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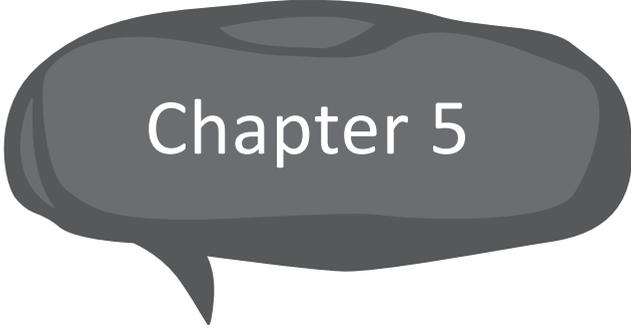


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## Chapter 5

IL-10-producing CD1d<sup>hi</sup> regulatory B cells from *Schistosoma haematobium*-infected individuals induce IL-10-positive T cells and suppress effector T-cell cytokines

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## Abstract

**Background.** Chronic schistosome infections are associated with T-cell hyporesponsiveness and a strong regulatory network. Murine studies have shown that schistosome infections can induce regulatory CD1d<sup>hi</sup> B cells, which inhibit inflammatory responses. Here, we evaluated the influence of regulatory B cells (Bregs) on T-cell cytokines *in vitro* in human schistosomiasis.

**Methods.** Gabonese young adults were recruited from areas where *Schistosoma haematobium* (*S.h*) infections were high or low endemic. The study participants were categorized as infected or uninfected from a high endemic area or uninfected from a low endemic (nonendemic) area. Their B cells were studied for Breg subset markers and cocultured with allogenic anti-CD3-stimulated CD4<sup>+</sup> T cells, followed by T-cell cytokine analysis.

**Results.** A greater percentage of B cells from *S. haematobium*-infected donors expressed cytoplasmic interleukin(IL) 10 (IL-10) and membrane-bound latency-associated peptide/transforming growth factor  $\beta$ 1 (LAP/TGF- $\beta$ 1), compared with uninfected donors. T cells produced less interferon  $\gamma$  (IFN- $\gamma$ ), IL-4 and IL-17 upon coculture with B cells from schistosome-infected individuals only, while the conversion to CD25<sup>hi</sup>FoxP3<sup>+</sup> and the percentage of IL-10<sup>+</sup> T cells was enhanced. Interestingly, depletion of the prominent IL-10-producing B-cell subset, CD1d<sup>hi</sup> cells, resulted in less IL-10<sup>+</sup> T cells in the *S. haematobium*-infected group, while levels of FoxP3<sup>+</sup> regulatory T cells remained unaffected.

**Conclusions.** Schistosomes can induce functional Bregs in humans that may be instrumental in general T-cell hyporesponsiveness and may contribute to the increased regulatory milieu found in schistosomiasis.

## Introduction

Helminths such as schistosomes are master regulators of host immune responses, characterized by polarized T-helper type 2 (Th2) cells (1;2) and a general T-cell hyporesponsiveness in the chronic stage of infection (3). This is facilitated by a number of potent regulatory cell types and increased levels of immunosuppressive cytokines, such as interleukin 10 (IL-10) and/or transforming growth factor  $\beta$  (TGF- $\beta$ ) (4-6). While B lymphocytes possess a variety of immune functions, more recently they were shown to be a relevant source of IL-10 and/or TGF- $\beta$  (7;8). Such B cells were termed regulatory B cells (Bregs) as they were able to ameliorate various hyperinflammatory diseases (9-13). For example, IL-10-producing B cells suppressed experimental autoimmune encephalomyelitis (EAE), collagen-induced arthritis (CIA), and inflammatory bowel disease (IBD) (14-17). Furthermore, TGF- $\beta$ -producing B cells protected against acute allergic airway inflammation (8), and inhibited spontaneous Th1 autoimmunity in nonobese diabetic mice (18). Recently, Bregs have been considered relevant in infectious diseases, such as helminth infection. During *Schistosoma mansoni* infection, active Bregs were enriched in the splenic marginal zone, and suppressed allergic responses via elevated IL-10 production and/or FoxP3<sup>+</sup> T-regulatory (Treg) cell induction (19;20). Likewise, during chronic *Heligiosomoides polygyrys* infections CD5<sup>+</sup>CD23<sup>hi</sup> B cells from mesenteric lymph nodes also reduced allergic airway inflammation, but they did so independently of IL-10 (21).

Although the majority of studies on Bregs were conducted in mouse models, Bregs are now also identified in humans during infectious diseases. For example, during chronic hepatitis B virus infection IL-10-producing CD24<sup>hi</sup>CD38<sup>hi</sup> B cells suppressed antigen-specific CD8<sup>+</sup> T-cell responses (22), while in systemic lupus erythematosus, IL-10 production by this Breg subset was impaired compared with findings in healthy controls, and as a consequence Th1 cytokine production was not reduced (23). In human immunodeficiency virus (HIV)-infected individuals, *in vitro* depletion of CD24<sup>hi</sup>CD38<sup>hi</sup> B cells resulted in enhanced anti-HIV CD8<sup>+</sup> T-cell responses, which was dependent on IL-10 (24). Furthermore, CD24<sup>hi</sup>CD27<sup>+</sup> Bregs from healthy individuals reduced TNF- $\alpha$  production in monocytes *in vitro* via IL-10 (25), however this Breg subset is not yet studied in the context of infectious diseases. Lastly, during *Mycobacterium tuberculosis* infection CD1d<sup>+</sup>CD5<sup>+</sup> B-cell numbers were increased and inhibited IL-17 production by T cells (26). Although active TGF- $\beta$ -producing CD5<sup>+</sup> Bregs were described in healthy individuals in response to milk antigens (7), the role of human TGF- $\beta$ -expressing Bregs in infectious diseases has not yet been addressed.

Enhanced numbers of IL-10-producing CD1d<sup>hi</sup> B cells were first described in helminth-infected Argentine patients with multiple sclerosis (MS) and could inhibit autoreactive T-cell responses *in vitro* (27). We found a similarly elevated percentage of IL-10-producing B cells in *Schistosoma haematobium*-infected Gabonese children, which returned to baseline levels after anti-schistosome

treatment (20). However, the functional properties of these cells were not explored. In the current study, we investigated the regulatory characteristics of peripheral B cells of *S. haematobium*-infected individuals and the functional consequences for CD4<sup>+</sup> T-cell activation. We observed that CD1d<sup>hi</sup> B cells and CD24<sup>hi</sup>CD27<sup>+</sup> B cells of *S. haematobium*-infected adults express enhanced levels of cytoplasmic IL-10 and enhanced levels of surface latency-associated peptide (LAP), part of a latent TGF- $\beta$  complex, respectively. In coculture with T cells, reduced effector T-cell cytokine responses, as well as more Tregs, were associated with B cells from infected individuals, compared with B cells from uninfected controls who resided in areas where *S. haematobium* is nonendemic. Depletion of the CD1d<sup>hi</sup> B-cell population resulted in a loss of IL-10<sup>+</sup> T-cell induction but did not yield differences in the numbers of FoxP3<sup>+</sup> Tregs, suggesting that other CD1d<sup>-</sup> Bregs may be important for FoxP3<sup>+</sup> T-cell induction. Taken together, schistosome-induced Bregs exhibit regulatory activity that may contribute to strong regulatory responses found during chronic schistosomiasis.

## Materials and Methods

### Study population

Heparinized venous blood was obtained from *S. haematobium*-uninfected young adults living in Lambaréné, Gabon, a semiurban municipality where *S. haematobium* is not endemic and the prevalence of *S. haematobium* infection is low; these individuals composed the "nonendemic uninfected group". Infected donors and additional uninfected donors were recruited from the rural village Bindo, where *S. haematobium* is endemic, residents are likely to have a history of *S. haematobium* infections, and no intervention studies have been performed (28); these donor groups composed the "endemic infected group" and the "endemic uninfected group", respectively. Current users of praziquantel were excluded from study participation, as were people who had lived in either region for <1 year before enrollment.

The presence of helminth and protozoan infections was determined as previously reported (20) (Table I). Infections with *Plasmodium (P.) falciparum* and *P. malariae* were determined by polymerase chain reaction (PCR) (29) and *Loa Loa* and *Mansonella perstans* microfilariae by microscopy (30). Written informed consent was obtained from all participants, and they were treated according to local guidelines. The study was approved by the Comité d'Éthique Regional Independent de Lambaréné (CERIL N°08/10) and conducted according to the principles expressed in the Declaration of Helsinki.

### Cell isolation and characterization

Peripheral blood mononuclear cells (PBMCs) were isolated by Ficoll-Hypaque density gradient centrifugation from approximately 40 ml of blood in 50 ml tubes (Greiner Bio-One). CD19<sup>+</sup> B cells, untouched CD4<sup>+</sup>, and CD4<sup>+</sup>CD25<sup>-</sup> T cells were isolated by magnetic-bead purification (purity, >96%). For CD1d<sup>hi</sup> depletion, B cells were first incubated with CD1d-APC (Biolegend), washed, and incubated with 20 µl/10<sup>7</sup> anti-APC MicroBeads (Miltenyi Biotec). The cells were then split in 2 groups, CD1d<sup>hi</sup>-depleted B cells (86% depleted) and mock-depleted B cells, to which the eluted CD1d<sup>hi</sup> B cells were added back after depletion. Part of the B cells were cryopreserved in cryovials (Greiner Bio-One) and transported to Leiden University Medical Center, Leiden, The Netherlands.

In PBMCs, B cells were characterized using CD1d-PE, CD19-PacificBlue, CD27-APCeFluor780 (all eBioscience), CD24-PeCy7 (ITK), and CD38-FITC (BD). For PFA-fixed PBMCs: CD1d-PE, CD20-APCeFluor780 (eBioscience), CD24-PeCy7, CD27-Biotin (eBioscience) plus streptavidin-Qdot525 (Life Technologies), CD38-Horizon450 (BD), and LAP1-APC (Biolegend). For all flowcytometric measurements, all samples were stained in 96-well V-bottom plates (Greiner Bio-One), FcγR-binding inhibitor was added to the antibody mix, and FMOs were used for gate setting for all surface markers and cytokines.

**B-cell stimulation and intracellular staining**

Freshly isolated B cells ( $1 \times 10^5$ ) were stimulated in medium (RPMI; Life Technologies) supplemented with 10% FCS (Greiner Bio-One), 1 mM pyruvate, 2mM Glutamate and antibiotics (100 U/ml penicillin and 100  $\mu\text{g}/\text{ml}$  streptomycin; all Life Technologies), in the presence of 100 ng/ml PMA, 1  $\mu\text{g}/\text{ml}$  ionomycin, 10  $\mu\text{g}/\text{ml}$  BrefeldinA (all Sigma-Aldrich) and 10  $\mu\text{g}/\text{ml}$  ultrapure LPS (Life Technologies) (25). After 5 hours, the cells were fixated with 1.9% PFA (Sigma-Aldrich) and stained for CD1d-PE, CD20-APCeFluor780, CD24-PeCy7, CD27-APC (BD), CD38-FITC, IL-10-Biotin (Abd Serotec) plus streptavidin-Qdot525, and TNF- $\alpha$ -eFluor450 (eBioscience).

**Functional *in vitro* assays**

B cells ( $1 \times 10^5$ ) were cultured in 96-well U-bottom plates (Greiner Bio-One) with allogenic CD4<sup>+</sup> T cells ( $1 \times 10^5$ ) from 1 European donor in the presence of 1  $\mu\text{g}/\text{ml}$  soluble anti-CD3 (IXE M1654, Sanquin) for 3 days or, in parallel, with CD4<sup>+</sup>CD25<sup>-</sup> T cells for 6 days (23). After 3 days, cells were restimulated with PIB, PFA-fixed and stained in 0.5% saponin buffer for IL-4-PE (BD), IL-10-Biotin plus streptavidin-Qdot525, IL-17-FITC (Biolegend), IFN- $\gamma$ -HorizonV450 (BD), TNF- $\alpha$ -PeCy7 (eBioscience), CD3-PerCPCy5.5 (BD), and CD20-APCeFluor780. After 6 days, restimulated cells fixed with eBioscience FoxP3 fixation buffer were stained in permeabilization buffer for: CD1d-APC (Biolegend), CD3-PerCPCeFluor710 (eBioscience), CD20-APCeFluor780, CD25-PE (BD), IL-10-Biotin plus streptavidin-Qdot525, TNF- $\alpha$ -PeCy7, FoxP3-eFluor450 (eBioscience), and Helios-FITC (Biolegend).

**Statistical analysis**

For non-parametric analysis Kruskal-Wallis, Mann-Whitney *U*, and Wilcoxon matched pairs tests were used. For categorical data, the Pearson Chi-squared and Fisher exact tests was used. All tests were performed used Graphad Prism. Correlations were analyzed using SPSS Statistics 20 (IBM). A *P* value <.05 was considered statistically significant.

## Results

### Elevated B-cell IL-10 and LAP/TGF- $\beta$ 1 expression in *S. haematobium*-infected subjects

Peripheral blood specimens was collected from 14 young adults in the endemic infected group and compared to specimens from 12 individuals in the endemic uninfected group and from 12 individuals from the nonendemic uninfected groups. As described in Table I, no significant differences were found between the 3 groups in the prevalence and infection intensity of other parasitic infections such as malaria, *Ascaris lumbricoides* infection, *Trichuris trichiura* infection, or hookworm infection. Furthermore, sex was equally distributed, but the age of the *S. haematobium*-infected individuals was slightly lower than that of uninfected individuals. Age was not regarded as a confounder, however, as the real age differences were very small (3-4 years). The average number of isolated PBMCs ( $1.1 \times 10^6$ ,  $1.0 \times 10^6$ ,  $1.0 \times 10^6$  cells/ml,  $P = .5$ ) and the average percentage of isolated B cells (8.3%, 8.3%, 6.2%,  $P = .4$ ) were comparable between the endemic infected group, endemic uninfected group, and nonendemic uninfected group, respectively.

CD19<sup>+</sup> B cells isolated from peripheral blood specimens were studied for the presence of IL-10-competent B cells by a short, 5-hour stimulation period with PMA, ionomycin, Brefeldin A and LPS (PIB+LPS) as described by Iwata *et al.* (25). TNF- $\alpha$  was included as an inflammatory control. The percentage of B cells expressing cytoplasmic IL-10 was the highest in the endemic infected group and lowest in the nonendemic uninfected group (Fig. 1A). TNF- $\alpha$  production was similar in all groups (Fig. 1A, Appendix S1A). The percentage of IL-10-producing B cells was not correlated with age, helminth burden (based on *S. haematobium* egg counts; data not shown), or *Plasmodium* parasites (Fig. 1C) (31). Of note, PIB-only stimulation also showed a greater percentage IL-10-producing B cells in the endemic infected group, compared to the other 2 groups, although in each group, levels after PIB-only stimulation were lower than those after PIB+LPS stimulation (data not shown).

We also analyzed the expression of LAP/TGF- $\beta$ 1. Dissociation of this complex results in bioactive TGF- $\beta$ 1. B cells from the endemic infected group expressed significantly more LAP, while B cells from endemic uninfected individuals showed an increased trend, compared with the nonendemic uninfected group (Fig. 1B, Appendix S1B). We did not study the coexpression of TGF- $\beta$ 1 and IL-10 in B cells because this investigation requires different methods (ie, B cells need to be stimulated for IL-10, but not for LAP/TGF- $\beta$  detection). These results indicate that schistosome infections or exposure induces elevated numbers of circulating IL-10-producing and/or LAP/TGF- $\beta$ 1-expressing B cells.

### Phenotypic characterization of IL-10 and LAP/TGF- $\beta$ 1-expressing B cells

To determine which Breg subset was responsible for the increased IL-10 production and/or LAP/TGF- $\beta$ 1 expression, PBMCs were analyzed for the presence of 3 different human Breg subsets: CD1d<sup>hi</sup> B cells, CD24<sup>hi</sup>CD38<sup>hi</sup> B cells and CD24<sup>hi</sup>CD27<sup>+</sup> B cells (23;25;27). Only the CD1d<sup>hi</sup> subset was elevated during infection (from 1.9% to 2.5%), whereas the other 2 subsets were similar among the 3 study groups (approximately 5% CD24<sup>hi</sup>CD38<sup>hi</sup> and 21% CD24<sup>hi</sup>CD27<sup>+</sup>) (Fig.

**Table I: Characteristics of study population**

	Endemic Infected (n=14)	Endemic Uninfected (n=12)	Nonendemic Uninfected (n=12)	P
<b>Study area</b>	Bindo	Bindo	Lambaréné	
<b>Gender male/ female</b>	7/7	6/6	6/6	
<b>Mean age in yrs (range)</b>	18.7 (16-23)	23 (18-28)	21.8 (18-27)	0.002 <sup>#</sup>
<b><i>S. haematobium</i></b>	14 (100%)	0	0	
Mean egg count (range)	165 (1-798)	-	-	
<b>Other helminths</b>				
Any helminths	6 (43%)	4 (33%)	3 (25%)	0.631*
<i>Ascaris</i> (%)	4 (29%)	2 (17%)	1 (8%)	0.407*
<i>Trichuris</i> (%)	4 (29%)	2 (17%)	1 (8%)	0.407*
<i>Hookworm</i> (%)	3 (21%)	2 (17%)	1 (8%)	0.656*
<i>Entamoeba histolytica</i> (%)	0 (0%)	0 (0%)	1 (8%)	0.329*
<b>Malaria (PCR)</b>				
Any malaria	4 (29%)	7 (58%)	3 (25%)	0.172*
<i>P. falciparum</i>	4 (29%)	6 (50%)	3 (25%)	0.372*
<i>P. malariae</i>	0 (0%)	1 (8%)	1 (8%)	0.540*
<b>Microfilaria</b>				
Any microfilaria	3 (21%)	5 (42%)	2 (17%)	0.332*
<i>Loa loa</i>	3 (21%)	2 (17%)	0 (0%)	0.248*
<i>Mansonella perstans</i>	0 (0%)	3 (25%)	2 (17%)	0.155*

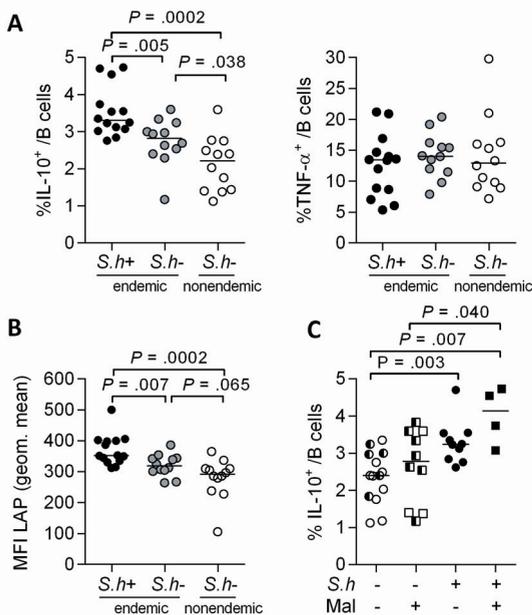
The study was initiated in January 2011, and participants were from 2 villages in Gabon.

<sup>#</sup> By the Kruskal-Wallis test for age. Mann-Whitney test shows the following differences:  $P = .002$  between endemic infected and -uninfected group,  $P = .006$  between endemic infected and nonendemic uninfected group, and  $P = .255$  between endemic uninfected and nonendemic uninfected groups.

\* By the Pearson Chi-squared test. Because of sample size, the Fisher exact test was performed between the groups, and results showed no differences between the groups for all co-infections.

2A/B, Appendix S1C). Interestingly, in the endemic infected group, the subset with the greatest percentage of cells producing IL-10 was the CD1d<sup>hi</sup> B cells (25 %), and only a small percentage of CD24<sup>hi</sup>CD27<sup>+</sup> B cells produced IL-10 (2.7 %; Fig. 2C). Within the IL-10-producing B-cell pool, approximately 10-20% of the B cells shared a CD1d<sup>hi</sup> or CD24<sup>hi</sup>CD27<sup>+</sup> phenotype. However, only the percentage of CD1d<sup>hi</sup> B cells within the IL-10-producing B-cell population significantly increased during infection, again indicating that mostly these cells are induced during schistosome infections. The percentage of the different Breg subsets was not correlated with helminth burden (data not shown).

Increased levels of LAP/TGF- $\beta$ 1 expression in total B cells from endemic infected individuals was mainly attributed to the CD24<sup>hi</sup>CD27<sup>hi</sup>, and partly to the



**Figure 1.** Intracellular interleukin 10 (IL-10) and tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ) detection in total B cells.  $1 \times 10^5$  blood-derived B cells from *Schistosoma haematobium*-infected individuals from a *S. haematobium*-endemic area (S.h+), uninfected individuals from a *S. haematobium*-endemic area (S.h-), and uninfected individuals from a *S. haematobium*-nonendemic area (S.h-) were directly stimulated for 5 hours with PMA, ionomycin, and brefeldinA (PIB) in combination with ultrapure lipopolysaccharide (LPS). (A) Intracellular IL-10 production and TNF- $\alpha$  production were analyzed using flow cytometry (FACS Canto II, BD Biosciences). (B)  $1 \times 10^5$  PFA-fixed peripheral blood mononuclear cells were stained for latency-associated peptide (LAP)/transforming growth factor  $\beta$ 1. The geometric mean fluorescence intensity (MFI) of LAP expression on total B cells was determined. (C) Intracellular IL-10 in B cells after 5 hours of PIB + LPS stimulation. Donors were grouped according to the presence of malaria parasite infection (Mal) and/or *S. haematobium* infection. The malaria-negative *S. haematobium*-negative group contains 14 individuals (5 of whom were from Bindo), the malaria-positive *S. haematobium*-negative group contains 10 (7 of whom were from Bindo), the malaria-negative *S. haematobium*-positive group contains 10, and the malaria-positive *S. haematobium*-positive group contains 4. Donors from Bindo are depicted in Figure 1C as half-shadowed circles or squares and donor from Lambaréné as open circles or squares. Each individual is represented as a separate dot, with a line showing the median value for the group. Data were analyzed using the Mann-Whitney *U* test.

CD24<sup>hi</sup>CD38<sup>hi</sup> B-cell population, but not to the CD1d<sup>hi</sup> B cells (Fig. 2D). The difference in IL-10 expression between the endemic infected group and the uninfected groups was lost when CD1d<sup>hi</sup> B cells were removed from the CD24<sup>hi</sup>CD27<sup>+</sup> B-cell gate, but the difference in LAP/TGF- $\beta$ 1 expression was maintained (Appendix S2). Therefore, these data suggest that the CD1d<sup>hi</sup> B cells are the dominant subset that increases in number and capacity to produce IL-10 during *Schistosoma* infection, while increased LAP/TGF- $\beta$ 1 has a stronger association with CD24<sup>hi</sup>CD27<sup>+</sup> B cells, although the frequency of this subset did not change during schistosomiasis.

### **Schistosome-induced B cells dampen the pro-inflammatory T-cell cytokine profile**

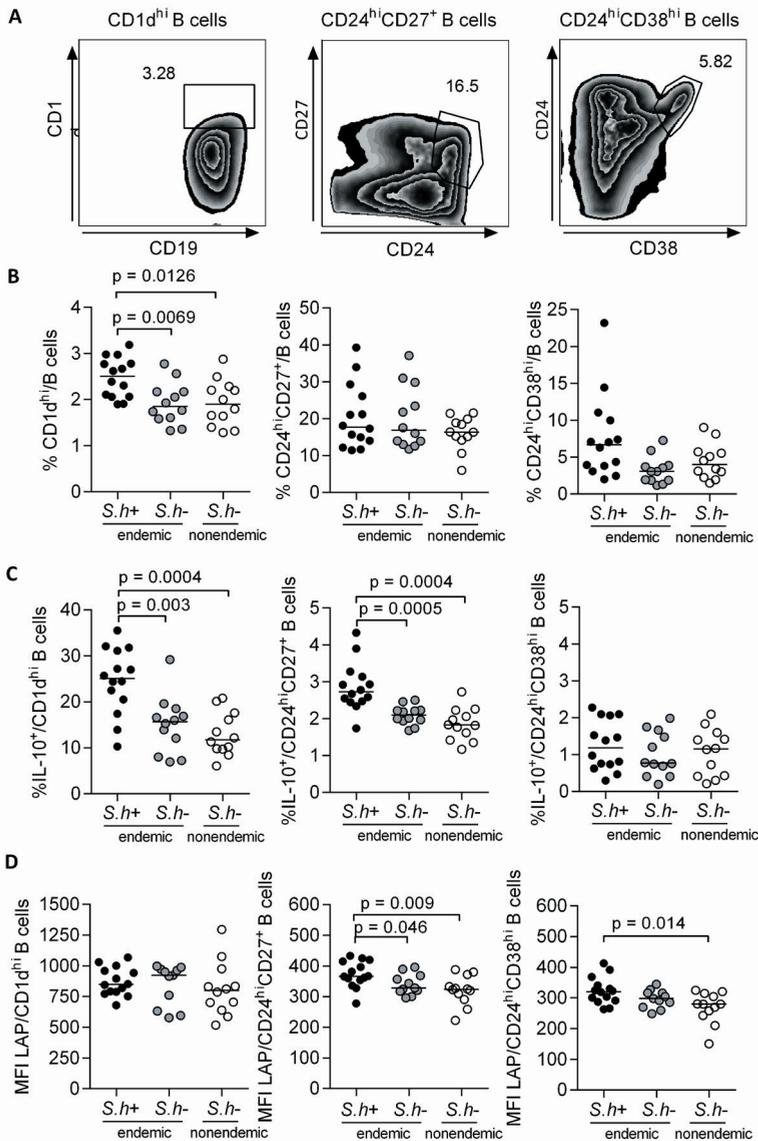
We next performed *in vitro* CD4<sup>+</sup> T and B cell cocultures in the presence of anti-CD3 to investigate whether the T-cell stimulatory activity of B cells from the endemic infected group is altered. Allogenic CD4<sup>+</sup> T cells from a single donor were used, to prevent differences due to variation in T-cell donors. After 3 days, the cells were restimulated with PIB, and we observed a slight but significantly lower percentage of CD4<sup>+</sup> T cells producing IFN- $\gamma$ -, IL-4- and IL-17 when stimulated in the presence of B cells from the endemic infected group, compared to nonendemic uninfected group, whereas TNF- $\alpha$  production was not affected (Fig. 3A-D). T cells stimulated with anti-CD3 only, in the absence of B cells, showed a variable degree of viability and were therefore excluded from the analysis. No significant differences were detected between endemic infected and endemic uninfected individuals, suggesting that past exposure to *Schistosoma* infections can yield changes in B-cell function, reflected in the slightly enhanced IL-10 and TGF- $\beta$  expression (Fig. 2), which may be sufficient to control effector T-cell responses.

### **CD1d<sup>hi</sup> B cells induce IL-10-producing T cells**

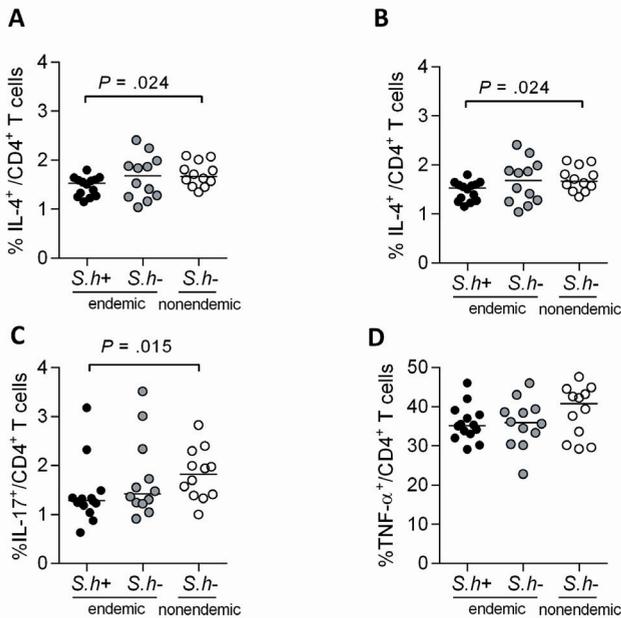
To investigate whether schistosome-induced B cells can prime the regulatory functions in T cells, either by enhancing IL-10 production (as in Tr1 cells) or by inducing CD25<sup>hi</sup>FoxP3<sup>hi</sup> Tregs, CD4<sup>+</sup>CD25<sup>-</sup> T cells were cocultured with B cells for 6 days. Interestingly, the frequency of IL-10-producing T cells was significantly higher after coculture with B cells from the endemic infected group, compared with the uninfected groups (Fig. 4A). In addition, more FoxP3<sup>hi</sup> Tregs were found in cocultures with B cells from the endemic infected group, compared to uninfected groups, although the variation in FoxP3 induction was greater than the variation in IL-10 induction (Fig. 4B, Appendix S3). When Helios expression was analyzed (as an additional marker for putative functional FoxP3<sup>+</sup> Tregs), the frequency of Helios<sup>+</sup>CD25<sup>hi</sup>FoxP3<sup>hi</sup> T cells still remained significantly higher in the endemic infected group (data not shown).

To explore the underlying mechanism, we next focused on the dominant IL-10-producing B-cell subset, CD1d<sup>hi</sup> B cells, to examine the effect of helminth-induced IL-10-producing B cells on regulatory functions of T cells (Fig. 4C). Because of the limited cell numbers available, we used a CD1d<sup>hi</sup> B cell depletion strategy instead of isolating CD1d<sup>hi</sup> B cells. Similar to findings for the total B-cell population

from the endemic infected group, the mock-depleted B cells induced higher levels of IL-10 and/or CD25<sup>hi</sup>FoxP3<sup>hi</sup> T cells compared to B cells from uninfected

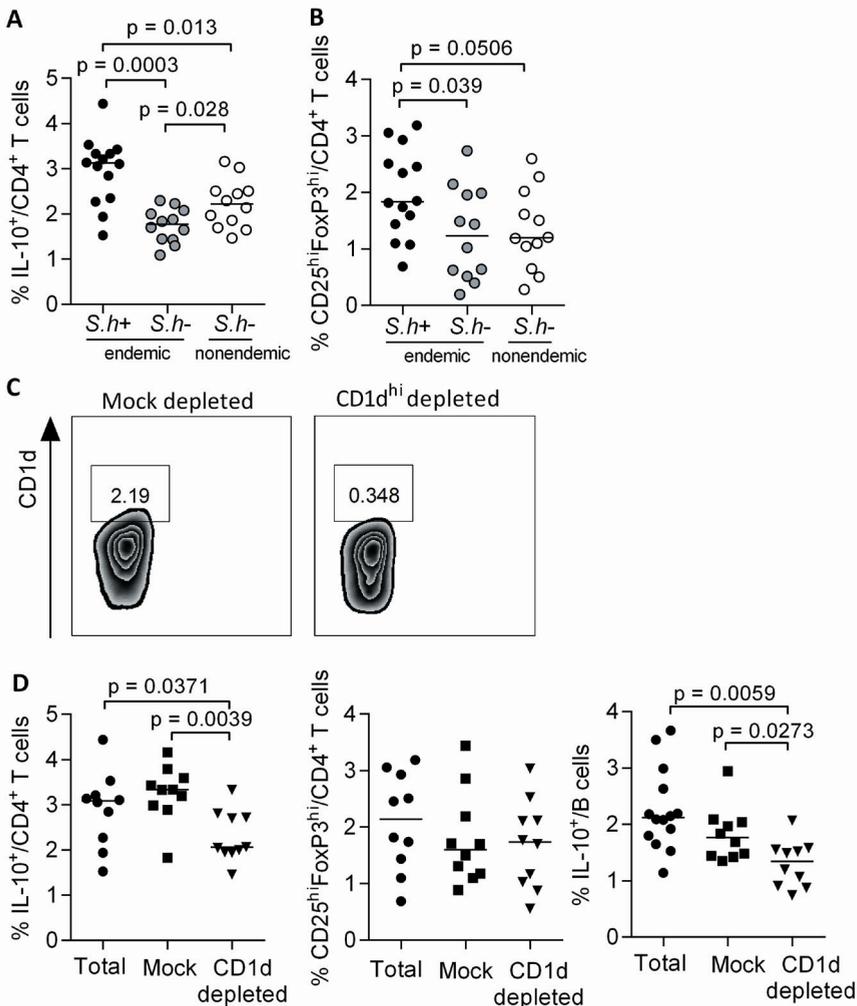


**Figure 2.** The frequency of different regulatory B cells in the blood specimens from *Schistosoma haematobium*-infected individuals from a *S. haematobium*-endemic area (S.h+), uninfected individuals from a *S. haematobium*-endemic area (S.h-) and uninfected individuals from a *S. haematobium*-nonendemic area (S.h-). (A) *Ex vivo* PBMCs were gating for CD19<sup>+</sup>, following by specific Breg subset gating as indicated in the figure by a representative gating example involving a *S. haematobium*-infected donor. (B) The percentage of CD1d<sup>hi</sup>, CD24<sup>hi</sup>CD27<sup>+</sup>, and CD24<sup>hi</sup>CD38<sup>hi</sup> B cells. (C) IL-10 production of each B-cell subset after 5 hours of stimulation with PIB+LPS. (D) The geometric MFI expression of LAP/TGF-β1 on the 3 Breg subsets. For all data, the Mann-Whitney *U*-test was used for analysis.



**Figure 3.** T-cell cytokine expression after coculture with B cells from *S. haematobium*-infected (*S.h+*) and uninfected individuals (*S.h-*) from a *S. haematobium*-endemic area, and uninfected individuals from a *S. haematobium*-nonendemic area (*S.h-*). Isolated B cells were cultured at a ratio of 1:1 with allogenic CD4<sup>+</sup> T cells (from 1 European donor) in the presence of soluble anti-CD3 (1  $\mu$ g/ml) for 3 days. After coculturing, the cells were restimulated with 100  $\mu$ g/ml PMA and 1  $\mu$ g/ml ionomycin for 6 hours, in the presence of 10  $\mu$ g/ml BrefeldinA for the last 4 hours. The cells were fixed with 1.9% PFA. Single cytokine-producing T cells were gated for pro-inflammatory cytokines IFN- $\gamma$  (A), IL-4 (B), IL-17 (C) and TNF- $\alpha$  (D). The Mann-Whitney *U*-test was used for data analysis.

groups. After the depletion of CD1d<sup>hi</sup> B cells, the percentage of T cells expressing IL-10 returned to levels similar to those induced by B cells from the uninfected groups, whereas no effect on the (elevated) level of CD25<sup>hi</sup>FoxP3<sup>hi</sup> T cells was observed (Fig. 4D, Appendix S3A-B). Indeed, B-cell IL-10 was also reduced to levels found in uninfected donors following CD1d depletion (Fig. 4D, Appendix S3C), suggesting that B-cell expression of IL-10 may play a role in the induction of IL-10-producing T cells but not the induction of FoxP3<sup>+</sup> Tregs. TGF- $\beta$  is known to be important for the generation of FoxP3<sup>+</sup> Tregs (32). To investigate its role in Treg induction, cryopreserved Gabonese B cells were cocultured with freshly obtained T cells and blocking anti-TGF- $\beta$  at our Dutch research institute. In contrast to the culture of fresh specimens performed at the field site, isotype-cultured B cells from infected individuals did not enhance Treg induction, suggesting that freezing and thawing the cells may have affected their function, despite a good viability after thawing (> 94 %; data not shown). Therefore, the role of TGF- $\beta$ -expressing B cells in the induction of FoxP3<sup>+</sup> Tregs could not be addressed within the current study. Nevertheless, it is tempting to suggest that CD24<sup>hi</sup>CD27<sup>+</sup> B cells may be involved in FoxP3<sup>+</sup> Treg induction via their enhanced LAP/TGF- $\beta$ 1 production (Fig. 2). Altogether, we showed in a small study set of *Schistosoma*-infected individuals that B cells not only modify effector T-cell responses by reducing their ability to produce pro-inflammatory cytokines, but also increase Treg functions by enhancing the development of FoxP3<sup>+</sup> Tregs and IL-10-producing T cells, with the latter depending on the activity of CD1d<sup>hi</sup> B cells.



**Figure 4.** Induction of IL-10<sup>+</sup> T and FoxP3<sup>+</sup> regulatory (Tregs) cells by B cells. Isolated B cells were cultured at a ratio of 1:1 with allogenic CD4<sup>+</sup>CD25<sup>+</sup> T cells (from 1 European donor) in the presence of soluble anti-CD3 (1 µg/ml) for 6 days. After coculturing, the cells were restimulated with 100 µg/ml PMA and 1 µg/ml ionomycin for 6 hours, in the presence of 10 µg/ml BrefeldinA for the last 4 hours. Cells were then fixed with FoxP3 fixation buffer (eBioscience). (A) T cells were gated for FoxP3<sup>+</sup> T cells and subsequently gated for IL-10<sup>+</sup>(TNF-α) T cells (Tr1 cells). (B) The induction of CD25<sup>hi</sup>FoxP3<sup>hi</sup> T cells after coculture with B cells from endemic infected, endemic uninfected or nonendemic uninfected individuals. The Mann-Whitney *U*-test was used for data analysis. Median values are denoted by horizontal lines, with dots representing individual donors. (C) A representative example of CD1d<sup>hi</sup> depletion, using Miltenyi Biotec isolation beads, is shown. In each group, total untouched B cells (total), mock-depleted total B cells (mock) and CD1d<sup>hi</sup>-depleted B cells were cultured with CD25<sup>+</sup>CD4<sup>+</sup> T cells (See Appendix S3 for uninfected groups). (D) The induction of IL-10<sup>+</sup> T cells, FoxP3<sup>+</sup> Tregs and IL-10 production by B cells after the depletion of CD1d<sup>hi</sup> B cells in *Schistosoma*-infected individuals. We were not able to perform CD1d depletion for all donors because of a limited number of B cells available. Therefore, 4 donors are missing in the *S.haematobium*-infected group (n=10). The Wilcoxon matched pairs test was used to compare data within the group.

## Discussion

In this study, we demonstrated that a greater percentage of B cells from schistosome-infected individuals produce IL-10 and/or LAP/TGF- $\beta$ 1, compared with B cells from uninfected individuals. In coculture with CD4<sup>+</sup> T cells, these B cells reduced the production of effector T-cell cytokines and more FoxP3<sup>+</sup> and IL-10<sup>+</sup> T cells were found, compared to cultures from uninfected controls. It is suggested that IL-10-producing Bregs detected following *in vitro* stimulations may not only be derived from existing circulating IL-10-producing Bregs but may also be derived from progenitor Bregs (25;33). Indeed, it was demonstrated that these progenitor cells can develop into IL-10-competent Bregs during a 2-day *in vitro* stimulation by CD40 ligand and/or BCR ligation (34). Earlier studies exploring Breg frequencies have not investigated the presence of existing IL-10-competent Bregs, as IL-10 production was only studied following long-term *in vitro* stimulation, which does not permit discrimination between these 2 B-cell populations (20;27). Here, we specifically focused on the presence of already competent IL-10 Bregs and found that a greater percentage of peripheral blood B cells from schistosome-infected individuals could readily produce cytoplasmic IL-10, without long-term *in vitro* priming or maturation of B cells. Presumably, these processes already occurred *in vivo*, indicating that schistosome infections indeed prime for the development of readily active IL-10-producing Bregs in humans.

Several studies demonstrated that Bregs may simultaneously produce various regulatory cytokines, such as IL-10 and TGF- $\beta$ . For example, LPS induced both IL-10 and TGF- $\beta$  secretion in B cells, and these LPS-activated B cells suppressed the development of diabetes by promoting apoptosis of effector T cells. However, the responsible cytokine was not addressed in this study (18). Furthermore, in food-allergic mice, allergen-specific T-cell proliferation was suppressed by activated CD5<sup>+</sup> B cells secreting both TGF- $\beta$  and IL-10 (35). Interestingly, similar IL-10- and TGF- $\beta$ -producing CD5<sup>+</sup> B cells were found in healthy individuals upon milk antigen stimulation (7;36). Lastly, co-expression of IL-10 and TGF- $\beta$  was also demonstrated in human CD25<sup>hi</sup>CD1d<sup>hi</sup>CD27<sup>hi</sup> B cells (37). However, other studies have suggested that IL-10 and TGF- $\beta$  are not always expressed by the same Breg subset. For example, CD5<sup>+</sup> B cells from mice chronically exposed to ovalbumin and tolerant for allergic airway inflammation, predominantly expressed IL-10, while TGF- $\beta$  was not restricted to CD5<sup>+</sup> B cells only (8). This is in agreement with the data presented here, as we also observed that IL-10 and LAP/TGF- $\beta$ 1 were expressed by different Breg subsets: the infected group had a greater percentage of CD24<sup>hi</sup>CD27<sup>+</sup> B cells expressing LAP/TGF- $\beta$ 1 and had a greater percentage of CD1d<sup>hi</sup> B cells producing IL-10 compared with the uninfected group. This may suggest that helminth infections can promote the activity of 2 different B-cell subsets.

Enhanced Treg activity is one of the hallmarks of chronic *Schistosoma* infections (38). Because part of schistosome-induced Breg activity in mice is

centered on their capacity to induce Tregs and modulate T-cell cytokines, we focused on these aspects here. However, during a natural infection it is also possible that other cells, such as antigen-presenting cells, may also be influenced by Bregs (25;39). Here, we find a clear induction of FoxP3<sup>+</sup> Tregs and IL-10<sup>+</sup> T cells by schistosome-induced B cells, and we tried to identify the responsible Bregs. Previous studies showed that both IL-10 and TGF- $\beta$  can induce FoxP3<sup>+</sup> Tregs (37;40). However in our study, cocultures with B cells depleted for CD1d<sup>hi</sup> B cells, the main source of IL-10, mainly reduced IL-10<sup>+</sup> T cells without affecting FoxP3<sup>+</sup> T-cell levels. This suggest that the IL-10-producing B cells mainly induced IL-10<sup>+</sup> T cells but not FoxP3<sup>+</sup> Tregs, although we cannot exclude the contribution of other (co)factors to this process. At this stage, it is unclear which schistosome-induced Breg subset is responsible for the enhanced Treg induction, although it is tempting to suggest that, given the role for TGF- $\beta$  in FoxP3<sup>+</sup> Treg induction, TGF- $\beta$ -producing B cells (ie, the CD24<sup>hi</sup>CD27<sup>+</sup> B cells) are involved here.

We previously reported that 6 months after helminth treatment, B cells still showed increased IL-10 secretion upon *in vitro* stimulation with soluble egg antigens (SEA), suggesting the presence of a persistent pool of memory schistosome-specific B cells after clearance of infection (20). Our *S.haematobium*-uninfected donors are coming from a schistosome-endemic area and former infections are likely to have occurred, which may have promoted Breg activity in these individuals at an earlier stage. Indeed, B cells from endemic uninfected group express more IL-10 and TGF- $\beta$  compared to uninfected individuals from a close-by nonendemic area. In addition, this may explain the similar levels of effector T-cell cytokines found in cocultures by B cells from schistosome-infected and uninfected individuals from the same area. Interestingly, the Treg-inducing and IL-10<sup>+</sup> T-cell-inducing capacity was not enhanced in B cells from endemic uninfected group, indicating that some of the activities or some of the Breg subsets are longer lasting after infection, while others may diminish with time.

Several mouse models showed that parasite products from either *Leishmania major* or *Brugia malayi* and the milk glycan LNFPII, containing LeX motifs similar to several schistosome glycoconjugates, can enhance B-cell-derived IL-10 via Toll-like receptor 4 ligation (41;42). Furthermore, *in vitro* stimulation of splenocytes by live schistosome worms induced IL-10-producing B cells, which dampened allergic airway inflammation upon transfer (19). Lastly, B cells from helminth-infected patients with multiple sclerosis stimulated with SEA resulted in a Toll-like receptor 2-mediated IL-10 production (43). These findings suggest that the development of Bregs, as observed in schistosome-infected individuals, may be specifically related to the presence and activity of certain helminth-derived molecules. However, we cannot exclude that Breg development may also occur as a bystander process resulting from continuous chronic infection in which the immune system desperately tries to counterbalance the inflammation induced by the continuous presence of helminths. Indeed, also other chronic infections with viral or bacterial agents show enhanced IL-10-producing B cells (22;24;26). However, what is interesting and what may point to a helminth-

specific Breg population, is that the affected Breg subsets in those studies were not characterized by CD1d<sup>hi</sup>, suggesting that this population is mostly restricted to helminth infections (20;27).

Reports on patients with autoimmune (23;44;45) or allergic diseases (7;36) have shown impairments in Breg frequencies or function, suggesting a role for Bregs in the maintenance of peripheral tolerance in healthy individuals. Therapies aiming to enhance Breg numbers or activity may have potential for future treatment of inflammatory diseases. Interestingly, upregulation of Breg activity was observed in patients with bee venom allergy following successful allergen-specific immunotherapy, reaching similar levels as found in tolerant beekeepers (46). As we here observed that schistosome-induced Bregs can affect T-cell responses (ie, reduced effector T-cell cytokines and more Treg induction), applying helminth-derived molecules may form an interesting novel therapy for hyperinflammatory disorders by driving potent Bregs.

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### **Conflict of interest**

All authors declare no conflict of interest.

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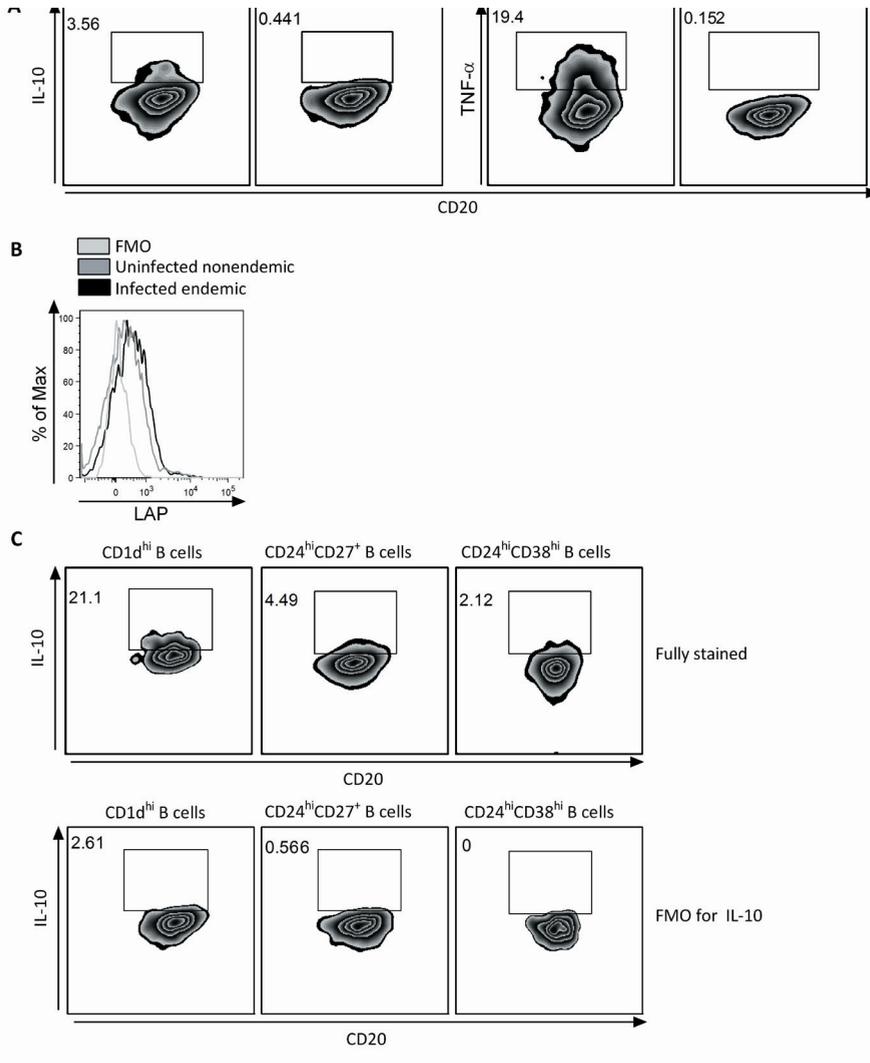
This work was supported by EU-funded project Immunological Interplay between Poverty Related Diseases and Helminth infections: An African-European Research Initiative 'IDEA' [Health-F3-2009-241642], EU-funded project 'The SchistoVac' [Health-2009-242107] and the Deutsche Forschungsgemeinschaft-funded project Deutsch-Afrikanische Kooperationsprojekte in der Infektiologie (DFG-Projekt KR 1150/6-1)

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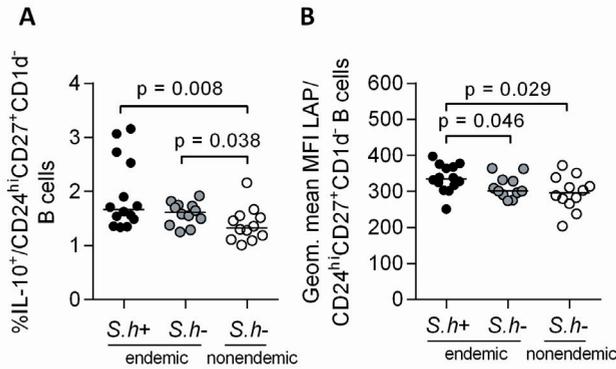
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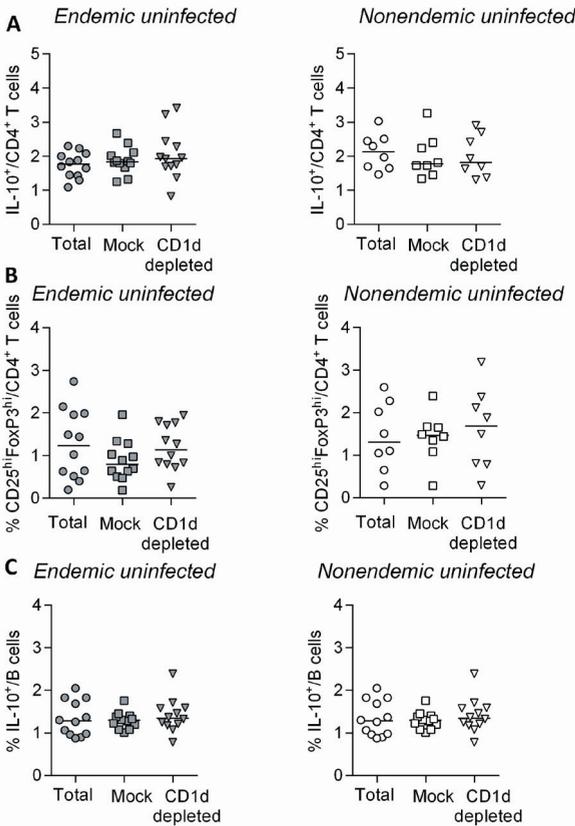
## Supplementary figures



**Figure S1.** (A) A representative gating example of intracellular IL-10 and TNF- $\alpha$  production upon 5 hours PIB+LPS stimulation in total B cells from a endemic infected (*S.h+*) individual. Fluorescence minus one (FMO) shows background levels of IL-10 and TNF- $\alpha$ . B, A representative gating example for LAP/TGF- $\beta$ 1 expression on *ex vivo* total B cells from endemic infected (including FMO) and nonendemic uninfected (*S.h-*) individual. C, Intracellular IL-10 staining upon 5 hours PIB+LPS stimulation in CD1d<sup>hi</sup>, CD24<sup>hi</sup>CD27<sup>+</sup>, CD24<sup>hi</sup>CD38<sup>hi</sup> B cells from a endemic infected individual. Fluorescence minus one (FMO) shows background levels of IL-10 in the different Breg subsets.



**Figure S2.** (A) IL-10 production and (B) LAP/TGF- $\beta$ 1 membrane-bound expression of CD24<sup>hi</sup>CD27<sup>+</sup> B cells after gating out the CD1d<sup>hi</sup> B cells in this population. The Mann-Whitney *U*-test was used for data analysis.



**Figure S3.** Induction of IL-10<sup>+</sup> T and FoxP3<sup>+</sup> Treg cells by B cells. Isolated B cells were cultured at a ratio of 1:1 with allogenic CD25<sup>+</sup>CD4<sup>+</sup> T cells (from 1 European donor) in the presence of soluble anti-CD3 (1  $\mu$ g/ml) for 6 days. After coculturing, the cells were restimulated with 100  $\mu$ g/ml PMA and 1  $\mu$ g/ml ionomycin for 6 hours, in the presence of 10  $\mu$ g/ml BrefeldinA for the last 4 hours. The cells were then fixed with FoxP3 fixation buffer (eBioscience). (A) The induction of IL-10-producing T cells after coculture with B cells from endemic uninfected and nonendemic uninfected donors. (B) The induction of FoxP3<sup>+</sup> Tregs and C, IL-10 production by B cells. We were not able to perform CD1d depletion for all donors due to a limited number of B cells available. Therefore, the total individuals in the endemic uninfected group were  $n=12$  (0 missing), and in the nonendemic uninfected group  $n=8$  (4 missing). For all data, Wilcoxon matched pairs test was used for data analysis.