

Published in final edited form as:

Trends Parasitol. 2013 December ; 29(12): . doi:10.1016/j.pt.2013.10.002.

Regulation of immunopathogenesis during *Plasmodium* and *Toxoplasma* infections: more parallels than distinctions?

Noah S. Butler^{1,*}, Tajie H. Harris^{2,*}, and Ira J. Blader³

¹Department of Microbiology and Immunology, University of Oklahoma Health Sciences Center, Oklahoma City, OK 73104, USA

²Department of Neuroscience, University of Virginia School of Medicine, Charlottesville, VA 22908, USA

³Department of Microbiology and Immunology, The University at Buffalo, The State University of New York, Buffalo, NY 14214, USA

Abstract

Toxoplasma and *Plasmodium* parasites exact a significant toll on public health. Host immunity required for efficient control of infection by these Apicomplexans involves the induction of potent T cell responses, which sometimes results in immunopathological damage. Thus, protective immune responses must be balanced by regulatory networks that limit immunopathology. We review several key cellular and molecular immunoregulatory networks operational during *Toxoplasma* and *Plasmodium* infections. Accumulating data show that despite differences in how the immune response controls these parasites, many host immunoregulatory pathways and cellular networks are common to both. Thus, understanding the cellular and molecular circuits that prevent or regulate immunopathological responses against one parasite is likely to inform our understanding of the host response to the other parasite.

Keywords

Plasmodium; *Toxoplasma*; immunopathology; IL-10; IL-27; TGF- β

Protective immunity and immunopathology after *Plasmodium* or *Toxoplasma* infection

Plasmodium and *Toxoplasma* represent two of the most prevalent and successful parasites. Reasons for this include the complex life cycle of each parasite and a limited understanding of the interplay between the parasites and host immune response. Although these organisms infect different tissues and cause distinct patterns of disease (Boxes 1 and 2), one feature common to both parasites is that some disease manifestations are directly linked to the highly inflammatory nature of the host immune response (Box 3). Moreover, hosts that lack key immunoregulatory molecules, cell types, or pathways cannot control parasite growth and succumb to lethal immunopathology [1–3]. Thus, several manifestations of malaria and toxoplasmosis are likely to be a consequence of the highly inflammatory nature of the innate and T cell mediated immune responses triggered during the acute phases of infection that develop to limit parasite replication.

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Corresponding authors: Butler, N.S. (noah-butler@ouhsc.edu); Harris, T.H. (tjiharris@virginia.edu).

* These authors contributed equally to this work.

Box 1**Induction of cell mediated immunity after *Plasmodium* infection**

Plasmodium infection begins with mosquito deposition of sporozoites in the mammalian dermis. Motile sporozoites enter the circulation, passively transit to the liver, and initiate an asymptomatic period of differentiation in hepatocytes. *Plasmodium* merozoites are released from hepatocytes and subsequently infect host erythrocytes. The blood stage of *Plasmodium* infection is responsible for all clinical symptoms of malaria. During this phase, asexual replication of merozoites in erythrocytes stimulates potent, highly inflammatory immune responses [76]. Early activation of host immunity is associated with accumulation of parasite-infected erythrocytes in the spleen. There, innate immune cells including inflammatory monocytes, macrophages, DCs, NK cells, and $\gamma\delta$ T cells release several proinflammatory cytokines and pyrogens, including LT- α , TNF- α , IL-1, IFN- γ , and IL-6 (See Figure 1 in main text) [77]. IL-12-mediated induction of highly activated, parasite-specific CD4 T cells expressing IFN- γ (Th1) is also central to protection against blood stage *Plasmodium* infection [78–81].

Box 2**Induction of cell mediated immunity after *Toxoplasma gondii* infection**

Human infection with *T. gondii* results from the ingestion of oocysts from the environment, the ingestion of tissue cysts from infected animals, or through vertical transmission of parasites from infected mothers to their fetus [82]. Once digested, parasites rupture from the cyst, infect intestinal cells where they transform into tachyzoites, and trigger the recruitment of numerous leukocytes including monocytes and DCs [52]. The parasite can also infect phagocytes and use them to initiate their dissemination to a wide variety of tissues including immune-privileged sites such as the brain or retina [83]. In the tissue, the parasite converts from the tachyzoite form to the slowly replicating bradyzoite form that resides within tissue cysts. Bradyzoites periodically reactivate to rapidly replicating tachyzoites, and an immune response must be mounted to control the reactivated infection [82]. Resistance to *T. gondii* in both the gut and CNS involves innate immune activation coupled with the development of highly polarized T cell responses necessary to limit parasite survival and persistence [84]. Initial recognition of parasites by APCs triggers the expression of chemokines and inflammatory cytokines including IL-12, IL-6, and TNF- α . Recent studies have shown that CD8⁺ DCs are the critical source of IL-12 during *T. gondii* infection [85]. IL-12 polarizes CD4 helper cells towards the Th1 lineage [86] and along with other inflammatory cytokines, such as IL-18 and IL-1, can further amplify inflammation by stimulating the release of IFN- γ by NK cells [87,88].

Box 3**Inflammation and immunopathology during toxoplasmosis and malaria**

T. gondii and *Plasmodium* parasites activate innate phagocytic cells via interactions between parasite-expressed pathogen-associated molecular patterns (PAMPs) and pathogen recognition receptors (PRRs) on monocytes, macrophages, and DCs. Appropriately activated phagocytes respond by secreting proinflammatory cytokines (e.g., IL-12, TNF- α , IL-6, and IL-1), engulfing parasites or parasite-infected host cells and migrating to regional draining lymph nodes or the spleen [88–92]. Local cytokine production also activates other innate immune cells, such as NK cells and $\gamma\delta$ T cells

[93,94]. In secondary lymphoid tissues, proinflammatory cytokine expression continues, and antigens from the parasites are processed and presented to CD4 and CD8 T cells, which triggers T cell activation and proliferation [81,95,96]. Specific cytokines, notably IL-12, induce activated and proliferating CD4 T cells to differentiate into potent Th1 effector cells that secrete copious amounts of IFN- γ [86,88,97,98]. High levels of IFN- γ potentiate macrophage activation by driving the production of TNF- α , reactive oxygen species (NO $^-$, O $_2^-$), and chemokine expression [99,100]. However, sustained high levels of IFN- γ are also associated with acute illness via the further induction and release of TNF- α , IL-6, and other pyrogenic factors responsible for fever, suppression of hematopoiesis, and anemia [27,101]. Th1 cells and proinflammatory cytokines are additionally associated with activation of brain endothelial cells and subsequent recruitment and activation of pathological CD8 T cells in rodent models of ECM [37,56,77,102]. During toxoplasmosis, local proinflammatory cytokine release can trigger apoptosis or necrosis of normal tissue. Systemically, proinflammatory cytokines trigger fever and alter hematopoiesis. Immunopathology is clinically evident during acute toxoplasmosis when parasites infect the gut and in cases of toxoplasmic retinochoroiditis, where unresolved inflammation threatens vision [103–105].

Clinical and experimental study of these two Apicomplexan parasites has provided critical insight into basic cellular and molecular circuits that regulate immunopathogenesis. We highlight the parallel pathways of immunoregulation that are operational after *Plasmodium* and *Toxoplasma gondii* infections. We focus on inhibitory receptors, regulatory cytokines, and functionally distinct immune cell subsets (Figure 1). Insight gained from the study of one parasite infection is likely to shed light on mechanisms of immunoregulation during infection by the other, as well as reveal fundamental insight into the biology of immunoregulation. A better understanding of the molecular and cellular factors that regulate immunopathogenesis should aid in the identification of novel opportunities to intervene and improve health outcomes after these or other microbial infections.

Immunoregulation by cell surface inhibitory receptors

After acute microbial infection or vaccination, professional antigen presenting cells (APCs) capture and present antigen to naïve T cells in secondary lymphoid tissues. Appropriately activated T cells undergo clonal expansion and transiently express an array of inhibitory receptors including cytotoxic T lymphocyte antigen-4 (CTLA-4) and programmed death-1 (PD-1). Inhibitory receptors counterbalance exuberant T cell activation and are essential for preventing immunopathologies. The ligands for inhibitory receptors and the mechanisms by which these receptors suppress T cell activity vary. For example, CTLA-4 out-competes CD28 (an activating receptor) for binding B7 family member ligands (CD80 or CD86) expressed by APCs. Thus, sequential expression of CD28 followed by CTLA-4 results in a shift from activating to inhibitory signals in effector T cells. PD-1 is a second inhibitory receptor, its ligation by CD80, programmed death ligand 1 (PD-L1), or PD-L2 on APCs, or PD-L1 on non-hematopoietic cells attenuates T cell receptor signaling. Inhibitory receptors are generally downregulated during the transition from effector to memory T cells, whereas sustained expression can lead to their functional impairment or ‘exhaustion’ [4]. Functional T cell exhaustion has been reported during both prolonged *Plasmodium* [5] and chronic *T. gondii* [6] infection in mice (reviewed in [7]).

Multiple reports highlight the critical roles for CTLA-4 and PD-1 in preventing immunopathology during acute *Plasmodium* and *Toxoplasma* infection. In *Plasmodium berghei*-infected mice, blockade of CTLA-4 enhances central nervous system (CNS) and liver immunopathology in experimental cerebral malaria (ECM)-susceptible C57BL/6 mice

[8], and blockade of CTLA-4 and PD-1 triggers the development of interferon-gamma (IFN- γ ; see Glossary) and T cell mediated, ECM-like disease in normally resistant BALB/c mice [9–11]. Furthermore, CTLA-4 expression by regulatory CD4 T cells (Tregs) can also limit effector T cell activation and pathology in ECM-susceptible mice [12]. During experimental ocular toxoplasmosis, intravitreal delivery of *Toxoplasma* tachyzoites triggered IFN- γ -dependent upregulation of major histocompatibility complex (MHC) class II and PD-L1 on infiltrating hematopoietic and resident retinal cells. Importantly, infiltrating CD4 T cells from *Toxoplasma*-infected retinas expressed high levels of PD-1, and retinal cells suppressed CD4 T cell activation in a PD-L1-dependent manner [13]. These reports illustrate the important roles that CTLA-4 and PD-1 play in T cell activation and immunopathology in toxoplasmosis and malaria. Future work should address the factors that control the expression of CTLA-4, PD-1, and other inhibitory receptors during parasitic infections. It is also of interest to determine which factors influence the cellular, temporal, or spatial patterns of inhibitory receptor–ligand expression, and ultimately whether this information can be exploited to regulate T cell mediated inflammation.

Immunoregulation by secreted factors

Interleukin-10 (IL-10)

IL-10 is a pleotropic cytokine that acts to suppress the activity of several immune cell types, including APCs, B cells, and T cells (Table 1). Initially characterized as a cytokine expressed by Type 2 T helper cells (Th2), it is now clear that most lymphocyte subsets [natural killer (NK) cells, CD8 T cells, and Th1, Th17, and Tregs] and several myeloid-derived cells [including macrophages, dendritic cells (DCs), and neutrophils] express IL-10 [14]. The finding that IL-10-deficient mice develop spontaneous immunopathology due to T cell responses directed against endogenous gut flora revealed the essential role for IL-10 in maintaining immune homeostasis [15]. Subsequently, IL-10 was shown to prevent immunopathology after infection with several pathogens, including *Toxoplasma* and *Plasmodium* [2,3,16–21].

The earliest and clearest example of the importance of IL-10 in limiting immunopathogenesis after microbial infection was shown by infection with *Toxoplasma* in IL-10-deficient mice. Compared with wild type (WT) mice, IL-10-deficient mice had greater Th1 responsiveness, elevated IFN- γ levels, and superior control of *Toxoplasma* growth. Despite better parasite control, these mice had more severe disease and died from fulminant immunopathology. Importantly, depletion of CD4 T cells rescued *Toxoplasma*-infected IL-10-deficient mice demonstrating that CD4 T cells are responsible for immunopathological responses [2]. Although an initial report suggested that follicular B cells were an important source of IL-10 in *Toxoplasma*-infected mice [22], subsequent work demonstrated that IFN- γ -expressing Th1 effector cells were the major source of IL-10 during infection and that IL-10 acted by antagonizing the expression co-stimulatory molecules and IL-12 by APCs [18,19]. This was an important shift in the view of immunoregulation by IL-10, because the long-standing model suggested that effector T cells responsible for immunopathology or parasite clearance did not express regulatory cytokines. Identifying additional cellular sources of IL-10 and determining whether IL-10 expression by Th1 cells is regulated by APCs or other cytokines remain areas of intense investigation.

In mice acutely infected with *Plasmodium chabaudi*, Th1 effectors also appear to be a critical source of IL-10 that limits immunopathology [20,21]. Studies using specific ablation of IL-10 expression in B cells, myeloid cells, or T cells in combination with adoptive transfer of WT or IL-10-deficient Tregs elegantly showed that highly activated Th1 effectors, but not Tregs, were essential sources of IL-10 that minimized weight loss, hypothermia, and anemia in mice [23]. Notably, the absence of IL-10 did not improve

parasite clearance in these animals, suggesting that the effects of IL-10 do not include suppression of protective immunity against *P. chabaudi*. By contrast, IL-10 in *Plasmodium yoelii*-infected mice acted to both limit immunopathology and to impede the induction and maintenance of highly activated T cells necessary for efficient parasite clearance [17]. In that report, the majority of IL-10 was produced by an infection-induced population of Foxp3-negative Tregs, termed Tr1 cells, which are distinct from Foxp3-expressing natural and peripherally induced Tregs. Thus, depending on the *Plasmodium* species and rodent host, both the source of IL-10 and its impact on immunopathogenesis and parasite control can be fairly distinct.

Consistent with data from experimental rodent model studies, data from studies on humans infected with *Plasmodium falciparum* indicate that IL-10 may have variable roles. For example, significant correlations between increased serum IL-10 levels and reduced risk of developing severe malarial anemia have been established [24], whereas other studies have either failed to identify links [25] or demonstrated that high IL-10 levels correlate with higher parasitemia or worse clinical outcomes [26–28]. Recent data also showed that sustained circulating levels of IL-10 in *P. falciparum*- or *Plasmodium vivax*-infected individuals correlated with reduced frequencies of mature DCs. Decreased numbers of DCs were due to DC apoptosis, and *in vitro* blockade of IL-10 inhibited DC apoptosis [29]. IL-10-mediated DC apoptosis represents a novel pathway of immunoregulation during *Plasmodium* infection; earlier data from experimental *Plasmodium* infection suggested that reduced DC survival was associated with direct interactions between DCs and infected erythrocytes [30]. Thus, these new data show that the loss of circulating mature DCs can largely be a consequence of the immune response directed against the parasite, rather than a direct effect of the parasite on the survival or maintenance of mature DCs.

Transforming growth factor- β (TGF- β)

TGF- β is expressed by and acts upon a diverse range of cells and tissues (Table 1). In the immune system, TGF- β primarily acts to suppress effector T cells and promote Treg maintenance and differentiation [31,32]. In this context, TGF- β has proven essential for regulating T cell mediated inflammation and maintaining immune homeostasis and tolerance. Given its critical role in regulating inflammatory Th1 cells and cytotoxic CD8 T cells, it is not surprising that TGF- β is linked to regulating the immunopathogenesis of both *Plasmodium* and *Toxoplasma* infections.

TGF- β was initially implicated as a regulator of *Plasmodium* infection-induced inflammation in studies comparing *P. berghei* infection in ECM-resistant and -susceptible strains of mice [33]. In these studies, analyses of cytokine profiles revealed that ECM-susceptible mice had markedly reduced expression of TGF- β , suggesting that it might be critical for maintaining the balance between T cell mediated protection and immunopathology. Circulating TGF- β is also a key correlate of improved outcomes during severe malarial anemia and cerebral malaria (CM) in *P. falciparum*-infected humans [34,35]. Although the precise pathways are unknown, development of CM is associated with proinflammatory cytokine storms that include release of lymphotoxin- α (LT- α), IL-6, and tumor necrosis factor- α (TNF- α), as well as the expansion of cytotoxic CD8 T cells and inflammatory Th1 cells [36–38]. Thus, suppression of effector T cell activation is probably one way that TGF- β regulates and/or reduces the severity of CM. A recent report demonstrated that TGF- β limits effector T cell survival via suppression of the antiapoptotic molecule Bcl-2 [39]. Future studies aimed at determining the precise contribution of TGF- β to regulating T cell mediated immunopathology during malaria could reveal potential avenues for therapeutic intervention during severe *Plasmodium* infections.

During *Toxoplasma* infection of the CNS, neuronal cells directly respond to IL-6-driven inflammation by expressing the regulatory cytokines TGF- β and IL-27. When IL-6 signaling was specifically ablated in neurons by deleting a component of the IL-6 receptor (gp130), mice failed to express either TGF- β or IL-27 at the site of infection and died as a result of an encephalitis associated with Th1 and Th17 cell accumulation in the CNS [39,40]. Because gp130 is also a component of a functional IL-27 receptor, the role of IL-27 in this study remains a query. An essential immunoregulatory role for TGF- β has also been described during intestinal *Toxoplasma* infection. Resident gut intraepithelial CD8 T cells were the major source of TGF- β , the expression of which was essential for reducing *Toxoplasma*-induced inflammation and mucosal immunopathology, and for maintaining homeostasis in the intestine [41].

These examples highlight the similar and important functions of TGF- β in regulating immunopathogenesis after *Plasmodium* or *Toxoplasma* infection. Moreover, these studies reveal that the cellular sources of TGF- β during these infections can be diverse. In addition to understanding how TGF- β expression is regulated, it is important to identify the cellular targets of TGF- β as well as define how the inflammatory environment influences the activity and effects of TGF- β . For example, studies have shown that in the presence of IL-6, TGF- β triggers inflammatory Th17 differentiation instead of suppressing effector T cell activity or promoting Treg differentiation [42]. Thus, TGF- β can function via complicated circuits to integrate environmental cues to both suppress T cell activity and potentiate inflammation.

Interleukin-27

IL-27 is a member of a family of cytokines that includes IL-6 and IL-12, two factors associated with inducing proinflammatory immune responses during infection with either *Plasmodium* or *Toxoplasma*. Early studies demonstrated that IL-27 could skew the differentiation of naïve CD4 T cells towards a Th1 functional profile via the induction of the key Th1 transcription factor, T-bet [43]. Thus, IL-27 was originally thought to function solely as a proinflammatory cytokine. More recently, IL-27 has been linked to functional suppression of effector CD4 T cells, including Th1, Th2, and Th17 cell populations [44]. The suppression of Th2 and Th17 cell activity is linked to the ability of IL-27 to antagonize IL-2 production [45], a cytokine necessary for proliferation and survival of antigen-specific T cells during an immune response.

The specific mechanisms by which IL-27 suppresses Th1 responses are also becoming clear; in mice, Th1-driven intestinal immunopathology that develops in the absence of IL-27 appears linked to reductions in unique Treg populations [46]. Oral high-dose *Toxoplasma* infections were associated with expansions of phenotypically and functionally distinct Tregs that also expressed molecules associated with effector Th1 cell trafficking and function, including CXCR3 and T-bet. After adoptive transfer, these T-bet⁺ CXCR3⁺ Treg populations appeared to act most efficiently in the gut, and their suppression of Th1-associated immunopathology was functionally linked to their ability to produce IL-10 [46]. Thus, IL-27 can induce highly specialized Tregs that limit *Toxoplasma*-induced immunopathology in a tissue-specific manner. Several important questions remain regarding IL-27-mediated immunoregulation during intestinal *Toxoplasma* infection, including the cellular source(s) of this regulatory cytokine and whether Tregs require signals from cytokines other than IL-27 to acquire and exert their suppressive activity.

Plasmodium infection has also been linked to the induction and activity of IL-27. Mice lacking a functional IL-27 receptor were highly sensitive to *Plasmodium*-induced immunopathology [46,47]. Despite the ability of these mice to efficiently control parasite replication, IL-27 receptor deficiency resulted in dysregulated Th1 responses, systemic inflammation, immunopathology, and disruption of normal liver physiology. The

mechanisms underlying the ability of IL-27 to regulate Th1 inflammatory T cells during *Plasmodium* infection are just now becoming clear. In contrast to its ability to induce specific populations of Tregs in the *Toxoplasma*-infected gut, IL-27 signaling in effector T cells during blood stage *Plasmodium* infection appears to suppress inflammatory Th1 cell responsiveness to specific chemokines and cytokines [48,49]. A recent report showed that IL-27 signaling in highly polarized Th1 cells led to marked reductions in the expression of the CCR5 chemokine receptor. The authors contend that dysregulation of CCR5-dependent T cell chemotaxis contributes to the ability of IL-27 to suppress inflammatory Th1 T cell migration to the spleen during *Plasmodium* infections [49]. Moreover, a subsequent study examining pathogenic Th1 cells induced by *P. berghei* infection found that IL-27 limited T cell responsiveness to IL-12 [48], which is a critical positive regulator of Th1 cell differentiation and survival as discussed above.

Through a variety of experimental models, these results highlight the importance of IL-27 in regulating Th1-mediated immunopathogenesis during infections with either *Toxoplasma* or *Plasmodium* and influencing the outcome of disease. In addition to inducing specific populations of Tregs or limiting responsiveness to cytokines or chemo-kines, IL-27 is also a well-known trigger of IL-10 expression by T cells [50,51]. For example, the generation of IL-10-expressing effector T cells during *P. chabaudi* infection requires IL-27 [23]. By contrast, a recent report showed that IL-27 is also important for IL-10-independent protection against immunopathology in *P. berghei*-infected mice [47]. The heterogeneous patterns of expression and responsiveness to IL-27 *in vivo* underscore the complexity of immunoregulation by this cytokine.

Immunoregulation by myeloid cells

Bone marrow-derived myeloid precursors differentiate into several functionally distinct innate immune cells including neutrophils, eosinophils, basophils, and monocytes. Significant interest in understanding the function and regulation of monocytes has grown from early reports showing that these cells are essential for control of both *Toxoplasma* and *Plasmodium* [52–54]. After infection, monocytes traffic from the blood to inflamed tissues where they can further differentiate into phagocytic macrophages and DCs. Thus, monocytes contribute to resistance by clearing pathogens through phagocytosis, releasing proinflammatory cytokines, and providing a pool of APCs to aid in promoting T and B cell responses.

In addition to these protective inflammatory responses, infiltrating monocytes and bone marrow-derived myeloid cells play immunoregulatory roles during both toxoplasmosis and malaria. For instance, intestinal infection by specific strains of *Toxoplasma* is associated with an influx of inflammatory monocytes that adopt regulatory properties characterized by IL-10 and prostaglandin E2 (PGE2) expression. Unexpectedly, IL-10 and PGE2 were expressed in response to infection with commensal bacteria but not other inflammatory cues such as *Toxoplasma* lysates, and in the absence of these secreted regulatory factors mice develop neutrophil-dependent intestinal pathology [55]. Thus, monocytes respond to diverse signals to regulate *Toxoplasma*-induced immunopathology.

Similarly, *Plasmodium* infections are associated with antimalarial, monocyte-driven inflammatory cascades [54]. These protective activities, which include phagocytosis of infected erythrocytes and secretion of cytokines, have been recently reviewed [56]. By contrast, *Plasmodium* blood stage infection has been linked to the accumulation of distinct subsets of myeloid cells that exhibit potent regulatory functions. In rodents infected with blood stage *Plasmodium* parasites, a striking inversion occurs in the ratio of proinflammatory DCs and regulatory DCs that express IL-10 themselves and also induce

IL-10 expression in CD4 T cells [57]. Finally, *in vitro* studies have shown that interactions between circulating monocytes and *P. falciparum*-infected erythrocytes can trigger Treg differentiation. Notably, the *in vitro* induction of these Tregs was associated with the capacity of monocytes to secrete IL-10 and TGF- β [58,59]. These latter observations are consistent with data showing that splenic inflammatory monocytes express IL-10 during acute *P. chabaudi* infection in mice [54]. Further study is required to determine whether such cellular interactions and regulatory pathways occur in *P. falciparum*-infected individuals. Given the critical role of IL-10 in regulating immunity during *Plasmodium* infection, these data suggest that infiltrating monocytes and resident myeloid cells can also function within regulatory networks that act to prevent or limit immunopathology.

The precise origins and relationships between parasite-induced myeloid cells exhibiting inflammatory or regulatory functions are not entirely clear, and in each of the examples cited above the regulatory function of monocytes was dependent on local cytokine, chemokine, or tissue microenvironments. Thus, it will be necessary to not only determine developmental relationships between inflammatory and regulatory myeloid cells but to also understand how specific microenvironments shape their differentiation or function. Such information could reveal novel strategies for regulating pathological inflammatory responses during malaria or toxoplasmosis.

Immunoregulation by lymphoid cells

T regulatory cells

Multiple subsets of CD4 T cells exhibiting regulatory function have been described, including Foxp3⁺ natural Tregs (nTreg) that arise from the thymus, Foxp3⁺ activation-induced Tregs (iTreg) that develop in the periphery during immune activation, and Foxp3-negative Th1 cells expressing IL-10 (Tr1) that also develop after peripheral activation [60]. In addition to Foxp3 expression, Tregs are also typified by expression of the high-affinity IL-2 receptor, CD25. nTreg and iTreg function via release of soluble factors, such as IL-10 and TGF- β , and via direct cell–cell contact involving inhibitory molecules such as CTLA-4 or the glucocorticoid-induced TNF receptor (TNFR) family related gene (GITR) [60,61], whereas the suppressive function of Tr1 cells appears limited to IL-10 and TGF- β secretion. In addition to their critical role in maintaining immune homeostasis and preventing autoimmunity, Tregs have become an important contributor in controlling immunopathogenesis during microbial infections [62].

The appearance and expansion of Tregs has been described during infection with either *Toxoplasma* or *Plasmodium*. For instance, malaria has been linked to a higher ratio of circulating Tregs relative to effector T cells [63]; however, their relative contribution to parasite control and disease pathogenesis is not fully understood and remains controversial. Some studies have shown that Tregs paradoxically correlate with enhanced immune-mediated clearance of parasites and protection against pathological inflammatory responses [64,65]. By contrast, other studies have reported that Treg suppression of protective immunity correlated to higher parasite burdens [65]. Similarly, conflicting results have been observed in rodent models (primarily murine ECM models), in which mechanistic dissection of the role of Tregs is possible (reviewed in [66,67]). Thus, the biological relevance of Treg expansions in humans and other species remains an unanswered question.

Inconsistencies surrounding the role of Tregs in modulating immunity against *Toxoplasma* are also apparent. Multiple studies reported that the depletion of Tregs using the anti-CD25 monoclonal antibody (clone PC61) in rodents revealed limited roles for these cells during the chronic phases of infection [68]. Conversely, adoptive transfer of FoxP3⁺ Tregs was shown to potentially limit Th1-driven immunopathology in mice acutely infected with

Toxoplasma [46]. Yet, DCs and effector Th1 cells were shown to simultaneously impair Treg suppressive function and induce Treg differentiation into Th1-like effector cells that contribute to immunopathogenesis via secretion of IFN- γ [69]. Further adding to this debate, recent data also suggest a harmful role for Tregs via their ability to constrain protective immunity; rapid proliferative expansion of Th1 effector cells after *Toxoplasma* infection is reportedly linked to a transient decrease in Tregs [70]. Mechanistically, this effect was attributed to Treg deprivation of IL-2, because exogenous IL-2 could restore Treg numbers, which suppressed Th1 effector responses and resulted in loss of parasite control. Thus, a transient reduction in Tregs appears necessary for optimal Th1 effector responses against *Toxoplasma* [70]. This latter study demonstrates that the numerical expansion and suppressive activity of Tregs, if not restrained, can significantly limit protective immunity after acute infection. Furthermore, the treatment of chronically infected mice with IL-2 complexes leads to increased cyst burdens in the CNS, suggesting that increased numbers of Tregs can limit protection in the CNS [69,70].

Tregs appear to play a role in either limiting immunopathogenesis or controlling antiparasitic immunity during *Toxoplasma* and *Plasmodium* infections. Of key importance will be defining how or whether local secretion of regulatory cytokines in specific tissues and microenvironments influences disease outcomes. For example, it is of interest to determine the role of Tregs and the contribution of their suppressive effector molecules in the CNS of *Toxoplasma*- and *Plasmodium*-infected hosts. *Toxoplasma* can breach the gut and disseminate throughout the host, including the eyes and brain. Within these immune-privileged tissues, highly activated T cells and IFN- γ are required to limit parasite replication. Similarly, accumulation of Th1 cells in the microvasculature of the CNS after *Plasmodium* infection is linked to the development of ECM. Thus, it is essential to tightly regulate these polarized cellular reactions in these critical tissues to prevent immune-mediated pathology. Tregs could act in numerous ways to balance protection and pathology; it is possible that Tregs suppress antigen presentation in *Toxoplasma*-infected neural tissues, the vasculature of the CNS during CM, or limit the local activity of effector T cells. Dissecting these mechanisms *in vivo* will require the use of powerful technologies, including intravital microscopy, to identify and study cellular interactions that may prove critical for understanding the regulation of immunopathogenesis during *Toxoplasma* infections. Finally, it is also of interest to define the roles of nTregs and iTregs, and determine whether true parasite-specific iTregs are expanded after *Toxoplasma* or *Plasmodium* infection. Given the critical roles for Tregs to both limit immunopathology and constrain protective immunity after infection, modulation of the number, localization, or function of Tregs may hold promise as interventional immune-based strategies for toxoplasmosis or malaria.

B regulatory cells

Another regulatory immune cell subset that has been the focus of attention recently is the B regulatory (B10) cell, so-called because of its propensity to regulate effector CD4 T cell activity through the secretion of IL-10. B10 cells are phenotypically defined as CD19⁺CD1d^{hi}CD5⁺. CD1d is a non-classical MHC class I molecule necessary for presentation of lipid antigens, and CD5 has been functionally linked to antagonizing both T cell and B cell receptor signaling [71,72]. Originally defined as being critical regulators of Th1-driven inflammation in autoimmune disease [73], B10 cells are now a focus of attention as important regulators of inflammation during infection. Recent data show that B10 cells numerically expand and modulate immunity during parasitic infections, including babesiosis and schistosomiasis [74,75]. Follicular (B-2) B cell derived IL-10 was reported as a potent inhibitor of host immunity during *Toxoplasma* infection [22], although the precise ontogeny of B10 cell development is unknown. Even though no formal reports of B10 cell expansion during either *Plasmodium* or *Toxoplasma* infection exist, given the critical mutual

counterbalance between inflammatory Th1 cells and IL-10-mediated immunoregulation reported for these two Apicomplexan infections, it would not be surprising if B10 cells play a functional role.

Concluding remarks

During *Toxoplasma* and *Plasmodium* infections, several overlapping immunoregulatory pathways maintain the balance between health and disease. Indeed, the potent immune responses that serve to limit parasite persistence within the host can also be responsible for local or systemic pathologies associated with these infections. Multiple regulatory factors can independently and coordinately act to limit immunopathology during the acute stages of toxoplasmosis and malaria. The striking regulatory potential of monocytes is now appreciated, and the developmental relationships, similarities, and functional distinctions between naturally occurring and peripherally induced Tregs is growing clearer. Secreted factors, such as IL-10, TGF- β , and IL-27, are known potent regulators of immunity, and data from experimental models and human clinical studies highlight the critical and complex role that these cytokines play in regulating immunity against *Toxoplasma* and *Plasmodium*. Finally, several cell surface-expressed inhibitory receptors are critical for limiting T cell mediated immunopathology during ocular toxoplasmosis and ECM.

Although much information exists regarding the ability of these cellular and secreted factors to regulate immunopathology after infection, many questions remain (Box 4). Regulatory cytokines and inhibitory receptor ligands are expressed by and act on a diverse range of cell types, underscoring the complexity of these regulatory circuits. TGF- β can both promote and prevent immunopathology, depending on whether IL-6 and specific T cell subsets are present, and IL-27 has the capacity to either potentiate or inhibit Th1 responses. Whether or how Tregs modulate inflammatory effector T cells in local tissue environments, such as the eye or brain, during toxoplasmosis, remains unknown. Understanding how these cytokines are temporally regulated after infection, identifying the specific cellular sources, and determining whether the anatomy of interactions between inflammatory and regulatory cell subsets determines disease outcomes remain important and unanswered questions.

Box 4

Outstanding questions

- Much information has been learned about these regulatory cells, circuits, and pathways from experimental models, but which features of immunoregulation are operational during clinical malaria or acute ocular or intestinal toxoplasmosis?
- What are the key cellular sources for secreted regulatory factors such as IL-27, IL-10, and TGF- β , and do the key cellular sources of regulatory cytokines temporally shift as infection progresses?
- How does the tissue microenvironment influence the expression of regulatory cytokines and inhibitory receptor ligands, or the manner in which they exert suppressive effects on target cells?
- Are parasite-specific Tregs induced during toxoplasmosis or malaria, and what are the critical signals driving their activation, differentiation, or proliferation?
- Do monocytes exhibit specific regulatory functions before differentiating into macrophages or DCs in *Toxoplasma*-infected tissues or the spleen of *Plasmodium*-infected hosts, and what are the signals that stimulate the acquisition of suppressive function by infiltrating monocytes?

- Given that both IL-6 and TGF- β are coexpressed after infection with either *Toxoplasma* or *Plasmodium*, why is there so little evidence for an immunopathological role for Th17 cells during toxoplasmosis or malaria?

Rodent models of *Toxoplasma* and *Plasmodium* infection have been extensively used to dissect host–pathogen interactions and pathways of immunoregulation during acute and chronic protozoan infections. The power and utility of these models relate to the numerous *Toxoplasma* genotypes and rodent-specific species of *Plasmodium* parasites available for study, as well as the vast number of strains and genotypes of susceptible and resistant laboratory rodents. The use of multiple independent models has revealed important information about immunoregulatory pathways during these infections. Depending on the model system, the contribution of a given immunoregulatory factor can be different. For example, the immunoregulatory biology of IL-27 may be fairly distinct when expressed in the *Toxoplasma*-infected intestine and the spleen of *Plasmodium*-infected mice. Although unique experimental systems can reveal seemingly disparate mechanisms of action for these regulatory cytokines and cellular subsets, the mass of accumulating data will ultimately translate into a more comprehensive understanding of the biology of immunoregulation. A more complete understanding of the contribution of regulatory pathways to both limiting immunopathology and impeding potent immunity against *Toxoplasma* and *Plasmodium* infections will help shape future treatment and therapy against these and other infections.

Acknowledgments

The authors acknowledge members of their laboratories for their helpful discussions and Dr. Kristina Wasson-Blader for editorial assistance. We also offer apologies to the many investigators whose contributions we were unable to discuss owing to space limitations. Work in the Butler laboratory is supported by grants from the National Institutes of Health (NIH, AI099070) and the American Heart Association (13BGIA17140002). Work in the Blader laboratory is supported by grants from the NIH (AI069986 and EY021259).

Glossary

Bradyzoites	slow dividing form of <i>Toxoplasma</i> that encysts in tissues of mammalian hosts
CCR5	cell surface expressed receptor for chemokines CCL4 and CCL5
Chemokine	secreted factor that mediates chemotaxis of cells expressing appropriate receptors
CXCR3	cell surface receptor for chemokines CXCL9, CXCL10, and CXCL11
Cytotoxic T cells	T cell subsets characterized by expression of the CD8 coreceptor, restricted to association with antigen/MHC class I complexes, and critical for protective immunity against intracellular pathogens via induction of apoptosis in pathogen-infected cell targets
Dendritic cells (DCs)	key antigen presenting phagocytic cells responsible for activating naïve T cells and initiating cellular adaptive immunity
Effector T cells	recently activated T cells that have acquired the capacity to traffic to sites of infection and exhibit potent antimicrobial activity via cytokine secretion or cytolysis of infected cells

Foxp3	transcription factor essential for the differentiation and maintenance of natural and subsets of peripherally induced T regulatory cells
$\gamma\delta$ T cells	a small subset of T cells that expresses a T cell receptor composed of γ and δ chains. $\gamma\delta$ T cells are found in mucosal tissues and contribute to innate and adaptive responses
Helper T cells	T cell subsets characterized by expression of the CD4 coreceptor and restricted to association with antigen/MHC class II complexes. These cells are critical for the activation of macrophages, orchestration of antibody-secreting B cell responses, and regulation of immunity via their differentiation into one of several functionally distinct subsets (e.g., Th1, Th2, Th17, Treg, Tr1, etc.)
Interferon-γ (IFN-γ)	inflammatory cytokine mainly expressed by T cells that promotes parasite clearance through activation of phagocytes and antibody isotype switching in B cells
Interleukin-2 (IL-2)	an essential mitogen and T cell growth factor
Interleukin-6 (IL-6)	inflammatory cytokine and pyrogen that promotes IL-17 expression by T cells, B cell activation and differentiation, and the antimicrobial properties of phagocytes
Interleukin-10 (IL-10)	regulatory cytokine that directly suppresses professional APC activation, which indirectly suppresses the induction of potent T cell responses
Interleukin-17 (IL-17)	inflammatory cytokine that promotes recruitment, activation, and differentiation of phagocytes and neutrophils
Interleukin-27 (IL-27)	regulatory cytokine that suppresses the activation and proliferation of IL-17-secreting T cells and promotes the generation of IL-10-secreting T cells
Macrophages	tissue resident phagocytes essential for control of <i>Plasmodium</i> and <i>Toxoplasma</i> via direct engulfment of parasite or parasite-infected host cells
Merozoites	non-motile, asexually reproducing form of the <i>Plasmodium</i> parasite responsible for infection and destruction of host red blood cells
Monocytes	circulating cells of the myeloid lineage that are rapidly recruited to sites of inflammation where they differentiate into macrophages or DCs
Naturally occurring T regulatory cells (nTregs)	regulatory Foxp3 ⁺ CD4 T cell that develops in the thymus, exhibits specificity for self-antigens, and exerts suppressive function through both contact-dependent mechanisms and/or secretion of IL-10 and/or TGF- β
Peripherally induced T regulatory cells (iTregs)	regulatory Foxp3 ⁺ CD4 T cell that develops in the periphery after infection or immune insult, exhibits specificity for either self- or non-self-antigens, and exerts suppressive function through both contact-dependent mechanisms and secretion of IL-10 and/or TGF- β

Polarized T cell	a T cell that is activated and expresses molecules associated with a particular T helper cell subtype
Pyrogens	proteins produced by phagocytes that induce fever
Secondary lymphoid tissue	an essential immune tissue (draining lymph nodes and spleen) that serves as the site of naïve T cell activation and induction of adaptive immune responses
Sporozoites	form of <i>Plasmodium</i> deposited during mosquito blood meal feeding. <i>Plasmodium</i> sporozoites infect hepatocytes to establish infection in mammalian hosts. Also an infectious form of <i>Toxoplasma</i> excreted by felids while encased in oocysts. Sporozoites are released from ingested oocysts and develop into bradyzoites or tachyzoites
Tachyzoites	rapidly replicating form of <i>Toxoplasma</i> responsible for direct cell death, tissue pathology, and systemic dissemination in mammalian hosts
T-bet	transcription factor functionally linked to the differentiation and activity of Th1 cells and inhibition of Th2 and Th17 cell differentiation
Tr1	peripherally induced T regulatory cells, regulatory population of CD4 T cell that develops in the periphery after infection or immune insult, exhibits specificity for non-self-antigens and exerts suppressive function through secretion of IL-10 and/or TGF- β . Unlike iTregs, Tr1 cells do not express Foxp3
Transforming growth factor-β (TGF-β)	pleiotropic anti-inflammatory cytokine that suppresses T helper cell activity and promotes the differentiation and maintenance of Tregs
Tumor necrosis factor-α (TNF-α)	inflammatory cytokine and pyrogen expressed primarily by macrophages but also by T cells and B cells. TNF- α promotes apoptotic cell death and contributes to pathogen control through activation, proliferation, and differentiation of leukocytes
Type 1 T helper cell (Th1) cells	functionally distinct subset of CD4 T helper cells associated with IFN- γ and TNF- α expression and resistance to intracellular pathogens
Type 2 T helper (Th2) cells	functionally distinct subset of CD4 T helper cells associated with IL-4, IL-5, and IL-13 expression and the induction of allergic responses
Type 17 T helper (Th17) cells	functionally distinct subset of CD4 T helper cells associated with IL-17 expression and resistance to bacterial and fungal pathogens

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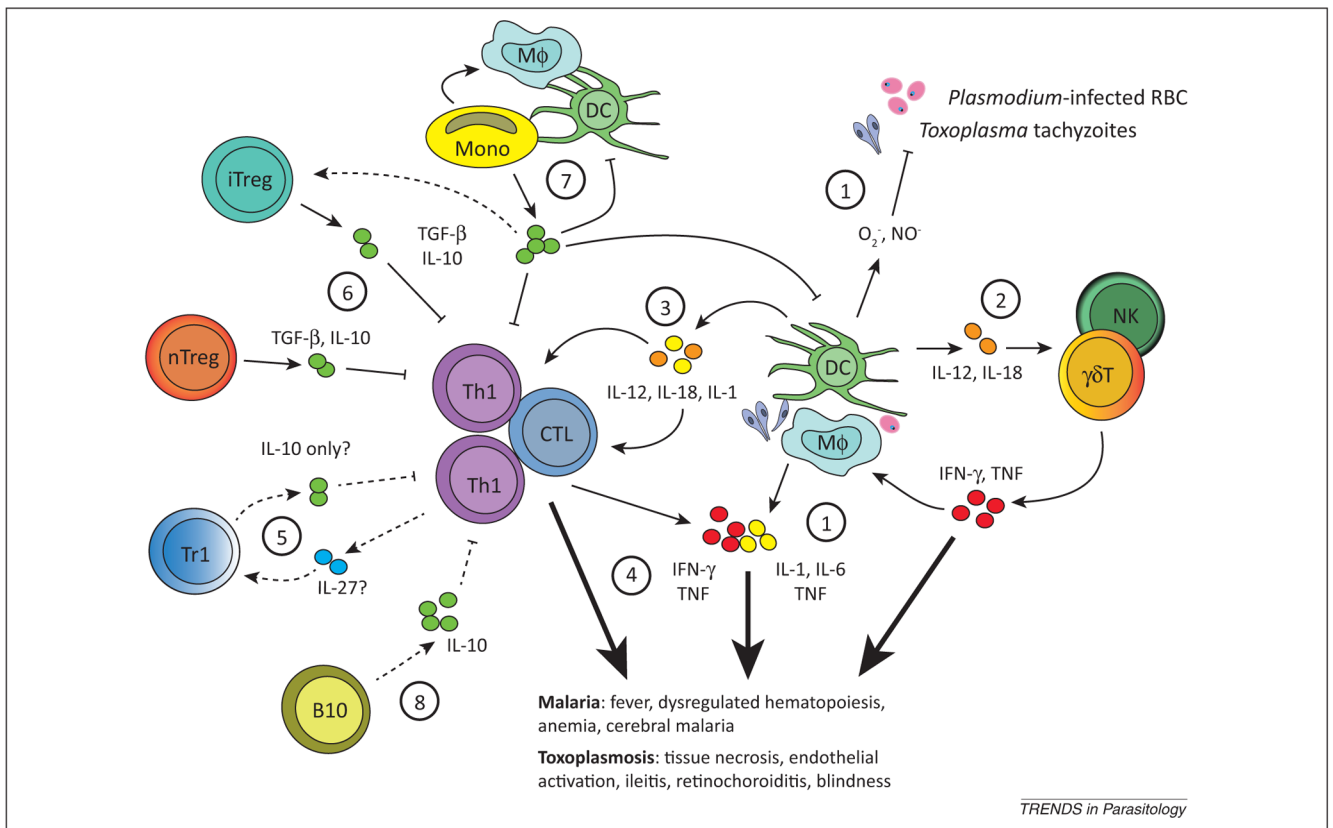


Figure 1.

Common regulatory networks limit *Toxoplasma* and *Plasmodium* immunopathogenesis. (1) Recognition of parasites or parasite-infected cells by macrophages (Mφ) and dendritic cells (DCs) triggers the production of antiparasitic reactive oxygen species and synthesis of inflammatory cytokines [88–92]. These inflammatory agents can act directly on parenchymal cells to induce necrotic or apoptotic pathways causing tissue damage, or (2) indirectly through the activation of innate natural killer (NK) or $\gamma\delta$ T cells ($\gamma\delta$ T) [93,94]. (3) Cytokines such as IL-12, IL-1, and IL-18 also play a critical role in shaping the inflammatory nature of adaptive T cell subsets, which include IFN- γ secreting T helper cells (Th1) and cytotoxic cells (CTL) [85,93]. (4) Effector T cells express several inhibitory receptors including programmed death-1 (PD-1) and cytotoxic T lymphocyte antigen-4 (CTLA-4) that act to limit their activation, proliferation, and cytokine expression, which helps maintain immune homeostasis and limit immunopathology after infection [5,6,11,13]. Generally, and despite inhibitory receptor expression, during acute intestinal or ocular toxoplasmosis and malaria, highly activated T cells appear to directly contribute to pathology [37,46,98,104]. Pathological inflammatory reactions are themselves counterbalanced by several secreted regulatory cytokines and functionally distinct immune cell subsets. Regulatory cytokines such as IL-10 and TGF- β can be secreted by multiple cell types, including B cells, T cells, monocytes (Mono), Mφ, and DCs [2,17,18,24,26,29,51,67]. (5) Polarized IFN- γ -secreting Th1 cells also have the capacity to secrete IL-10 after their differentiation into Treg-like (Tr1) cells. Tr1 differentiation may involve IL-27 [23]. (6) Natural, thymus-derived Tregs (nTregs) suppress the activity of inflammatory CTL and Th1 cells through direct contact or secretion of IL-10 and TGF- β [12,46,106]. The precise role of nTreg, inducible Treg (iTreg), and Tr1 in limiting immunopathology during malaria and toxoplasmosis is not well understood. (7) Monocytes that infiltrate *Toxoplasma*-infected

tissues or the spleen of *Plasmodium*-infected hosts exhibit dual roles in pathology and protection, by either secreting inflammatory cytokines (e.g., IL-12, TNF- α , IL-1, IL-6) or regulatory cytokines (IL-10 or TGF- β), and possibly by driving the differentiation of Tregs [52–56]. The cues that determine whether monocytes exhibit an inflammatory or regulatory function are unknown. Additional questions include the specific role of IL-27 and the cellular source(s) of this cytokine, as well as **(8)** the contribution of IL-10-secreting regulatory B cells (B10). Well-described pathways are shown as unbroken lines, and unknown or speculative pathways are shown as broken lines.

Table 1

Regulatory cytokines that limit immunopathology during toxoplasmosis and malaria

Cytokine	Key cellular sources	Cellular targets	Predominant effect <i>in vivo</i>	Refs
IL-10	T cells, macrophages, DCs, B cells	DCs, B cells, macrophages, epithelial cells, endothelial cells	Suppress activation and antigen presentation by DCs, B cells, and macrophages	[2,17,18,24,26,29,51,67]
TGF- β	T cells, B cells, NK cells, macrophages, epithelial cells, endothelial cells, platelets, neurons	T cells, DCs, macrophages, B cells	Induction of iTreg and Th17 differentiation, trigger effector T cell apoptosis	[26,32,33,40,63,107–109]
IL-27	Unknown	CD4 T cells, DCs, macrophages	Induction of Tr1 cell differentiation	[23,40,46–49,110]