blockage of phagosome maturation and increased presentation of mycobacterial peptides to T cells and upregulated nitric oxide synthase 2 (iNOS2), an enzyme that produces nitric oxide (NO) and is involved in bacterial killing<sup>116,117</sup>. The role of NO in the control of mycobacterial intracellular growth is well established in the mouse model where *Nos2*<sup>-/-</sup> mice displayed increased susceptibility to *M. tuberculosis* infection. In humans, iNOS and NO production has been reported in alveolar macrophages in response to infection<sup>106,118-122</sup>. However, the role of NO in protection during human infection is controversial.

In the addition to the effect of IFN- $\gamma$  on macrophages, it also was found in mouse radiation chimeras that responsiveness to IFN- $\gamma$  by non-hematopoietic cells is involved in the protective immunity to *M. tuberculosis*<sup>55</sup>.

Although the relevance of IFN- $\gamma$ -producing T cells in protection is not up for debate, the numbers of IFN- $\gamma$ -secreting T cells after vaccination do not always correlate with increased protection against *M. tuberculosis* infection<sup>123,124</sup>. This

suggests that more factors than IFN- $\gamma$  production by CD4<sup>+</sup> T cells should be assessed to determine protective immune responses to *M. tuberculosis* infection<sup>49</sup>.

#### 1.2.4 Interleukin-6

IL-6 is involved in innate and adaptive immunity and is produced by mononuclear phagocytes, endothelial cells, fibroblasts, adipocytes and muscle cells<sup>73</sup>. It is induced in response to pathogens, TNF and IL-1. IL-6 signals are transmitted via a receptor complex consisting of IL-6R $\alpha$  and gp130, the common IL-6 family co-receptor. Binding of IL-6 to the receptor complex initiates activation of JAK1 followed by recruitment of several signaling molecule such as STAT3 and SHP2. STAT3, once phosphorylated, dimerizes and acts as a transcription factor in the nucleus, and SHP2 links IL-6 to the Ras-MAP kinase pathway. Depending on the target cell, IL-6 stimulates secretion of acute phase proteins by hepatocytes, neutrophil production in the bone marrow, growth of B lymphocytes and generation of Th17 cells<sup>73</sup>.

Historically, IL-6 has been regarded as a pro-inflammatory cytokine, but recently anti-inflammatory functions of IL-6 have also been described, which will be discussed in detail later<sup>125,126</sup>.

*M. tuberculosis*-infected human and murine macrophages secrete IL-6 and elevated IL-6 serum levels are found in tuberculosis patients<sup>127-129</sup>. However, the role of IL-6 during infection with *M. tuberculosis* is not well understood.

IL-6-deficient mice displayed increased susceptibility to *M. tuberculosis* infection compared to control mice, but only after a systemic high dose infection<sup>130</sup>. In the low-dose aerosol model, which mimics the natural route of infection, *Il-6*<sup>-/-</sup> mice only initially showed elevated bacterial levels<sup>131</sup>. In contrast, *in vitro* studies have shown that high levels of IL-6 impair T cell proliferation of antigen-specific T cells, reduce the responsiveness to IFN- $\gamma$  of infected macrophages and counteract TNF-mediated mycobactericidal activities in macrophages<sup>132-134</sup>.

#### 1.2.5 Interleukin-17

IL-17 can be produced by  $\gamma\delta$  T cells and Th17 cells and has been shown to be involved in the attraction of neutrophils to the site of infection<sup>135-138</sup>. The

development of Th17 cells depends on STAT3 activation (by IL-6, IL-21, IL-23) and TGF- $\beta$  and is counter-regulated by Th1 cells<sup>139,140</sup>.

In mice, IL-17 is mainly secreted by  $\gamma\delta$  T cells during *M. tuberculosis* and early BCG infection<sup>141,142</sup>. Additionally, numbers of IL-17 producing  $\gamma\delta$  T cells were elevated in tuberculosis patients compared to healthy donors<sup>142,143</sup>. Although the lack of IL-17 or IL-17R does not impair the ability of mice to control a low dose aerosol *M. tuberculosis* infection, deficient mice were unable to control *M. tuberculosis* after a high dose intratracheal infection<sup>144,145</sup>. Thus, the relevance of IL-17 for the control of *M. tuberculosis* infection needs to be further investigated.

However, the impact of IL-17 on granuloma formation and pathology have been suggested since BCG-infection of *Il-17a*<sup>-/-</sup> mice showed impaired granuloma maturation<sup>146,147</sup>. On the other hand, repeated BCG-immunizations of *M. tuberculosis*-infected mice led to increased infiltration of neutrophils and tissue damage in the lungs associated to elevated IL-17 levels<sup>146</sup>.

Interestingly, high numbers of neutrophils have been associated with susceptibility to *M. tuberculosis* infection in mice<sup>55,146-148</sup>. In *Ifn- $\gamma$* <sup>-/-</sup> mice infiltrating neutrophils could be related to high levels of IL-17<sup>55</sup>. Another hint for a detrimental effect of high IL-17 levels in *M. tuberculosis* infection was found in IL-27R deficient mice. These mice showed lower bacterial burdens compared to control mice but developed enlarged granulomas and succumbed earlier to infection most likely due to enhanced IL-17 responses<sup>148</sup>.

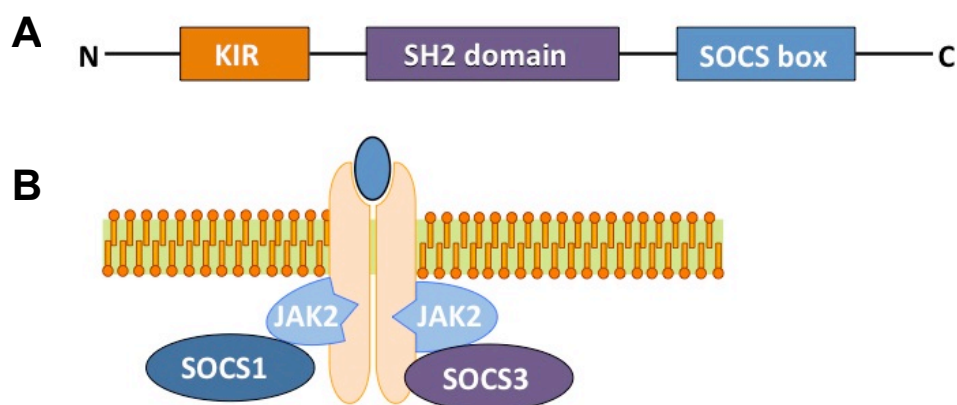
### 1.3 SOCS PROTEINS

The family of suppressor of cytokine signaling (SOCS) proteins consists of 8 members (cytokine inducible SH2 protein, CIS, SOCS1-7), and all are main regulators of intracellular signal transduction mediated through the JAK-STAT pathway. All SOCS proteins share a modular organization with a C-terminal SOCS box, a central SH2 domain and an amino-terminal domain of variable length (Fig. 4). The SOCS-box was shown to interact with additional proteins to form an E3 ubiquitin ligase complex, initiating ubiquitination of target proteins, followed by proteosomal degradation of bound signaling complexes as JAKs and cytokine receptors<sup>149-151</sup>. The SH2 domain of SOCS proteins determines the specificity for their receptors<sup>152,153</sup>.

Both deletion and overexpression of SOCS proteins in animal models provided insights into their importance in regulating the responsiveness to cytokines, growth

factors and hormones but also showed the specificity of SOCS proteins for certain receptors<sup>154</sup>. In particular, SOCS1 and SOCS3 were found to balance immune functions by influencing sensitivity to cytokines, modulating the differentiation of immune cell populations as well as their activation.

To date, SOCS1 and SOCS3 are the most studied members of the SOCS family. They both contain an additional N-terminal kinase inhibitory region (KIR) that is absent in other SOCS proteins. The KIR of SOCS1 and SOCS3 can directly inhibit JAK tyrosine kinase activity by acting as a pseudo-substrate, and thereby blocks the interaction of JAK with their substrate STAT molecules. Although SOCS1 preferably binds directly to the JAK activation loop, the SH2 domain of SOCS3 binds to the cytokine receptor. Using this mechanism, JAK inhibition by SOCS1 and SOCS3 takes place even in the absence of the SOCS box<sup>155,156</sup>.



**Figure 4: SOCS proteins.** A: Schematic structure of SOCS1 and SOCS3 proteins is shown. B: SOCS1 and SOCS3 binding to JAK or cytokine receptor.

## 1.4 SOCS1

SOCS1 expression can rapidly be induced by many cytokines, especially IFNs, and serves as a classical feedback loop inhibiting its inducing pathways<sup>157,158</sup>. Importantly, *Socs1* mRNA expression increases even in response to microbial molecules such as LPS, Pam<sub>3</sub>Cys and CpG oligonucleotides that signal via TLR<sup>159-162</sup>. Furthermore, hormones like insulin<sup>163</sup>, cardiotrophin<sup>164</sup> or glucocorticoids<sup>165</sup> have been shown to stimulate SOCS1 expression.

SOCS1 can bind to the catalytic domain of JAK2 and to TYK2, another molecule of the JAK family, which mediates IFN- $\alpha/\beta$  signaling. SOCS1 has also been shown to bind directly to type I IFN receptors (IFN $\alpha$ 1)<sup>166</sup> and to the IFN- $\gamma$  receptor, efficiently inhibiting STAT1-mediated signaling<sup>154,166,167</sup>.



#### 1.4.1 SOCS1<sup>-/-</sup> mice

Mice deficient for SOCS1 die within 3 weeks after birth due to fatty degeneration and necrosis of the liver<sup>168,169</sup>. Additionally, these mice show retarded growth, lymphopenia and multi-organ haematopoietic infiltrates<sup>168,169</sup>. This inflammatory disease could be related to hyper-responsiveness of *Socs1*<sup>-/-</sup> mice to IFN- $\gamma$ , because mice with an additional knockout of IFN- $\gamma$  or STAT1 survive until adulthood<sup>170,171</sup>. Moreover, *Socs1*<sup>-/-</sup> mice that were treated with neutralizing IFN- $\gamma$  antibodies from birth reached adulthood but died of myocarditis<sup>172</sup>. Indications for an involvement of T cells and NK T cells as a probable source of enhanced levels of IFN- $\gamma$  secretion were found in T and NK T cell-deficient *Rag2*<sup>-/-</sup>/*Socs1*<sup>-/-</sup> mice that are healthy to at least 3 month of age, and in the finding that *Socs1*<sup>-/-</sup> mice transgenic for SOCS1 in T cells survive<sup>171</sup>. However, conditional deletion of SOCS1 in T and NKT cells or in macrophages is not sufficient for the hyper-inflammation<sup>173</sup>. Requirement of non-hematopoietic cells for detrimental inflammation could be excluded, as chimeric mice, which received *Socs1*<sup>-/-</sup> bone marrow after radiation, showed decreased survival<sup>174</sup>. Altogether, lethality caused by SOCS1-deficiency is mainly due to an exacerbated secretion of IFN- $\gamma$  by T and NKT cells during the neonatal period and to an increased sensitivity of myeloid cells to IFN- $\gamma$ .

In contrast, *Socs1*<sup>-/-</sup> mice crossed to *IFNar1*<sup>-/-</sup> mice survived beyond weaning, indicating that even type I IFN contributes to the lethal inflammation<sup>166</sup>. Besides the rescue of SOCS1-deficient mice by IFN depletion, *Socs1*<sup>-/-</sup> mouse strains crossed to *Stat1*<sup>-/-</sup>, *Stat4*<sup>-/-</sup> or *Stat6*<sup>-/-</sup> mice are partially protected from neonatal lethality, but nevertheless displayed chronic inflammation with aberrant T cell activation<sup>175-177</sup>. STAT6 and STAT4 are two STAT family members that specifically mediate signals that emanate from the IL-4/ IL-13 and IL-12 receptors, showing that SOCS1 also affects IL-4 and IL-12 signaling. Uncontrolled responses to IL-12, which is an important activator of IFN- $\gamma$  secretion by T cells, could contribute to increased IFN- $\gamma$  levels and therefore to the development of inflammation in *Socs1*<sup>-/-</sup> mice<sup>175</sup>.

### 1.4.2 SOCS1 in immunity

*Socs1*<sup>-/-</sup> mice are hypersensitive to LPS-induced endotoxic shock, associated with increased levels of IL-12 and TNF<sup>178,179</sup>. Surprisingly, even though *Ifn-γ*<sup>-/-</sup> mice are highly resistant to endotoxic shock<sup>180</sup>, additional deletion of IFN-γ or STAT1 does not rescue *Socs1*<sup>-/-</sup> mice from lethal LPS injection, demonstrating that SOCS1 even attenuates IFN-γ-independent mechanisms<sup>178-180</sup>. Furthermore, elevated sensitivity to endotoxin shock was observed in *Socs1*<sup>fl/fl</sup> *LysM-cre* mice (see description of mouse strain in material and methods) lacking SOCS1 specifically in macrophages and neutrophils<sup>181</sup>.

Increased sensitivity to LPS suggests an interaction of SOCS1 with components of the TLR signaling. In fact, binding of SOCS1 to IRAK1 and the p65 subunit of NF-κB has been shown to destabilize and limit NF-κB activation<sup>178,179,182,183</sup>. SOCS1 can also bind to the apoptosis-regulating kinase 1 and regulates mitogen-activated protein kinases JNK and p38<sup>184</sup>. Moreover, SOCS1 was shown to mediate the degradation of the adaptor protein MAL that is involved in TLR2 and TLR4 signaling<sup>185</sup>. However, results from other investigators could not confirm a direct effect of SOCS1 on TLR signaling<sup>186,187</sup>. SOCS1 overexpression did not affect TLR signaling; instead the inhibition of IFN-α/β-mediated STAT1 activation by SOCS1 may account for the observed sensitivity to LPS in *Ifn-γ*<sup>-/-</sup>/*Socs1*<sup>-/-</sup> mice<sup>186,187</sup>. Thus, further work is required to clarify whether SOCS1 is directly involved in the regulation of TLR signaling.

### 1.4.3 SOCS1 in infectious diseases

A wide range of pathogens including parasites, bacteria and viruses are potent stimulators of SOCS1 expression. Despite obvious differences in the type of protective or deleterious immune response elicited by different pathogens, for most intracellular infections studied, SOCS1 expression is associated to an increase in susceptibility to the infection by facilitating pathogen replication. However, SOCS1 may on the other hand also reduce the pathological outcome of infections by hampering inflammatory reactions.

Several viral infections including herpes simplex virus, human respiratory virus, HIV and hepatitis C virus were found to induce SOCS1 expression<sup>188-191</sup>. Importantly, silencing of SOCS1 increased type I IFN signaling and resulted in inhibited viral replication in a human respiratory virus model<sup>189</sup>. Accordingly, *Socs1*-

deficient mice showed increased type I IFN-mediated resistance to Semliki Forest-virus<sup>166,170</sup>. However, unrestrained immune responses cause damaging immunopathology in other virus infections. In these cases, impairment of cytokine responses with SOCS1 may prevent inflammatory damage. Thus, in a vaccinia virus infection model, mice could be protected against lethal virus infection by the administration of a peptide mimicking the SOCS1 KIR region<sup>192</sup>.

In parasite infections, SOCS1 has been shown to inhibit responsiveness to IFN- $\gamma$ , as in an *in vitro* model *Socs1*<sup>-/-</sup> macrophages were capable of killing *Leishmania major* at significantly decreased IFN- $\gamma$  concentrations compared to SOCS1-sufficient macrophages<sup>170</sup>. However, increased pro-inflammatory signaling was not beneficial *in vivo* as *Socs1*<sup>+/-</sup> mice infected with *L. major* showed worsened pathology without a reduction in numbers of parasites<sup>193</sup>.

Studies on the role of SOCS1 in bacterial infections mainly focused on intracellular infections, in which IFN- $\gamma$  plays a major protective role. Research from our laboratory demonstrated that *Chlamydia pneumoniae* infection of macrophages induced SOCS1 expression<sup>194</sup>. Infected *Socs1*<sup>-/-</sup> macrophages displayed lower bacterial titers and higher levels of IFN-regulated genes as *iNos* and indoleamine dioxygenase that participate in the control of intracellular bacteria. *Rag1*<sup>-/-</sup>/*Socs1*<sup>-/-</sup> mice showed 10-fold lower bacteria numbers in lungs than controls 6 days after infection. However, *Rag1*<sup>-/-</sup>/*Socs1*<sup>-/-</sup> mice died within seven days after infection with *C. pneumoniae* showing a severe pulmonary inflammation, whereas *Rag1*<sup>-/-</sup> mice survived for more than 60 days. Thus, SOCS1 has a crucial role in preventing acute lethal inflammation in *C. pneumoniae* infection.

Infections with different mycobacterial species such as *M. bovis*<sup>195</sup>, *M. avium*<sup>196</sup> and *M. tuberculosis* were found to induce SOCS1 expression in macrophages<sup>195-198</sup>. This SOCS1 induction was shown to be mediated by different innate immune receptors as TLR2 and DC-SIGN in mycobacterial-infected myeloid cells<sup>198</sup>. Knockdown of SOCS1 using siRNA in mouse DCs resulted in increased killing of virulent *M. tuberculosis*<sup>198</sup>. However, whether deletion of SOCS1 is beneficial for the outcome of mycobacterial infections *in vivo* remains to be investigated and will be discussed in this thesis.

## 1.5 SOCS2

SOCS2 is induced in response to growth hormone (GH), prolactin, insulin, lipoxin, LPS and several cytokines<sup>199,200</sup>. Compared to SOCS1 and SOCS3, SOCS2 shows a complete different spectrum of functions, with GH and insulin-like growth factor signaling as main targets of SOCS2 inhibition. Additionally, several reports indicate that SOCS2 not only regulates GH receptor signaling but may also be involved in responses to infection<sup>201,202</sup>. In contrast to SOCS1 and SOCS3, SOCS2 lacks the N-terminal KIR domain so that the inhibitory function of SOCS2 is dependent on competitive binding via its SH2 domain and, even more importantly, on the ubiquitination and proteasomal degradation of the interacting receptor<sup>200</sup>.

In contrast to *Socs1*<sup>-/-</sup> mice, *Socs2*<sup>-/-</sup> mice are viable. However, *Socs2*<sup>-/-</sup> mice show a 40% increase in body weight alike GH-transgenic mice, and overgrowth of *Socs2*<sup>-/-</sup> mice could be attenuated by an additional deletion of STAT5, the STAT molecule that mediates GH signal transduction<sup>203,204</sup>.

### 1.5.1 SOCS2 in immunity and infection

Silencing of SOCS2 in human immature DCs demonstrated that the expression of SOCS2 was required for maturation of DCs after LPS stimulation<sup>199</sup>. Furthermore, *Socs2*<sup>-/-</sup> DCs were refractory to anti-inflammatory mediators like lipoxin (LXA4) and showed in turn increased expression of pro-inflammatory cytokines in response to microbial stimulation<sup>202</sup>. *Socs2*<sup>-/-</sup> mice infected with *Toxoplasma gondii* showed uncontrolled production of pro-inflammatory cytokines, aberrant leukocyte infiltration and elevated mortality that was most probably due to hyper-responsive DCs<sup>202</sup>.

Regarding T cell responses, SOCS2-deficient CD4<sup>+</sup> T cells displayed a Th2 predisposition, and *Socs2*<sup>-/-</sup> mice showed enhanced Th2 responses to helminthic infections, atrophic dermatitis and allergic lung inflammation<sup>201</sup>. This suggested SOCS2 as an inhibitor of Th2 development.

Even though SOCS2 is expressed in Th1 cells, *Leishmania major*-infected *Socs2*<sup>-/-</sup> mice showed the same level of disease progression than control mice, indicating that despite its potential involvement in Th1 immune responses, SOCS2 is not involved in the control of cutaneous leishmaniasis<sup>193</sup>. Yet, a role for SOCS2 in other Th1-controlled infections has not been studied and will be discussed later as part of this thesis.

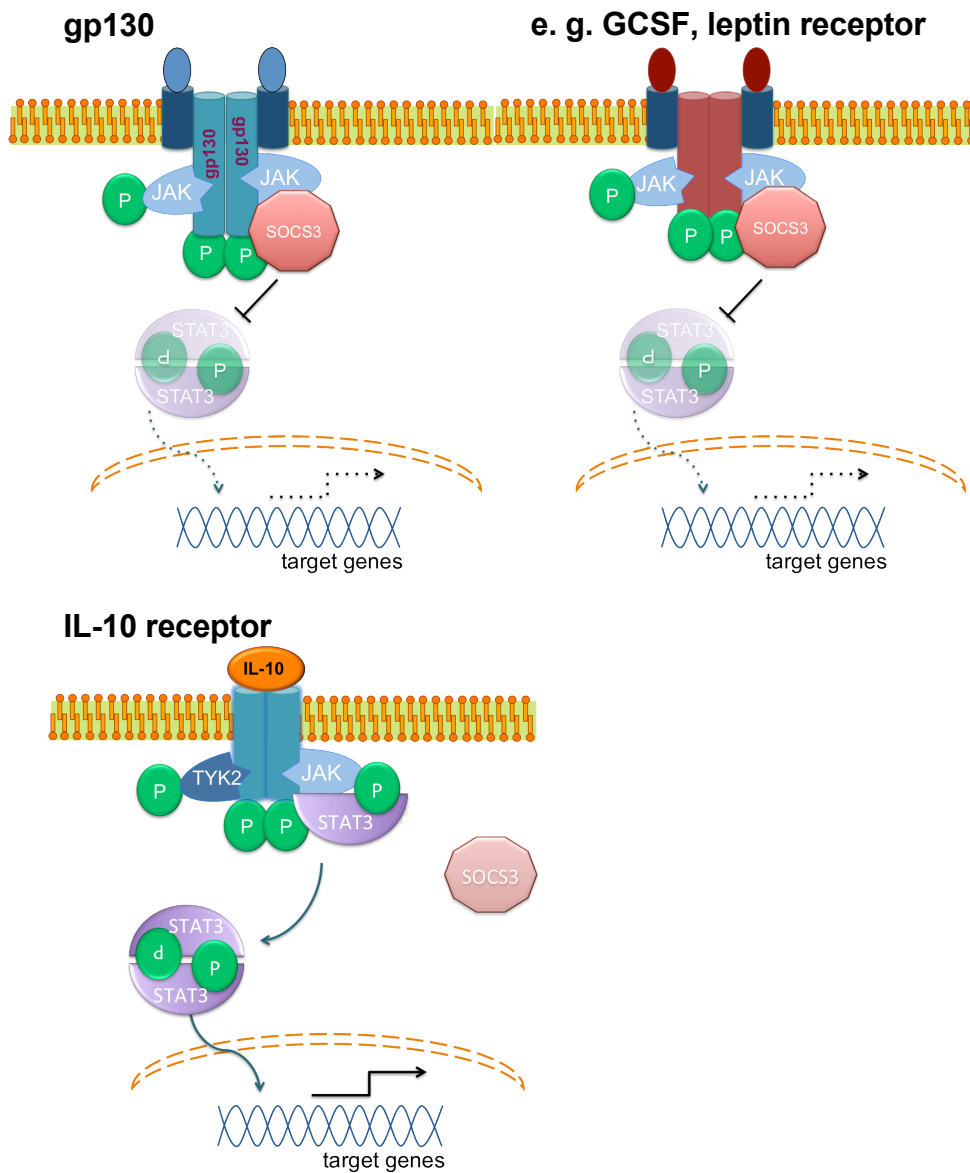
## 1.6 SOCS3 AND GP130

SOCS3 can be induced by different cytokines like leukemia inhibitory factor (LIF), IL-11, IL-10, IL-2, IL-6,<sup>158,205-208</sup> hormones including ciliary neutrophilic factor, leptin, prolactin and growth hormones<sup>209-212</sup> but also by microbial molecular patterns as LPS or CpG<sup>162,213</sup>.

SOCS3 is an important endogenous regulator of STAT3-mediated signaling. Interestingly, not all cytokine signaling pathways that activate STAT3 are SOCS3-regulated since SOCS3 binds to the gp130 and selected receptor subunits, whereas the signaling of IL-10 that also stimulates STAT3-activation is unaffected by SOCS3 (Fig. 5)<sup>77,125,126,214,215</sup>. This is explained by the ability of SOCS3 to bind to the IL-6 receptor subunit gp130 but not to the IL-10 receptor (Fig. 5)<sup>126,153,216</sup>. Furthermore, SOCS3 inhibits non-gp130-mediated cytokine and hormone receptor signaling pathways activated by G-CSF, leptin, IL-12 and IFN<sup>206,215,217-220</sup>.

The glycoprotein gp130 is a promiscuous cytokine receptor subunit that mediates signaling by IL-6 and all other cytokines belonging to the IL-6 family such as IL-11, IL-27, oncostatin M, cardiotropin 1, cardiotropin-like cytokine and LIF<sup>221</sup>. IL-6 binds to the IL-6R alpha chain, recruits gp130 and forms a hexameric complex consisting of two IL-6 bound to two IL-6R $\alpha$  and gp130 subunits<sup>222,223</sup>. It was found that gp130 could be present on hematopoietic cells depending on their activation status but also on non-hematopoietic cells<sup>224-226</sup>.

Binding of the ligand to gp130 induces activation of JAK1 kinases, which results in tyrosine phosphorylation of gp130. This creates binding sites for signaling transmitting factors at two different sites: phosphorylation of the four membrane-distal tyrosines leads to binding of STAT1 and STAT3, whereas phosphorylation of the membrane-proximal tyrosine (Y757) enables binding of the Src-homology tyrosine phosphatase-2 (SHP2)<sup>227-230</sup>. These bound molecules are in turn activated and trigger distinct intracellular signaling cascades.



**Figure 5: SOCS3-mediated inhibition of gp130 and non-gp130 signals but not of IL-10-signaling.**

Since SOCS3 cannot bind to the IL-10 receptor, STAT3 activation in response to IL-10 is unimpaired in the presence of SOCS3, whereas gp130, GCSF and leptin receptor signals are inhibited by SOCS3.

Interestingly, Y757 (SHP2-binding site) is also the binding site for SOCS3 and required for SOCS3-mediated inhibition of STAT3 activation<sup>231</sup>. Therefore, a mutation at Y757 has two effects: abrogation of SHP2 signaling and deletion of SOCS3-mediated STAT3 inhibition.

### 1.6.1 SOCS3 and gp130 mutations in mice

SOCS3 knockout mice die during embryogenesis due to placental defects<sup>232,233</sup>. Mortality is caused by enhanced LIF signaling in the absence of SOCS3 causing an altered trophoblast differentiation<sup>234</sup>. However, mice with a conditional knockdown for SOCS3 in specific tissues or cell lineages survive and have been used for targeted SOCS3 studies (see in materials and methods).

Since the knockdown of gp130 leads to embryonic or perinatal lethality, mice with conditional knockdowns for gp130 or mutations ablating either the STAT1/3 binding or SHP2/SOCS3 binding are used to dissect the role of gp130 in mouse models<sup>235</sup>. To study the role of SOCS3 within gp130 signaling, two different models can be used: One model, *gp130*<sup>F759</sup> mice, the mouse *gp130* is replaced with human *gp130* mutant cDNA, in which the crucial tyrosine for SOCS3 binding in position 759 (corresponds to 757 in the mouse) is replaced by phenylalanine (*gp130*<sup>F759</sup> mice), and the other model, in which tyrosine 757 is directly substituted by phenylalanine (*gp130*<sup>F/F</sup> mice)<sup>236,237</sup>. These models show similarities, a prolonged activation of STAT3 in response to gp130-activating factors was found in both models confirming the impairment of SOCS3 function by these mutations<sup>236,238</sup>. Phenotypically these mice developed splenomegaly that could be accounted to increased STAT3 activation by IL-6 as it is also found in IL-6/IL-6R $\alpha$  transgenic mice<sup>236,237,239</sup>. Furthermore, *gp130*<sup>F759</sup> mice showed autoimmune arthritis and lymphadenopathy, whereas in *gp130*<sup>F/F</sup> mice gastric adenoma formation and a reduced life span was found<sup>225,226,240</sup>. Combination of *gp130*<sup>F/F</sup> mice with deletion of IL-6 (*gp130*<sup>F/F</sup>/*Il-6*<sup>-/-</sup>) or reduction of STAT3 levels (*gp130*<sup>F/F</sup>/*Stat3*<sup>+/-</sup>), allows to study the underlying mechanisms caused by the changed gp130 signaling<sup>238</sup>.

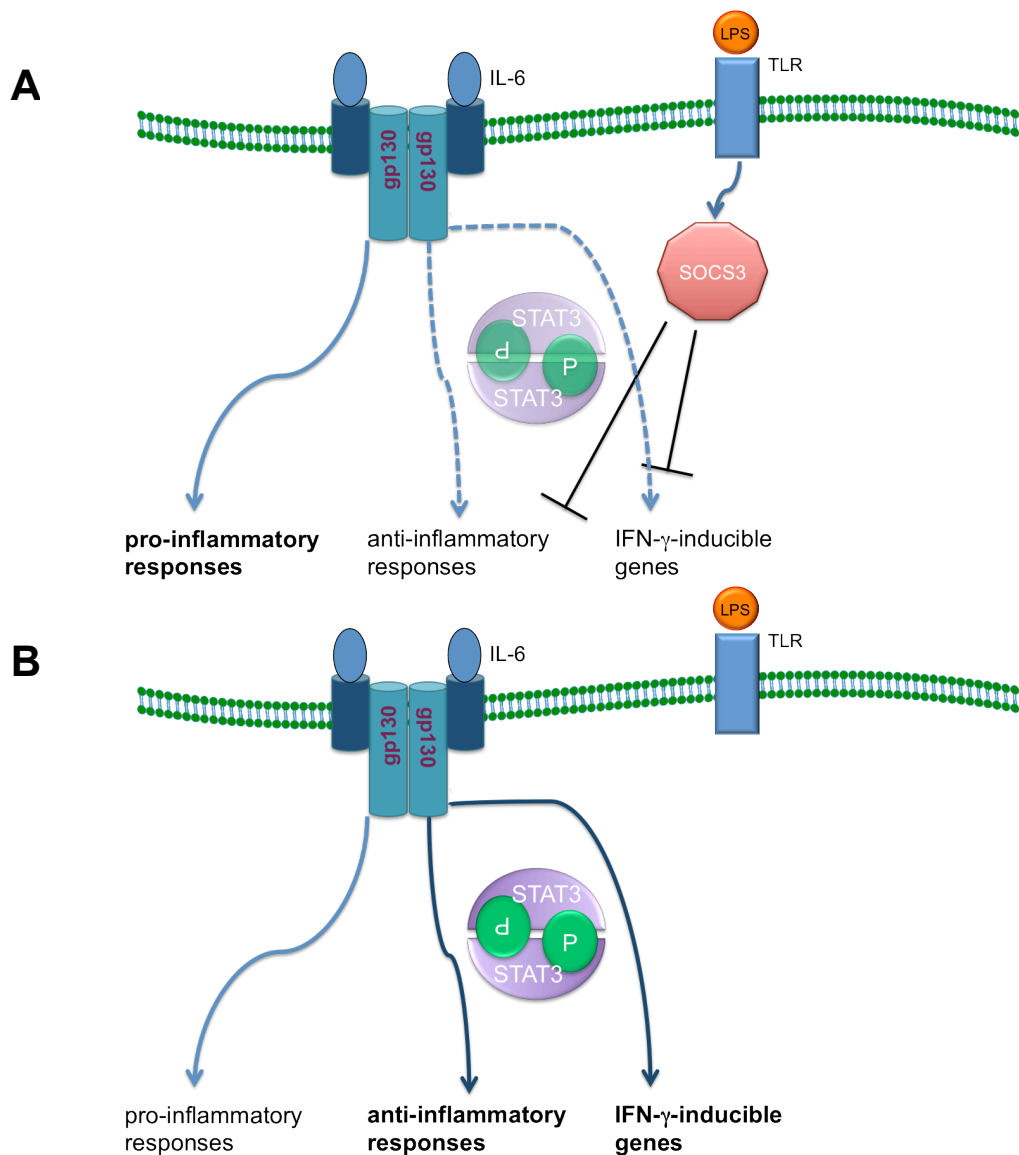
### 1.6.2 SOCS3 in myeloid cells

Mice with a conditional knockdown of SOCS3 in myeloid cells highlighted the crucial role of SOCS3 in suppression and termination of IL-6/gp130-mediated signaling. SOCS3-deficient macrophages stimulated with IL-6 displayed an increased magnitude and duration of STAT1 and STAT3 activation in comparison to controls<sup>125,126,241</sup>. This prolonged STAT3 activation is a hallmark of anti-inflammatory IL-10 signaling and LPS-induced pro-inflammatory responses like secretion of TNF and IL-12 were reduced in the presence of IL-6 in SOCS3-deficient macrophages (Fig. 6). The importance of SOCS3-regulated STAT3 activation was confirmed in *gp130<sup>F/F</sup>* macrophages that responded to IL-6 similar as to IL-10 under LPS stimulation<sup>242</sup>. Additionally, Croker (2003), Lang (2003) and co-workers found an increased STAT1 activation and elevated levels of IFN- $\gamma$  responsive genes in response to IL-6 using SOCS3-deficient macrophages (Fig. 6)<sup>125,241</sup>.

The anti-inflammatory properties of IL-6 in the absence of SOCS3 or SOCS3-binding may explain why mice deficient for SOCS3 in myeloid cells are resistant to LPS-induced endotoxic shock whereas *Stat3<sup>-/-</sup>* mice are highly susceptible<sup>126,243</sup>. Moreover, SOCS3-deficient DCs were found to be capable of suppressing the development of experimental autoimmune encephalitis (EAE)<sup>244,245</sup>.

Furthermore, deletion of SOCS3 in the myeloid or hematopoietic cells increased the numbers of neutrophils, which showed increased survival and proliferative capacity<sup>214,247</sup>. Following GCSF injection *in vivo*, SOCS3-deficient mice developed neutrophilia, and a spectrum of inflammatory pathologies characterized by neutrophil infiltration in multiples tissues<sup>248</sup>.





**Figure 6: SOCS3 inhibits anti-inflammatory and IFN-inducible genes in response to LPS in presence of IL-6 (adapted from <sup>246</sup>).** LPS-induced SOCS3 modulates IL-6 signaling leading to pro-inflammatory responses (A). In contrast in the absence of SOCS3, anti-inflammatory responses and expression of IFN- $\gamma$ -inducible genes are observed (B).

### 1.6.3 SOCS3 in T cells

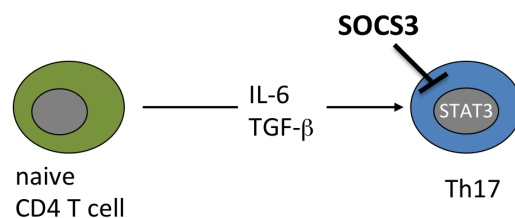
Even though T cell development in the thymus is unaffected by SOCS3<sup>249</sup>, several models suggest a role for SOCS3 in T cell proliferation. One indication for its regulatory role is the fact that SOCS3 levels are high in resting T cells, whereas they are low in activated T cells<sup>250</sup>. Furthermore, T cells overexpressing SOCS3 showed a reduced proliferation upon T cell receptor stimulation<sup>251,252</sup>, whereas SOCS3-deficient T cells were hyper-proliferative towards T cell receptor stimulation<sup>253,254</sup>. Interestingly, T cells from *gp130*<sup>F/F</sup> mice behaved alike SOCS3-deficient T cells,

suggesting that hypersensitivity to gp130 cytokines may drive this enhanced proliferation<sup>253</sup>.

SOCS3 protein levels may influence the Th1/Th2 balance, as expression was found to be low in Th1 cells and high in Th2 cells<sup>255,256</sup>. Accordingly, overexpression of SOCS3 in mice resulted in elevated Th2 responses and hypersensitivity to allergic diseases<sup>256</sup>. A possible explanation for the increased Th2 development under high SOCS3 levels might be that binding of SOCS3 to IL-12R $\beta$ 2 inhibits IL-12-mediated STAT4 activation that in turn impairs Th1 differentiation<sup>256,257</sup>.

Interestingly, a conditional knockdown for SOCS3 in T cells mice showed reduced Th2 responses together with increased numbers of Th3-like T cells secreting IL-10 and TGF- $\beta$ <sup>258</sup>. Unexpectedly, Th1 responses were not elevated in the absence of SOCS3, which may have been caused by the presence of immunosuppressive cytokines<sup>258</sup>.

Other reports indicated only minimal effects of SOCS3 deletion in T cells on the Th1/Th2 balance but found an increased Th17 generation and enhanced STAT3 activation in response to IL-23<sup>249</sup>. This data was confirmed in a study of an atherosclerosis model using mice with a T cell specific SOCS3 knockdown as well as in a rheumatoid arthritis model, in which mice lacking SOCS3 in hematopoietic and endothelial cells showed increased Th17 development<sup>259,260</sup>. These findings fit well with the known mechanism for Th17 differentiation, as STAT3 activation either by IL-23 or IL-6 is crucial for Th17 development (Fig. 7). Furthermore, TGF- $\beta$ , another cytokine necessary for Th17 generation, was shown to inhibit SOCS3 expression with the resulting prolonged STAT3 activation and thereby promoting Th17 development<sup>261</sup>.



**Figure 7: Role of SOCS3 in Th17 cell development.**

STAT3 activation is essential for the generation of Th17 cells and can be inhibited by SOCS3.

No SOCS3 protein expression was found in Tregs<sup>262</sup>. Accordingly, in SOCS3-deficient T cells no defects in Treg differentiation and function were reported<sup>254</sup>. Recently, SOCS3 has been assigned a role in development of memory CD8+ T cells since numbers of memory CD8+ T cells were dramatically reduced in the absence of SOCS3 in T cells<sup>263,264</sup>.

#### 1.6.4 SOCS3 in infectious diseases

Similar to SOCS1 and SOCS2, SOCS3 expression can be stimulated by both cytokines and TLR agonists. Additionally, several pathogens including viruses, bacteria and parasites have been shown to induce SOCS3 expression. Due to the diversity of described SOCS3 actions, it is hard to predict in which cases SOCS3 expression will be beneficial for the host.

SOCS3 was found to inhibit STAT1 activation in response to type I IFNs but was less effective than SOCS1 in such function<sup>215,219</sup>. Infections with several viruses stimulated SOCS3 expression that correlated with reduced STAT1 activation in response to type I IFNs. Silencing of SOCS3 significantly increased IFN signaling and hampered viral replication. Similar findings were obtained during different viral infections e.g. HSV, Influenza A, hepatitis C virus and Epstein Barr virus infection indicating that increased SOCS3 expression improves viral replication<sup>265-269</sup>.

Surprisingly, mice with a T cell-specific SOCS3 deletion showed increased T cell activation and viral clearance without development of immunopathology during the infection with lymphocytic choriomeningitis virus (LCMV)<sup>270</sup>. Treatment of LCMV-infected mice with IL-7 repressed SOCS3 expression and promoted IL-6 production, which resulted in enhanced T cell effector functions and viral clearance. On the contrary, *Leishmania major*-infected mice were more susceptible in the absence of SOCS3 in T cells. In this case, SOCS3 deficiency in T cells led to increased anti-inflammatory TGF- $\beta$  secretion promoting the infection<sup>258</sup>.

Infection of macrophages with *M. avium* and *M. bovis* raised SOCS1 and SOCS3 levels in a TLR2-NOTCH1 dependent pathway<sup>195,196,271</sup>. In clinical studies, patients with active tuberculosis were found to have higher SOCS3 expression levels in whole blood and T cells in comparison to latently infected controls<sup>272,273</sup>. Accordingly, SOCS3 expression decreased under chemotherapy treatment indicating that SOCS3 expression could be interpreted as an on-going infection response<sup>272,273</sup>.

However, to prove that SOCS3 plays a crucial role for tuberculosis control, more work needs to be done. This was one of the objectives of my thesis.

Regarding SOCS3-ablations due to gp130 mutations, *gp130*<sup>F759</sup> mice showed an attenuated early phase of defense against *Listeria monocytogenes* infection resulting in higher bacterial loads and mortality most probably due to insufficient elevation of IFN- $\gamma$  levels<sup>274</sup>. Furthermore, low IFN- $\gamma$  titers led to increased susceptibility to *Toxoplasma gondii* in mice deficient for SOCS3 in myeloid cells as well as in *gp130*<sup>F/F</sup> mice.<sup>275,276</sup> In both mouse strains, lower IL-12p40 and therefore lower IFN- $\gamma$  levels after infection compared to control mice were found. Interestingly, addition of IL-12 as well as neutralization of IL-6 could restore the wild type (WT) phenotype in both models demonstrating that an altered IL-6 signaling was responsible for the observed increased susceptibility<sup>275,276</sup>.

## 2 AIMS

*M. tuberculosis* is a highly successful pathogen. Even though the literature is extensive, detailed reasons why a latent tuberculosis patient will convert to active disease and the mechanisms that regulate the control of *M. tuberculosis* are not completely understood. More knowledge about protective immune responses during *M. tuberculosis* infection is required for the design of prophylaxis and therapy.

SOCS proteins are non-redundant regulators of immune responses through inhibition of cytokine signaling. Therefore, these molecules are interesting target candidates that may modulate the outcome of infections.

In my thesis work, I have studied the role of SOCS1, SOCS2 and SOCS3, which have different specific cytokine signaling pathways as their targets, in the outcome of mycobacterial infections.

Specific questions:

### SOCS1

- How is SOCS1 expression regulated in mycobacteria-infected macrophages?
- What is the role of SOCS1 in the regulation of intracellular bacterial levels and cytokine expression in mycobacteria-infected macrophages *in vitro*?
- What is the role of SOCS1 *in vivo* during *M. tuberculosis* or BCG infection regarding
  - Bacterial loads
  - Cytokine responses
  - Immunopathology?

### SOCS2

- How is SOCS2 expression regulated in response to TLR stimulation?
- How is SOCS2 expression regulated in mycobacteria-infected macrophages?
- Is there a role for SOCS2 during *M. tuberculosis* infection *in vivo*?

## SOCS3

- How is SOCS3 expression regulated in mycobacteria-infected macrophages?
- Which function has SOCS3 in macrophages/DCs during mycobacterial infections regarding the regulation of
  - Bacterial loads
  - Cytokine responses
  - Immunopathology?
- What is the role of SOCS3 in T cells during *M. tuberculosis* infection *in vivo* regarding the regulation of
  - Bacterial loads
  - Cytokine responses
  - Immunopathology?
- Which function has gp130 signaling in SOCS3-deficient models *in vivo* and *in vitro* during mycobacterial infections?

### 3 MATERIAL AND METHODS

The methods applied in our studies are described in detail in the published articles and in the manuscript. Here, I want to give an overview about the genetically modified mice that have been used.

#### SOCS-unrelated

mouse strain	type of modification	effect
<i>Rag1</i> <sup>-/-</sup>	knockout	disruption of the <i>Rag1</i> gene in all cells no mature B and T cells
<i>MyD88</i> <sup>-/-</sup>	knockout	disruption of the <i>MyD88</i> gene in all cells
<i>Irf3</i> <sup>-/-</sup>	knockout	disruption of the <i>Irf3</i> gene in all cells
<i>Ifn-γ</i> <sup>-/-</sup>	knockout	disruption of the <i>Ifn-γ</i> gene in all cells

#### SOCS proteins and SOCS-related

mouse strain	type of modification	effect
<i>Rag1</i> <sup>-/-</sup> / <i>Socs1</i> <sup>-/-</sup>	double knockout	no mature B and T cells disruption of the <i>Rag1</i> and <i>Socs1</i> gene in all cells
<i>Ifn-γ</i> <sup>-/-</sup> / <i>Socs1</i> <sup>-/-</sup>	double knockout	disruption of the <i>Ifn-γ</i> and <i>Socs1</i> gene in all cells
<i>Socs1</i> <sup>fl/fl</sup> <i>LysM cre</i> *	conditional knockdown	excision of the <i>Socs1</i> gene in myeloid cells
<i>Socs2</i> <sup>-/-</sup>	knockout	disruption of the <i>Socs2</i> gene in all cells
<i>Socs3</i> <sup>fl/fl</sup> <i>LysM cre</i> *	conditional knockdown	excision of the <i>Socs3</i> gene in myeloid cells
<i>Socs3</i> <sup>fl/fl</sup> <i>lck cre</i> *	conditional knockdown	excision of the <i>Socs3</i> gene in T cells
<i>gp130</i> <sup>F/F</sup>	knockin	mutation of the SOCS3 binding site in gp130 in all cells
<i>gp130</i> <sup>F/F</sup> / <i>Il-6</i> <sup>-/-</sup>	knockin/knockout	mutation of the SOCS3 binding site in gp130 in all cells disruption of the <i>Il-6</i> gene in all cells
<i>gp130</i> <sup>F/F</sup> / <i>Stat3</i> <sup>+/-</sup>	knockin/knockout	mutation of the SOCS3 binding site in gp130 in all cells heterozygous for <i>Stat3</i> gene knockout

#### \*Conditional knockdown:

Mice were generated using the Cre-lox combination system, in which the Cre DNA-recombinase is expressed under a cell-specific promoter. Although the target gene is flanked by *LoxP* sequences in all cells for recognition by the Cre enzyme, it will only be deleted specifically in Cre-expressing cells. The corresponding WT mice contain the floxed gene (*Socs1*<sup>fl/fl</sup> and *Socs3*<sup>fl/fl</sup>) but are negative for Cre expression.

## 4 RESULTS AND DISCUSSION

### 4.1 SOCS1 DURING *M. TUBERCULOSIS* INFECTION (PAPER I)

IFN- $\gamma$  is essential for protection against *M. tuberculosis* infection. SOCS1 is induced during infection of macrophages with *Mycobacterium avium* and *Mycobacterium bovis* BCG, and may thereby inhibit responses to IFN- $\gamma$  and play a central role in the outcome of mycobacterial infections<sup>195,196</sup>. Thus, we hypothesized that stimulation of SOCS1 expression could constitute an immune-escape mechanism for the establishment and chronicity of *M. tuberculosis* infection.

*How is SOCS1 expression regulated in mycobacteria-infected macrophages?*

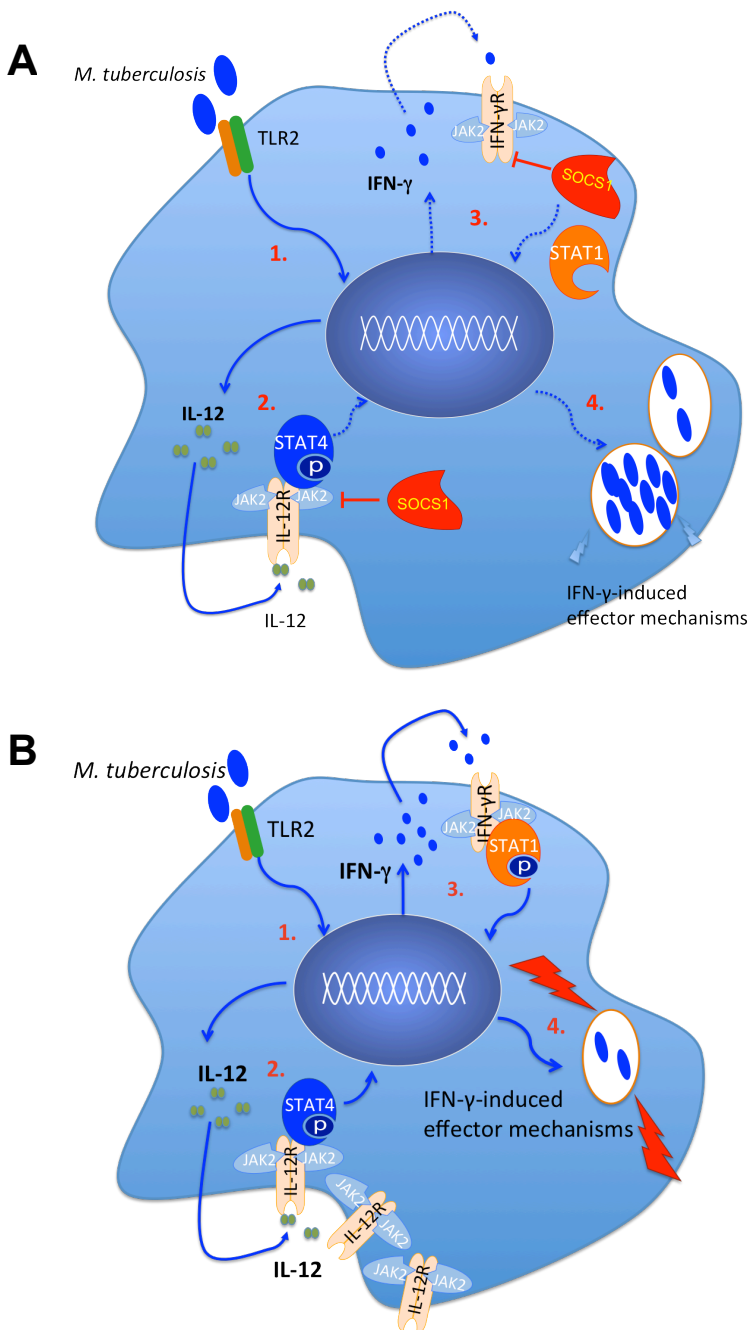
We found that *Socs1* mRNA was stimulated during BCG and *M. tuberculosis* infections *in vitro* in human and murine macrophages as well as in a mouse model in lung tissue during infections. Using macrophages from genetically manipulated mice, we found that the expression of *Socs1* mRNA during BCG infection required phagocytosis and the presence of NOD2 and IFN- $\alpha/\beta$  receptors, the adaptor protein MyD88 and the common transcription factor NF- $\kappa$ B. This shows that TLR and non-TLR signals cooperate in optimal *Socs1* mRNA expression in mycobacteria-infected macrophages. Furthermore, our data was confirmed by a recent publication, in which *M. tuberculosis*-induced *Socs1* mRNA expression was dependent on TLR and DC-SIGN receptor 1 in human DCs<sup>198</sup>. Moreover, our finding of SOCS1 induction in lungs tissue of BCG and *M. tuberculosis*-infected mice is further supported by clinical data that demonstrated increased SOCS1 expression in induced sputum of tuberculosis patients compared to healthy individuals and patients with other lung diseases<sup>277</sup>. Mechanisms of SOCS1 action in the mouse model may therefore provide insights into the role of the molecule in determining the onset and course of active tuberculosis in patients.

*What is the role of SOCS1 in the regulation of intracellular bacterial levels and cytokine expression in mycobacteria-infected macrophages in vitro?*

To study the function of SOCS1 *in vitro*, we took advantage of cells derived from *Socs1* knockout and conditional knockdown mice as well as SOCS1 mimetic peptides. We found that SOCS1 inhibited growth control of *M. tuberculosis* and BCG



in macrophages. Infected *Socs1*<sup>-/-</sup> macrophages showed lower mycobacterial numbers and expressed higher levels of IFN- $\gamma$ -responsive genes, but interestingly, also higher levels of IFN- $\gamma$  itself than infected control cells. In agreement, *Socs1*<sup>-/-</sup> macrophages secreted higher amounts of IFN- $\gamma$  in response to IL-12 that was associated with increased IL-12R $\beta$ 1 but not IL-12 expression in infected *Socs1*<sup>-/-</sup> macrophages. Neutralization of IL-12 reduced the elevated IFN- $\gamma$  expression in infected *Socs1*<sup>-/-</sup> macrophages. Of importance, the improved control of *M. tuberculosis* and BCG by *Socs1*<sup>-/-</sup> macrophages was lost in *Ifn- $\gamma$* <sup>-/-</sup>/*Socs1*<sup>-/-</sup> macrophages, demonstrating that IFN- $\gamma$  secretion mediated the improved mycobacterial control of *Socs1*<sup>-/-</sup> macrophages. Moreover, *M. tuberculosis*-infected WT macrophages were unimpaired in their response to addition of exogenous IFN- $\gamma$  despite SOCS1 expression. Altogether, our results suggest that SOCS1 regulates the secretion rather than the response to IFN- $\gamma$ , via controlling responses to IL-12 causing the increased mycobacterial resistance of *Socs1*<sup>-/-</sup> macrophages (Fig. 8). This mechanism is unexpected, as to our knowledge, all described effects of SOCS1 on infected host cells were explained by the inhibition of STAT1 signaling leading to impaired type I IFN-signaling for viral infections or IFN- $\gamma$ -signaling for intracellular bacteria and parasites<sup>166,170,189,195,196</sup>. Since SOCS1 binds to JAK2 that is also associated with the IL-12 receptor, this interaction probably influences IL-12 signaling. Furthermore, *Socs1*<sup>-/-</sup>/*Stat4*<sup>-/-</sup> mice, deficient for the IL-12 signal-transmitting molecule STAT4, live longer than *Socs1*<sup>-/-</sup> mice<sup>175</sup>. This suggests an inhibitory role of SOCS1 on STAT4 activation and increased responses to IL-12 in absence of SOCS1 have been found in T cells and DCs<sup>175,278</sup>. Thus, our findings broaden the inhibitory spectrum of SOCS1 from IFN responses in macrophages to IL-12 responses during infections (Fig. 8).



**Figure 8: *M. tuberculosis*-infected macrophages in presence (A) and absence (B) of SOCS1.**

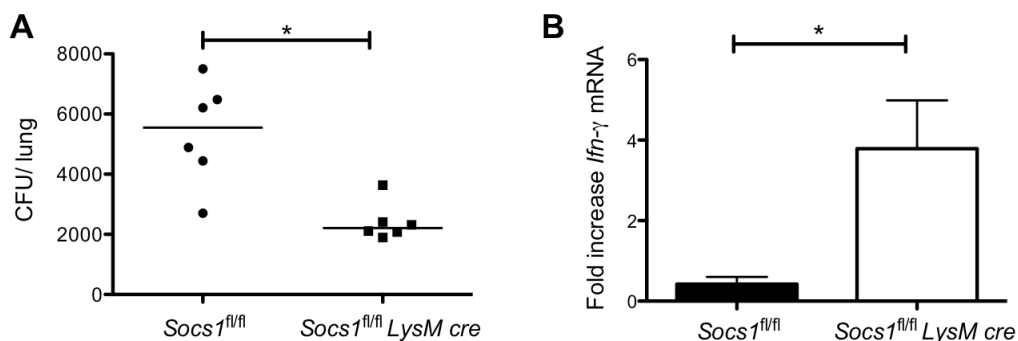
Infection of macrophages with *M. tuberculosis* stimulates innate immune responses (1) such as the production of IL-12. Responses to IL-12 are inhibited in the presence of SOCS1 (2) leading to reduced IFN- $\gamma$  induction and IFN- $\gamma$ -signaling (3). Therefore, presence of SOCS1 results in higher bacterial numbers (4) and reduced IFN- $\gamma$ -induced effector mechanisms.

To obtain a detailed overview about SOCS1-affected genes during *M. tuberculosis* infection in macrophages, we performed a genome wide expression microarray. In agreement with previous results, a significant fraction of the macrophage genome was altered after *M. tuberculosis* infection<sup>116,279</sup>. Unexpectedly,

the majority of infection-regulated genes was even stronger regulated in the absence of SOCS1, suggesting a major role of SOCS1 in the control of macrophage activity during *M. tuberculosis* infection. The majority of genes involved in the defense and immune responses and almost all IFN- $\gamma$ -regulated genes showed higher levels in infected *Socs1*<sup>-/-</sup> macrophages, whereas genes that were decreased in infected control cells were even further downregulated in infected *Socs1*<sup>-/-</sup> macrophages. Altogether, our data implicates that the presence of SOCS1 attenuates both the negative and positive regulation of the majority of genes that change their expression after infection.

*What is the role of SOCS1 during in vivo M. tuberculosis or BCG infection regarding bacterial loads, cytokine responses and immunopathology?*

Importantly, the improved bacterial control in *Socs1*<sup>-/-</sup> macrophages *in vitro* was reflected in the role of SOCS1 during *M. tuberculosis* infection *in vivo*. Aerosol infection with *M. tuberculosis* of *Rag1*<sup>-/-</sup>/*Socs1*<sup>-/-</sup> as well as *Socs1*<sup>fl/fl</sup> *LysM cre* mice showed lower bacterial loads in the lungs 7 days after infection compared to control mice (Fig. 9). Moreover, *Ifn- $\gamma$*  and *iNos* mRNA levels in the lungs of *Rag1*<sup>-/-</sup>/*Socs1*<sup>-/-</sup> and *Socs1*<sup>fl/fl</sup> *LysM cre* mice were elevated compared to control mice (Fig. 9). At this early stage of infection, mainly innate immune mechanisms are present, and no increase of IFN- $\gamma$  in lungs of infected control mice was detected. Consistent with our *in vitro* results, lungs from infected *Rag1*<sup>-/-</sup>/*Socs1*<sup>-/-</sup> and *Socs1*<sup>fl/fl</sup> *LysM cre* mice showed higher levels of IFN- $\gamma$  that are sufficient to impair bacterial growth *in vivo*, since no improved bacterial control was observed in *Ifn- $\gamma$* <sup>-/-</sup>/*Socs1*<sup>-/-</sup> mice compared to *Ifn- $\gamma$* <sup>-/-</sup> mice.



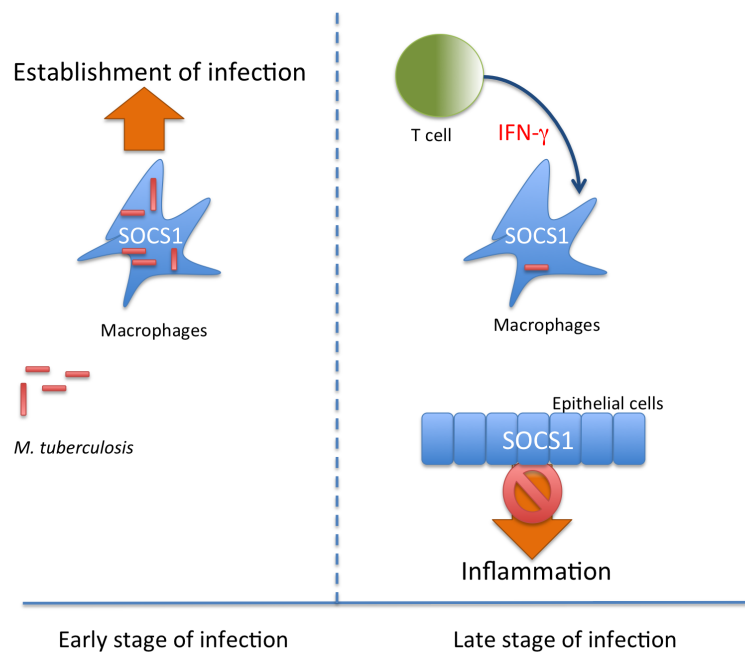
**Figure 9: SOCS1 in myeloid cells 7 days after *M. tuberculosis* infection.**

7 days after *M. tuberculosis* infection, bacterial loads in the lungs of individual *Socs1*<sup>fl/fl</sup> and *Socs1*<sup>fl/fl</sup> *LysM cre* mice and median were determined (A). Expression of *Ifn- $\gamma$*  mRNA in lungs is shown in (B) as mean  $\pm$  SEM.

Three weeks after infection, in conjunction with the appearance of IFN- $\gamma$  producing T cells in lungs of aerosol infected mice, similar bacterial counts were registered in the lungs of *Rag1<sup>-/-</sup>/Socs1<sup>-/-</sup>*, *Socs1<sup>fl/fl</sup> LysM-cre* and their respective control mice. We suggest that, at this later time point, the advantage of the SOCS-deficient host is lost when IFN- $\gamma$  secretion by T cells is prominent and macrophages even in the presence of SOCS1 are able to respond to IFN- $\gamma$ . Relevant to our observation, SOCS1 expression is elevated in the blood of patients with pulmonary tuberculosis that at the same time express an IFN-induced gene signature<sup>280</sup>. As we found no differences in IFN- $\gamma$  and IFN- $\gamma$ -regulated genes in control and *Rag1<sup>-/-</sup>/Socs1<sup>-/-</sup>* mice at these later stage, it is possible that the inhibitory effects of SOCS1 can be shadowed by a potent Th1-mediated immunity.

Additionally, both *Rag1<sup>-/-</sup>/Socs1<sup>-/-</sup>* and also *Ifn- $\gamma$ <sup>-/-</sup>/Socs1<sup>-/-</sup>* mice showed a very severe pulmonary inflammation 3 weeks after *M. tuberculosis* infection. This inflammatory response was mediated by SOCS1 deficiency in non-macrophage cells, because it was not observed in *M. tuberculosis*-infected *Socs1<sup>fl/fl</sup> LysM-cre* mice, which are lacking SOCS1 only in myeloid cells. We conclude that SOCS1-expressing non-macrophage cells are responsible for preventing a detrimental inflammation that is elicited at least partially in an IFN- $\gamma$ -independent manner. This is an interesting finding as in the absence of SOCS1 inflammation was reported to be mediated to a great extent by increased IFN- $\gamma$  secretion by T cells and elevated responsiveness of macrophages to IFN- $\gamma$ <sup>170,171,173</sup>. SOCS1 has also been shown to play a role in type I IFN, IL-4, IL-13 and IL-12 signaling, but apart from IL-12, these cytokines are not classically associated to resistance against *M. tuberculosis* infections<sup>281,282</sup>. However, whether these cytokines can contribute to mycobacterial infection-mediated inflammation remains to be examined.

Overall, we demonstrated that *M. tuberculosis*-induced SOCS1 by diminishing IL-12 responses impaired IFN- $\gamma$  secretion by macrophages. This resulted in lower levels of IFN- $\gamma$ -regulated genes and promoted *M. tuberculosis* infection leading to increased pulmonary bacterial levels in the lungs at early time points. Later during infection, mainly T cells produced IFN- $\gamma$  and SOCS1 in non-macrophages cells protected mice from severe inflammation thereby allowing the establishment of chronic infection (Fig.10).



**Figure 10: Role of SOCS1 at different stages of *M. tuberculosis* infection.**

SOCS1 in macrophages increases bacterial numbers at the early stage of infection, whereas it has no effect on bacterial control at later time points. SOCS1 in non-macrophage cells (e. g. epithelial cells) hampers inflammation at later stages.

## 4.2 SOCS2 (PAPER II)

Little is known about the regulation of SOCS2 expression in immune cells. In paper II we investigated the regulation of SOCS2 expression in response to TLR4 agonists and to *Mycobacterium bovis* BCG in macrophages.

*How is SOCS2 expression regulated in response to TLR4 stimulation?*

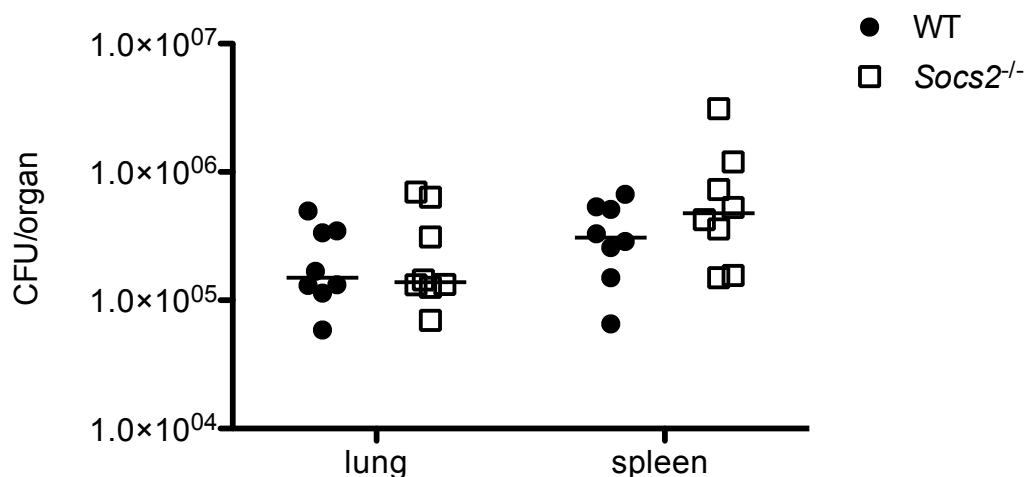
We demonstrated that LPS-induced SOCS2 expression in human monocyte-derived DCs was dependent on the transcription factors IRF1 and IRF3 as shown by silencing IRFs. The expression and activation of IRF1 and IRF3 is known to be stimulated by type I IFNs, and neutralization of type I IFNs abolished *Socs2* mRNA expression after TLR4 stimulation. Silencing of different STAT transcription factors revealed the requirement of STAT3 and STAT5 in LPS-induced *Socs2* mRNA expression. TLR-signaling has been reported to mediate a direct induction of SOCS2 expression<sup>283</sup> but we clearly could show that inhibition of IRF1, IRF3 and IFN- $\alpha/\beta$  receptors severely impaired LPS-stimulated SOCS2 induction.

### How is SOCS2 expression regulated in mycobacteria-infected macrophages?

*Socs2* mRNA levels increased in BMM after infection with *M. bovis* BCG. The reduction of *Socs2* transcript induction in macrophages derived from *Irf3*<sup>-/-</sup> and *MyD88*<sup>-/-</sup> mice suggested the involvement of signaling via TLR or other innate immune receptors.

### Is there a role for SOCS2 during *M. tuberculosis* infection in vivo?

Since *Socs2* mRNA expression was stimulated by *M. bovis* BCG, we studied whether SOCS2 played a role in the outcome of infection with *M. tuberculosis*. *Socs2*<sup>-/-</sup> and control mice were infected via aerosol with *M. tuberculosis* and bacterial burdens in lungs and spleens were determined 6 weeks after infection (Fig. 11).



**Fig. 11: SOCS2 expression does not change *M. tuberculosis* infection.**

WT and *Socs2*<sup>-/-</sup> mice were infected via aerosol with *M. tuberculosis* Harlingen. Mice were sacrificed 6 weeks after infection and bacterial colony forming units (CFU) in lungs and spleens determined. Individual counts and median of each group is shown.

No differences in bacterial titers were detected, indicating that SOCS2 is not involved in the control of *M. tuberculosis*. This result contrasts with the role of SOCS2 during infection with *Toxoplasma gondii* that showed increased cumulative mortality in *Socs2*<sup>-/-</sup> mice<sup>202</sup>. The heightened morbidity and mortality of *T. gondii*-infected *Socs2*<sup>-/-</sup> mice was explained by the finding that *Socs2*<sup>-/-</sup> DCs were refractory to the anti-inflammatory properties of the lipoxin LXA4<sup>202</sup>. Both *M. tuberculosis* and *T. gondii* induce LXA4, an eicosanoid mediator with potent anti-inflammatory properties, that e.g. reduces IL-12 secretion by DC<sup>284,285</sup>. However, only *T. gondii*-infected 5-lipoxygenase-deficient mice (that are deficient for LXA4) succumbed earlier to infection

than control mice due to lethal encephalitis, whereas *M. tuberculosis*-infected 5-lipoxygenase-deficient mice displayed an improved control of infection without concomitant immunopathology<sup>285,286</sup>. This indicates a vital role for LXA4 and its regulation by SOCS2 during *T. gondii* infection in prevention of fatal immune activation. However, our results suggest that SOCS2-mediated lipoxin LXA4 regulation is redundant for the control of infection with *M. tuberculosis*.

Regarding a possible role for SOCS2 in controlling T cells responses, enhanced generation of Th2 cells was described in helminth-infected *Socs2*<sup>-/-</sup> mice<sup>201</sup>. For *M. tuberculosis* control, specific Th1 cell-mediated immunity is fundamental 6 weeks after infection. Thus, we conclude that the absence of SOCS2 does not hamper Th1 responses during *M. tuberculosis* infection, confirming previous findings, in which SOCS2 was redundant for the control of *Leishmania major* infection<sup>193</sup>.

These results suggest that although SOCS2 is induced during mycobacterial infections, its role is minor or redundant for disease control.

#### **4.3 SOCS3 DURING *M. TUBERCULOSIS* INFECTION (PAPER III)**

SOCS3 was shown to be induced in macrophages in response to mycobacterial infection with *M. avium* and *M. bovis* and in T cells of patients with active tuberculosis<sup>195,196,272</sup>. Since SOCS3 has multiple functions in the regulation of different cytokine signaling pathways with some of them unapparent in their function of tuberculosis control, we investigated the role of SOCS3 in the regulation of *M. tuberculosis* infection.

##### *How is SOCS3 expression regulated in mycobacteria-infected macrophages?*

In a first set of experiments, we found that *Socs3* mRNA expression was increased in lung tissue of *M. tuberculosis* and BCG-infected mice with around 10-fold higher levels during *M. tuberculosis* than in BCG infection. *Socs3* transcript induction in infected BMM was dependent on MyD88 and NF-κB but not on IRF3. To study whether this SOCS3 induction is important during *M. tuberculosis* infection, we examined the outcome of infection in different mouse models.

*Which function has SOCS3 in macrophages/DCs during mycobacterial infections regarding the regulation of bacterial loads, cytokine responses and immunopathology?*

Mice deficient for SOCS3 in myeloid cells (*Socs3<sup>fl/fl</sup> LysM cre* mice) showed increased bacterial burdens in lungs and spleens from 16 days after infection with *M. tuberculosis*. *Socs3<sup>fl/fl</sup> LysM cre* mice also displayed enhanced immunopathology and succumbed 82 days after infection (median survival), whereas control mice survived for 150 days until the experiment was terminated. Moreover, increased bacterial loads in lungs and spleen were observed after BCG infection of *Socs3<sup>fl/fl</sup> LysM cre* mice, although differences to bacterial loads in control mice were smaller compared to those registered during *M. tuberculosis* infection.

Next, we studied whether SOCS3-deficient macrophages were hampered in their control of mycobacterial growth. Surprisingly, *Socs3<sup>fl/fl</sup> LysM cre* BMM displayed unimpaired control of intracellular of *M. tuberculosis* replication and reduced bacterial loads in response to exogenous IFN- $\gamma$ . Thus, defects of the bactericidal/ bacteriostatic capacity of SOCS3-deficient macrophages are unlikely to cause the increased *in vivo* susceptibility to *M. tuberculosis* of *Socs3<sup>fl/fl</sup> LysM cre* mice. In line with our data, previous work revealed that SOCS3-deficient macrophages could control *T. gondii* infection even though *Socs3<sup>fl/fl</sup> LysM cre* mice were highly susceptible to the infection with the parasite<sup>276</sup>.

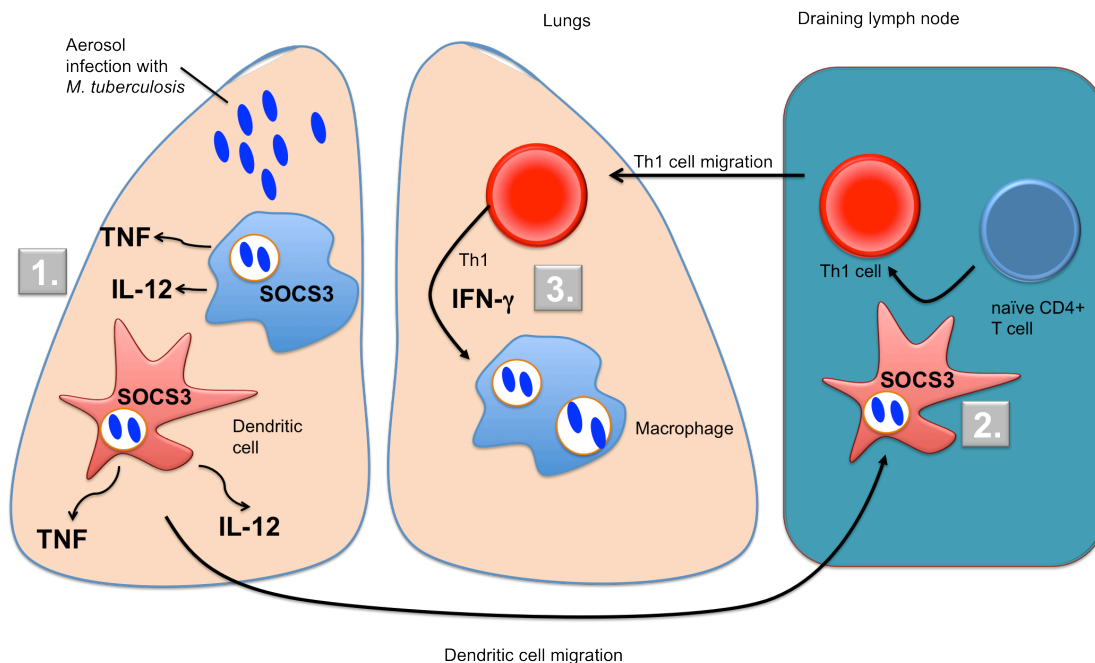
Thereafter, we analyzed the presence of cytokines and inflammatory mediators in supernatants of mycobacteria-infected macrophages. We found lower levels of TNF and IL-12p40 and higher levels of nitric oxide (NO) in supernatants of SOCS3-deficient macrophages after infection with *M. tuberculosis* or BCG and after stimulation with TLR-agonists in comparison to control cells. All three molecules – TNF, IL-12 and NO- have been demonstrated to be critical for control of clinical and experimental mycobacterial infections<sup>82,83,96,116,117</sup>. The elevated NO levels could explain the slightly improved bacterial growth control *in vitro*, whereas reduced IL-12p40 levels mainly would be important *in vivo* due to its crucial role in the generation of Th1 responses<sup>97,99,101</sup>. Previous studies showed reduced secretion of TNF and IL-12p40 in *Socs3<sup>fl/fl</sup> LysM cre* BMM in response to LPS, but only in co-culture with IL-6<sup>126</sup>. Interestingly, differences in TNF and IL-12p40 levels between mycobacteria-infected *Socs3<sup>fl/fl</sup> LysM cre* and control BMM were found even in absence of exogenously added IL-6. Addition of recombinant IL-6 to the cultures further diminished IL-12 and TNF concentrations in supernatants of mycobacteria-infected *Socs3<sup>fl/fl</sup> LysM cre* BMM. Furthermore, it has previously been shown that neutralization



of IL-6 in IFN- $\gamma$ /LPS stimulated cultures of *Socs3<sup>fl/fl</sup> LysM cre* BMM enhanced IL-12p40 production<sup>276</sup>. The increased NO and iNOS levels in BCG-infected *Socs3<sup>fl/fl</sup> LysM cre* BMM could reflect the described IFN- $\gamma$ -like response to IL-6 in *Socs3<sup>fl/fl</sup> LysM cre* BMM<sup>125,241</sup>.

Moreover, we found lower IL-12p40 expression levels in *M. tuberculosis* and BCG-infected SOCS3-deficient BMDC. Thus, we studied whether IFN- $\gamma$  levels and secretion were impaired in lungs of *M. tuberculosis*-infected *Socs3<sup>fl/fl</sup> LysM cre* mice. Sixteen days after infection, lungs from *Socs3<sup>fl/fl</sup> LysM cre* mice contained significantly lower levels of *Ifn- $\gamma$*  mRNA compared to control mice. At this time point, *Ifn- $\gamma$*  mRNA accumulation in lungs from *M. tuberculosis*-infected control mice was mediated by T cells because depletion of CD4<sup>+</sup> cells in control mice reduced IFN- $\gamma$  levels down to levels found in *Socs3<sup>fl/fl</sup> LysM cre* mice. Moreover, the differences in susceptibility to infection of SOCS3<sup>fl/fl</sup> *LysM cre* and control mice were abrogated under CD4<sup>+</sup> cell depletion.

Surprisingly, *Ifn- $\gamma$*  mRNA levels were only transiently reduced in *Socs3<sup>fl/fl</sup> LysM cre* mice, and no differences were detected 4 weeks after infection, although differences in bacterial levels at this time point were remarkable. However, a delayed recruitment of antigen-specific T cells to the lungs was shown to underlie increased bacterial burdens in susceptible mouse strains<sup>25</sup>. These results suggest that a delay of T cell-derived IFN- $\gamma$  secretion due to decreased IL-12p40 production in SOCS3-deficient DCs accounts for the increased susceptibility to *M. tuberculosis* infection of *Socs3<sup>fl/fl</sup> LysM cre* mice (Fig. 12).



**Figure 12: Role of SOCS3 in myeloid cells during *M. tuberculosis* infection.**

SOCS3 increases the secretion of IL-12 and TNF in *M. tuberculosis*-infected macrophages and DCs (1). Infected dendritic cells prime Th1 cells in the draining lymph node (2) and in the presence of SOCS3 we found earlier high levels of IFN- $\gamma$  in the lungs (3) activating infected cells.

In line with our results, the increased susceptibility of *Socs3<sup>fl/fl</sup> LysM cre* mice to *T. gondii* was due to reduced IL-12 responses that could be restored either by IL-6 neutralization or by administration of recombinant IL-12<sup>276</sup>. Since immune defense mechanisms to the intracellular *T. gondii* and intracellular *M. tuberculosis* resemble each other, it is not surprising that we found a similar setting during *M. tuberculosis* infection. Accordingly, in *T. gondii*-infected *Socs3<sup>fl/fl</sup> LysM cre* mice impaired IFN- $\gamma$  secretion by T and NK cells was found<sup>276</sup>, but unlike *M. tuberculosis* defense mechanisms that mainly rely on T cells, IFN- $\gamma$  produced by NK cells plays a crucial role in protection to *T. gondii*<sup>114,287,288</sup>.

Together with our data, this demonstrates a so far unappreciated role of IL-6 that, in the absence of SOCS3 in myeloid cells, becomes a potent antagonist of IFN- $\gamma$ -mediated protection.

*What is the role of SOCS3 in T cells during M. tuberculosis infection in vivo regarding the regulation of bacterial loads, cytokine responses and immunopathology?*

Mice with SOCS3-deficient T cells (*Socs3<sup>fl/fl</sup> lck cre* mice) displayed a dramatically increased susceptibility to *M. tuberculosis* infection. In comparison to *Socs3<sup>fl/fl</sup> LysM cre* mice, SOCS3 deficiency in T cells led to higher bacterial loads in the lungs and spleen, increased pathology, and earlier mortality. Transfer of SOCS3-deficient T cells in *Rag1<sup>-/-</sup>* mice did not convey resistance and suppressed protection by co-inoculated control T cells. Moreover, absence of SOCS3 in T cells in *Leishmania major*-infected mice increased parasite counts confirming a SOCS3-dependent protection mechanism<sup>258</sup>.

However, the observed susceptibility to *M. tuberculosis* seemed to be a pathogen-specific mechanism because detected bacterial numbers after BCG infection in *Socs3<sup>fl/fl</sup> lck cre* and control mice were similar. Surprisingly, BCG-vaccinated *Socs3<sup>fl/fl</sup> lck cre* mice showed the same protection to challenge with *M. tuberculosis* as control mice. This suggests that SOCS3 in T cells is not necessarily required for the resistance to *M. tuberculosis* if T cells are primed by BCG-immunization. In contrast to our results, *Socs3<sup>fl/fl</sup> lck cre* mice showed faster LCMV clearance than infected control mice demonstrating a very divergent function of SOCS3 depending on the pathogen<sup>270</sup>.

Since we detected higher levels of *Socs3* mRNA expression in *M. tuberculosis* than in BCG-infected lungs, we speculated that *Socs3* expression levels in T cells might be elevated during *M. tuberculosis* in contrast to BCG infection. Unexpectedly, we did not detect any upregulation of *Socs3* mRNA in pulmonary T cells neither after *M. tuberculosis* nor after BCG infection. Our data obtained from mycobacteria-infected mice is consistent with earlier observations that describe a down-regulation of SOCS3 in antigen-stimulated T cells in comparison to naïve T cells<sup>250</sup>. However, since the regulation of SOCS3 expression in T cells during BCG and *M. tuberculosis* infection is comparable, it does not explain the increased susceptibility to *M. tuberculosis* of *Socs3<sup>fl/fl</sup> lck cre* mice.

The immune responses of *Socs3<sup>fl/fl</sup> lck cre* mice were then investigated. We chose 2.5 weeks after *M. tuberculosis* infection as first examined time point as antigen-specific T cells were already present in the lungs, but bacterial loads in control and *Socs3<sup>fl/fl</sup> lck cre* mice were similar. We found that higher levels of IL-17 were secreted in response to antigen-specific stimulation of lung homogenates by *Socs3<sup>fl/fl</sup> lck cre* mice matching previously reported SOCS3-mediated inhibition of Th17 generation<sup>249,254,259</sup>. However, no differences in IL-17 secretion were detected after re-

stimulation of splenocytes from BCG-immunized *Socs3<sup>fl/fl</sup> lck cre* and control mice. Therefore, the increased IL-17 secretion and susceptibility to *M. tuberculosis* in *Socs3<sup>fl/fl</sup> lck cre* mice could be causally associated. The role of IL-17 in resistance to *M. tuberculosis* is controversial and the lack of IL-17 or IL-17R in a low dose model that is also applied in our experiments did not impair the ability of mice to control *M. tuberculosis* infection<sup>144,145</sup>. In contrast, high IL-17 levels may contribute to an increased bacterial dissemination and a neutrophil-mediated inflammation that may account for the early death and immunopathology of *M. tuberculosis*-infected *Socs3<sup>fl/fl</sup> lck cre* mice as described for other susceptible mouse strains<sup>55,289,290</sup>. In fact, increased neutrophil-derived molecules and necrosis could be observed in *M. tuberculosis*-infected *Socs3<sup>fl/fl</sup> lck cre* mice.

Moreover, IFN- $\gamma$  levels were elevated in *Socs3<sup>fl/fl</sup> lck cre* mice already 2.5 weeks after *M. tuberculosis* infection and might be explained by the described binding of SOCS3 to IL-12R $\beta$ <sup>257</sup>. However, increased IFN- $\gamma$  levels cannot justify the dramatic susceptibility to *M. tuberculosis* of *Socs3<sup>fl/fl</sup> lck cre* mice. Kinjyo *et al.* reported that SOCS3-deficient T cells show enhanced development of Th3 cells that produce anti-inflammatory cytokines such as IL-10 and TGF- $\beta$ <sup>258</sup>. Nevertheless, IL-10 levels were not increased in *Socs3<sup>fl/fl</sup> lck cre* mice.

Altogether, additional studies on IL-17 that is increased during *M. tuberculosis* but not during BCG infection in *Socs3<sup>fl/fl</sup> lck cre* mice may provide promising insights into its role during *M. tuberculosis* infection.

*Which function has gp130 signaling in SOCS3-deficient models in vivo and in vitro during mycobacterial infections?*

SOCS3 inhibits different signaling receptors. We investigated whether SOCS3 promoted resistance against *M. tuberculosis* via the regulation of gp130-dependent or independent receptor signals. We found that *gp130<sup>F/F</sup>* mice were highly susceptible to *M. tuberculosis* infection, which was at least in part caused by aberrant IL-6 and STAT3 responses since *gp130<sup>F/F</sup> Il-6<sup>-/-</sup>* and *gp130<sup>F/F</sup> Stat3<sup>+/-</sup>* mice displayed lower bacterial loads than *gp130<sup>F/F</sup>* mice. Cytokine responses of mycobacteria-infected *gp130<sup>F/F</sup>* BMM resembled *Socs3<sup>fl/fl</sup> LysM cre* BMM with decreased IL-12 and TNF levels compared to control cells. Previously, it had been demonstrated that *gp130<sup>F/F</sup>* BMM alike *Socs3<sup>fl/fl</sup> LysM cre* BMM induce reduced TNF and IL-12 levels in response to LPS and IL-6 compared to control BMM<sup>242</sup>. Moreover, we could demonstrate that

the reduction of cytokine concentration during mycobacterial infections was IL-6 dependent since IL-12 and TNF levels in *gp130<sup>F/F</sup>/Il-6<sup>-/-</sup>* BMM were restored to control levels. Accordingly, lower IL-12p40 expression levels were found in the lungs of *gp130<sup>F/F</sup>* compared to control mice, which were restored in *gp130<sup>F/F</sup>Il-6<sup>-/-</sup>* mice, implying that changed IL-6 signaling in the absence of SOCS3 inhibition of gp130 signaling may account for the increased susceptibility.

Taken together, we propose that, as in *Socs3<sup>fl/fl</sup> LysM cre* mice, delayed T cell priming due to impaired IL-12 induction contributes to the susceptibility to *M. tuberculosis* in *gp130<sup>F/F</sup>* mice. Similar results were obtained during *T. gondii* infection of *gp130<sup>F/F</sup>* mice that showed a transient reduction of IFN- $\gamma$  serum levels<sup>275</sup>. Treatment of *gp130<sup>F/F</sup>* mice with either neutralizing IL-6 antibodies or exogenous IL-12 improved parasite control demonstrating that SOCS3 provides similar protection mechanisms by modulation of gp130 signaling during *T. gondii* and *M. tuberculosis* infection<sup>275</sup>.

Interestingly, T cells from *gp130<sup>F/F</sup>* mice could transfer protection to *Rag1<sup>-/-</sup>* to the same extent as transferred control T cells, indicating that the protective SOCS3 expression in T cells is gp130-independent. Since the susceptibility of *gp130<sup>F/F</sup>* to *M. tuberculosis* was significantly higher than of *Socs3<sup>fl/fl</sup> LysM cre* and T cells were not involved, we suggest that non-hematopoietic cells may also contribute to SOCS3-mediated control to *M. tuberculosis* infection, a suggestion that remains to be tested.

Altogether, SOCS3 plays a critical and non-redundant role in myeloid and T cells during *M. tuberculosis* infection. Notably, mechanisms of resistance conveyed by SOCS3 are different in myeloid and T cells but in both cases necessary to enable a proper T cell differentiation.

## 5 SUMMARY

In summary, we define the following roles for SOCS proteins during *M. tuberculosis* infection:

### SOCS1

- ... reduces IFN- $\gamma$  induction in response to IL-12 in macrophages.
- ... inhibits IFN- $\gamma$ -mediated growth control during early infection.
- ... is a major controller of non-macrophage-mediated inflammation.

### SOCS2

- ... has no influence on mycobacterial growth control.

### SOCS3

- ... in myeloid and T cells reduces susceptibility to infection.
- ... in myeloid cells impairs IL-6 dependent reduction of IL-12 and TNF secretion and leads to early induction of IFN- $\gamma$  by CD4<sup>+</sup> cells in the lungs.
- ... in T cells hampers IL-17 and IFN- $\gamma$  secretion during *M. tuberculosis* infection.
- ... is redundant in T cells for *M. tuberculosis* protection if mice are BCG-vaccinated.
- ... mediates inhibition of gp130-transmitted signals and thereby reduces susceptibility that is dependent on IL-6 and STAT3.

## 6 CONCLUSIONS AND FUTURE PERSPECTIVES

Summarizing the studies on SOCS proteins presented in this thesis, I would like to give an evaluation of the role of the different SOCS proteins during mycobacterial infections and discuss under which circumstances a modulation of SOCS levels may be of importance in controlling the infection.

SOCS1 plays a two-faced role during mycobacterial infections. During the innate phase, when bacteria replicate in macrophages without any adaptive immunity present, SOCS1 suppresses IFN- $\gamma$  induction by macrophages and therefore enables mycobacterial growth. Deletion of SOCS1 could slow down this mycobacterial growth but did not lead to complete eradication of the bacteria. Since it has been proposed that a fraction of the *M. tuberculosis*-exposed individuals remain uninfected, it would be interesting to determine whether SOCS1 is involved in such innate control of infection. In contrast, SOCS1 expression protected from severe inflammation in the lungs without impairing immunity to infection during later stages of *M. tuberculosis* infection. Thus, SOCS1 mimetic peptides may be useful to limit immune-mediated pathology in tuberculosis as it was shown in a vaccinia virus infection model<sup>192</sup>.

Regarding our data on SOCS2, its role can be addressed as minor or redundant during *M. tuberculosis* infection. Although enhanced Th2 responses were reported in the absence of SOCS2, we observed similar bacterial burdens in presence and absence of SOCS2 suggesting a well functioning Th1 response. However, it is possible that under Th2-stimulating conditions, such as a common co-infection with helminths, absence of SOCS2 may further enhance Th2 responses and thereby impair *M. tuberculosis* control.

In contrast to SOCS1 and SOCS2, SOCS3 expression by either myeloid or lymphoid cells had a central role in controlling *M. tuberculosis* infection. Absence of SOCS3 in BMM has been described to prolong STAT3 activation in response to IL-6 that in turn decreased IL-12 and TNF transcription<sup>126</sup>. The knockdown of STAT3 in DCs has been shown to increase cytokine production, antigen-specific T cell activation and resistance to IL-10-mediated suppression<sup>291</sup>. Moreover, increased STAT3 activation prevents recognition of tumor cells by the immune system in different kinds of cancer and inhibition of STAT3 activation by administration of small inhibitory molecules has been shown to impair the growth of cancer cells *in vitro* and *in vivo*<sup>292</sup>. Therefore, it would be a promising approach to limit STAT3

activation to improve resistance to *M. tuberculosis* infection by increasing IL-12 levels and antigen-specific Th1 responses. However, since STAT3 is unique in its capacity to direct anti-inflammatory activities of IL-10 and mice with a knockdown for STAT3 in myeloid cells showed a pro-inflammatory phenotype with spontaneous development of colitis<sup>243</sup>, it will be a challenge to keep the balance between pro- and anti-inflammatory signals. Additionally, it has to be taken in consideration that a knockdown of *gp130* in myeloid cells did not improve control of *M. tuberculosis* infection in mice, demonstrating that ablation of all gp130-mediated signals may be contraindicated<sup>293</sup>. Even considering these aspects, our data convincingly argue for further explorations in targeting STAT3 to improve immune control during tuberculosis.

Since SOCS3 expression in T cells was not required to mount protective immunity after BCG-vaccination, it would be of high interest to understand why SOCS3 in turn is required for resistance to primary *M. tuberculosis* infection. Furthermore, the high IL-17 levels in the absence of SOCS3 in T cells may have contributed to the observed susceptibility since we did not detect any elevated anti-inflammatory immune responses that could account for increased bacterial growth. Understanding the role of these two factors, SOCS3 expression and elevated IL-17 secretion by T cells during *M. tuberculosis* infection, may provide important information how to achieve protection against *M. tuberculosis* and thereby how to improve vaccine design. Summarizing, we suggest that modulation of SOCS3-regulated pathways is a promising approach to improve vaccination and the control of mycobacterial infections.



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