

***Plasmodium falciparum*-specific interleukin-2 and tumor necrosis factor- α expressing-T cells are associated with resistance to reinfection and severe malaria in healthy African children**

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ABSTRACT. The frequency of *P. falciparum*-specific interleukin (IL)-2-, interferon (IFN)- γ -, tumor necrosis factor (TNF)- α - and IL-10-expressing CD3⁺ cells was studied in healthy Gabonese children segregated according to their clinical presentation at admission to a longitudinal study of severe and mild malaria. The percentage of IL-2- and TNF- α - expressing *P. falciparum*-specific CD3⁺ cells was significantly higher in the children with prior mild malaria and less frequent reinfections compared to the children with prior severe malaria and more frequent reinfections. No differences were shown for *P. falciparum*-specific IFN- γ and IL-10 expression within CD3⁺ cells and parasite-non-specific expression of IL-2, IL-4, IL-6, IL-10, IL-13, TNF- α , and IFN- γ within the CD4⁺, CD8⁺, TCR γ/δ ⁺ CD3⁺ and CD94⁺ CD3⁻ cell populations, indicating that immunological determinants regulating the susceptibility to malaria in age-matched children are parasite-specific. The ability of *P. falciparum*-specific T cells to mount a rapid IL-2 and TNF- α response might be of significance in preventing severe disease and reinfection.

Keywords: T cells, cytokines, *Plasmodium falciparum* malaria

INTRODUCTION

Malaria is responsible for up to two million deaths worldwide per annum [1]. In areas of Sub-Saharan Africa such as the Province of Moyen Ogooué, Gabon, where transmission of *Plasmodium falciparum* is stable and intense [2, 3], mortality is highest in infants younger than six years of age. Severe malaria in these largely non-immune Gabonese children is characterized by hyperparasitemia and severe anemia, with cerebral malaria being less frequently observed. The conditions that regulate the development of mild versus severe malaria in children are still incompletely understood, yet it is the delicate balance between protective and harmful immune responses that finally determines the outcome of malaria [4]. T cells and their cytokines have been implicated in both the pathogenesis of acute malaria attacks as well as in the development of protective immunity, which is, however, acquired only slowly even in endemic areas [5].

The pro-inflammatory cytokines IFN- γ and TNF- α , derived from both the innate and the adaptive arm of the immune system, are released early into the circulation after infection, and high levels have been associated with clinical

symptoms such as fever and malaise as well as with the development of cerebral malaria and severe anemia in susceptible hosts [6-12]. However, IFN- γ and TNF- α are also components of protective immunity and it has been shown in many animal models and some studies in humans that the cellular capacity to produce these cytokines during early phases of infection is essential for the control of parasitemia [13-22]. Thus, it is the relative contribution of these pro-inflammatory cytokines, counterbalanced by the action of anti-inflammatory cytokines such as interleukin (IL)-10 or transforming growth factor (TGF)- β , that may influence the outcome of infection in terms of clinical severity or susceptibility to reinfection. Indeed, in non-immune children from the Lambaréné area, IFN- γ responses to liver-stage antigen 1 (LSA-1) and merozoite surface antigens were associated with resistance to reinfection, whereas in adolescents and adults from western Kenya, IL-10 responses to LSA-1 were predictive of resistance [23, 24]. While this switch from a protective IFN- γ -skewed response in young children to protective IL-10 production in older, malaria-exposed individuals is still to be confirmed, both studies were able to demonstrate correlates of protection using a longitudinal study design.

However, in these studies no information was provided with regard to the cellular sources of the cytokines. In addition, well defined antigens instead of whole parasite preparations were used, which allows for characterization of antigen-specific immune responses but does not reflect so closely the *in vivo* situation when host cells are confronted with a large panel of diverse antigens.

Here we studied the phenotypes and frequencies of cytokine-expressing T cells after *P. falciparum*-specific and non-specific stimulation of PBMC from healthy Gabonese children with a history of either mild or severe *P. falciparum* malaria participating in a longitudinal study. The results obtained were related to initial clinical presentation and subsequent reinfection profiles.

STUDY SUBJECTS AND METHODS

Study site and participants

The study was performed at the Albert Schweitzer Hospital in Lambaréné, Gabon, where *P. falciparum* malaria is predominantly hyperendemic, with an estimated annual entomological inoculation rate of 10-100 [2, 3]. A cohort of 15 healthy children (three males, 12 females; median age, nine years, age range, eight - 14 years) with a history of mild malaria, and of 11 healthy children (three males, eight females; median age, 9.5 years, age range, seven - 14 years), with prior severe malaria were included into the study during spring 2002. All children were recruited from a prospective longitudinal case-control study of severe and mild malaria, details of which have been published previously [23, 25]. Briefly, patients entered the study in 1995 and 1996, severe malaria on admission being defined as hyperparasitemia ($> 250\,000$ parasites/ μL) and/or severe anemia (hemoglobin < 50 g/L) and other signs of severe malaria. Criteria for mild malaria included parasitemia 1000-50 000/ μL on admission, no schizontemia, hemoglobin > 80 g/L, circulating leucocytes containing malarial pigment $> 50/\mu\text{L}$, platelets $> 50/\text{nL}$, leucocytes $< 12/\text{nL}$, lactate < 3 mM, and blood glucose > 50 mg/dL. Reinfections and/or clinical malaria attacks were detected through active clinical and parasitological follow-up of individuals every two weeks following discharge from the hospital, at which times thick blood smears were routinely examined. Therefore, the period of active follow-up after admission of all children with either severe or mild malaria was at least six years. At the time of examination in 2002, all children were clinically healthy and physical examination was unremarkable. Malaria or any other severe illness within the two months prior to the study precluded participation. Blood smears were used to assure that participants were parasite-free. The rate of reinfection per year, calculated during the period of active follow-up of at least six years, was significantly higher in the group of children with prior severe malaria than in the group of children with mild malaria (median, 2.95, range, 1.19-4.79 *versus* median, 0.29, range, 0 - 4.5, $P < 0.001$) as shown by non-parametric analysis (Mann-Whitney U-test). Informed consent was obtained from the parents or guardians of participating children. Ethical clearance was given by the ethics committee of the Albert Schweitzer Hospital in Lambaréné.

Parasite preparation

P. falciparum strain S007, originally isolated from a child with severe malaria in Lambaréné, was cultured in human type 0 erythrocytes, adjusting the hematocrit to 5% and the parasitemia to 2-5% in complete parasite medium (CPM: RPMI-1640 supplemented with 25 mM HEPES, 2 mM L-glutamine, 50 $\mu\text{g}/\text{mL}$ of gentamicin, 0.5% Albumax II [Gibco, Paisley, UK], with 200 μM of hypoxanthine, and 2% AB⁺ serum). All cultures and media were regularly tested for *Mycoplasma* contamination by PCR amplification with genus-specific primers (GPO-1 5'-ACT CCT ACG GGA GGC AGC AGT A-3' and MGSO 5'-TGC ACC ATC TGT CAC TCT GTT AAC CTC-3') of 16S rDNA as described previously [26].

For synchronization and enrichment of *P. falciparum*-infected erythrocytes (PFE) a magnet-activated cell sorter (MACS) system was applied as described recently [27]. Parasite concentration could be increased up to 90% using CS columns and VarioMACS (Miltenyi BioTec, Bergisch-Gladbach, Germany). Yield and purity were assessed by microscopic examination of Giemsa - stained eluate and flow - through. The MACS-eluate of *P. falciparum* strain S007 was frozen at -80 °C, thawed once and used for stimulation of the PBMC from all children.

Detection of Plasmodium falciparum-specific T cell cytokine expression by flow cytometry

PBMC were isolated from heparinized blood by ficoll-diatrizoate centrifugation and plated out in 12-well plates at 2.5×10^6 /well. Cells were cultured in Ultra Culture Medium (UCM) (Bio Whittaker, Walkersville, MD, USA) supplemented with L-glutamine (2 mM/L; Sigma, St. Louis, MO, USA), gentamicin (170 mg/L; Sigma) and 2-mercaptoethanol (3.5 $\mu\text{L}/\text{L}$; Merck, Darmstadt, Germany) for 18 hours at 37 °C in 5% CO₂, and stimulated with or without 50 μL of the MACS-eluate containing the late-stage, schizont-rich parasite preparation at a ratio of 10:1 PFE:PBMC as described recently [28, 29]. To enhance the Ag-specific response, the co-stimulatory MAb CD28 (PharMingen, San Diego, CA, USA) was added at 10 μL to some wells (5 $\mu\text{g}/\text{mL}$ final concentration) [29, 30]. Uninfected erythrocytes were processed as described above and served as controls (with and without the addition of MAb CD28). Further controls included the addition of MAb CD28 to the PBMC preparation without PFE, and medium alone without MAb CD28 or PFE. MAb CD28 in the absence of Ag had no effect on cellular cytokine responsiveness, however, it increased the frequency of PFE-induced cytokine - expressing cells up to twofold as reported previously [29].

Brefeldin A (1 μM ; 10 $\mu\text{g}/\text{mL}$ final concentration, Sigma) was added after six hours to block protein secretion. Cells were then harvested on ice without scraping, washed twice in phosphate-buffered saline (PBS) and fixed with 2% formaldehyde (1mL per 2×10^6 cells, Merck) for 20 min. After two additional washes in PBS, cells were then resuspended in Hank's balanced salt solution (HBSS, supplemented with 0.3% bovine serum albumine [BSA] and 0.1% sodium-azide), and stored at 4 °C in the dark until staining. Fixed cells were washed twice with PBS and made permeable with saponin (0.1%; Sigma). They were then resuspended with 50 μL of saponin-buffer-diluted

antibodies and incubated for 25 min at room temperature in the dark.

The following monoclonal antibodies were used: cytokine-specific mouse anti-human monoclonal antibody (MAb) (IFN- γ [clone: B27], fluorescein-isothiocyanate [FITC]-labeled) and rat anti-human MAb (IL-2 [MQ1-17H12], phycoerythrin [PE]-conjugated; IL-10 [JES3-9D7], PE-labeled; TNF- α [Mab11], PE-labeled). All MAb were purchased from Pharmingen. The anti-CD3-MAb was peridinin chlorophyll (PerCP), the anti-CD69-MAb was allophycocyanin (APC)-labeled (Becton Dickinson, Mountain View, CA, USA). Four-colour staining was performed and at least 10⁴ cells were analyzed on a FACSCalibur (Becton Dickinson) equipped with a two laser system (488 nm and 633 nm wavelength, respectively). All cytokine combinations (IL-2/IFN- γ , IL-10/IFN- γ , TNF- α /IFN- γ) were stained in conjunction with CD3 and CD69.

T cells were defined by their side-scatter characteristics and anti-CD3 MAb staining. The specificity of cytokine staining in CD3⁺ cells was verified by counterstaining with CD69 as a marker for activated lymphocytes. Only cells clearly positive for CD69 were classified as cytokine-producing CD3⁺ cells. Data were analysed with CELLQuest software (Becton Dickinson), and results were expressed as the percentage of cytokine-producing cells in each CD3⁺ CD69⁺ cell population. Specificity of cytokine staining was confirmed by the absence of significant background in controls with isotype-matched irrelevant MAbs. Frequencies of background events within the CD3⁺ cell population (unstimulated cells; addition of MACS-processed uninfected erythrocytes with or without anti-CD28) were < 0.03%. These values were not subtracted from those obtained after stimulation with PFE.

Detection of non-specific, mitogen-induced T cell cytokine expression by flow cytometry

PBMC were isolated from heparinized blood and stimulated in UCM with phorbol 12-myristate 13-acetate (PMA, 10 ng/mL; Sigma) and ionomycin (1.25 μ M; Sigma) in the presence of brefeldin A (1 μ M; Sigma) for 4 hours at 37° C in 5% CO₂. Cells were then harvested and fixed as described above. For the staining procedure, the following monoclonal antibodies (MAbs) were used: cytokine-specific mouse anti-human MAb (IFN γ [clone B27], fluorescein isothiocyanate [FITC]-conjugated); rat anti-human MAb (IL-2 [MQ1-17H12], IL-4 [MP4-25D2], IL-6 [MQ2-13A5], IL-10 [JES3-9D7], IL-13 [JES10-5A2], TNF- α [Mab11], all phycoerythrin [PE]-conjugated), the anti-CD4 MAb, anti-TCR γ/δ MAb and anti-CD94 MAb

were allophycocyanin-conjugated, anti-CD3 MAb and anti-CD8 MAb were peridinin chlorophyll-conjugated; all cytokine-specific MAbs were purchased from Pharmingen (San Diego, CA, USA), the surface marker-specific MAbs from Becton Dickinson (Mountain View, CA, USA).

Samples were gated on lymphocytes according to their light scatter characteristics and the results were expressed as the percentage of cytokine-producing cells in the CD4⁺, CD8⁺, TCR γ/δ ⁺ CD3⁺ or CD94⁺ CD3⁻ cell population, respectively.

Statistical methods

Statistical analysis was performed using a standard statistical package (SPSS 10.0 for Windows; SPSS Inc., Chicago). The Mann-Whitney U-test was applied for group differences (children with prior mild *versus* severe malaria, children with < one reinfection *versus* children with > two reinfections per year). A P value of < 0.05 was considered significant.

RESULTS

Differences in the frequency of cytokine-expressing P. falciparum-specific CD3⁺ cells between healthy children with a history of either severe or mild malaria

The frequency of cytokine-expressing CD3⁺ cells was compared between 11 children with prior severe malaria and 15 children with prior mild malaria. At the time of PBMC stimulation, all patients were apparently healthy and free of parasites.

The frequency of CD3⁺ cells exclusively expressing IL-2 and TNF- α , as well as the frequency of CD3⁺ cells expressing IL-2 together with IFN- γ , were significantly higher in the mild malaria group when compared to the group with severe malaria (Figures 1, 2). Likewise, the frequencies of overall IL-2- and TNF- α -expressing CD3⁺ cells were significantly higher in the mild malaria than in the severe malaria group (Table 1, Figure 1). As shown previously [31, 32], most IL-10-expressing cells also stained positively for IFN- γ , however, no differences were observed between groups (Figure 1).

Frequency of cytokine-expressing, non-specifically stimulated PBMC in healthy children with prior severe or mild malaria

The frequencies of IL-2-, IL-4-, IL-6-, IL-10-, IL-13-, IFN- γ -, and TNF- α - expressing CD4⁺, CD8⁺, TCR γ/δ ⁺ CD3⁺ and CD94⁺ CD3⁻ cell populations after mitogenic

Table 1
Frequency of *P. falciparum*-specific cytokine expression within the CD3⁺ cell population in healthy children with prior mild or severe malaria*

Cytokines	Prior mild malaria (n = 15)	Prior severe malaria (n = 11)
IL-2 ⁺ (IL-2 ⁺ /IFN- γ ⁺ and IL-2 ⁺ /IFN- γ ⁻)	0.090 \pm 0.012 (0.02-0.17) ^{1**}	0.047 \pm 0.006 (0.02-0.09)
IFN- γ ⁺ (IL-2 ⁺ /IFN- γ ⁺ and IL-2 ⁻ /IFN- γ ⁺)	0.077 \pm 0.015 (0.01-0.19)	0.060 \pm 0.008 (0.03-0.11)
TNF- α ⁺ (TNF- α ⁺ /IFN- γ ⁺ and TNF- α ⁺ /IFN- γ ⁻)	0.140 \pm 0.022 (0.03-0.28) ²	0.074 \pm 0.014 (0.02-0.18)
IL-10 ⁺ (IL-10 ⁺ /IFN- γ ⁺ and IL-10 ⁺ /IFN- γ ⁻)	0.023 \pm 0.006 (0.00-0.11)	0.035 \pm 0.010 (0.00-0.11)

*PBMC were stimulated with MACS-separated, late stage *Plasmodium falciparum* isolates in the presence of mAb CD28.

**Values indicate mean percentages of cytokine-expressing CD3⁺ cells \pm SEM; the respective ranges are given in parentheses.

¹ significant difference between the children with prior mild or severe malaria, P < 0.01 as calculated by the Mann-Whitney U-test.

² significant difference between the children with prior mild or severe malaria, P < 0.05 as calculated by the Mann-Whitney U-test.

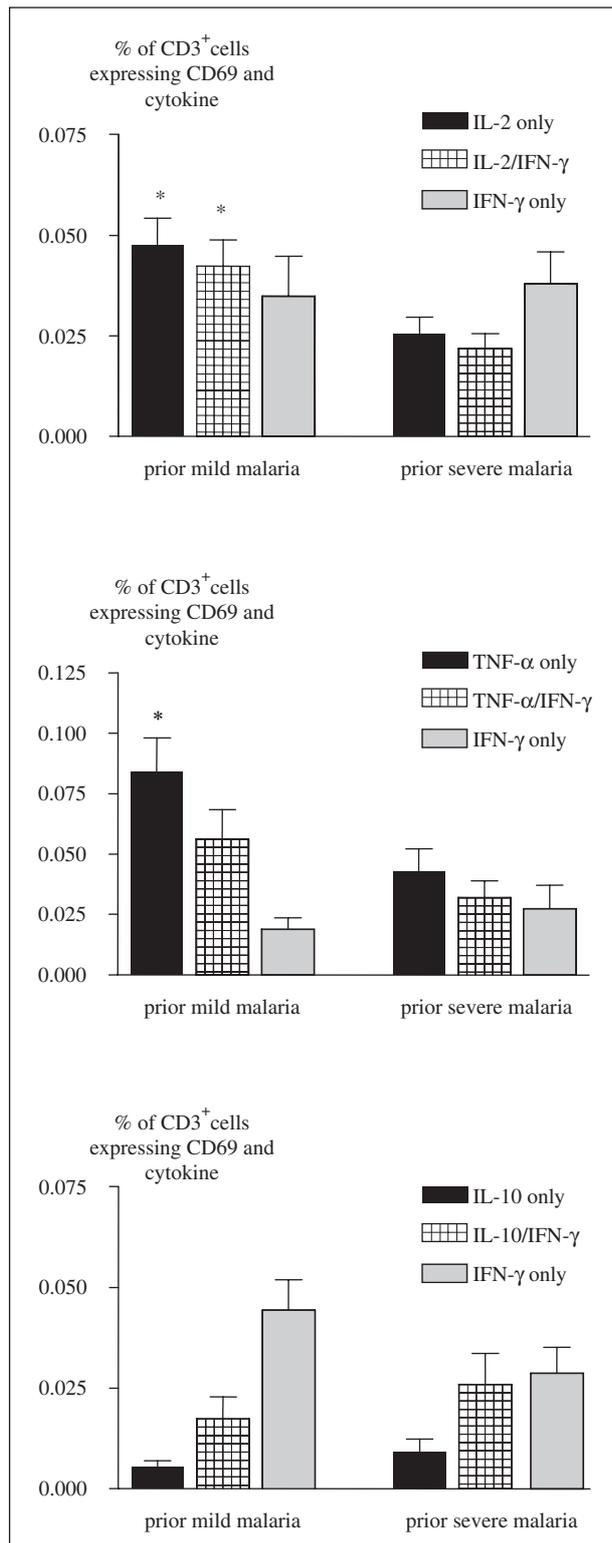


Figure 1

Frequency of cytokine-expressing, activated (CD69⁺) CD3⁺ cells obtained after stimulation with MACS-processed PFE in children with prior mild malaria compared to children with prior severe malaria. Each bar indicates the mean percentages \pm SEM. Values that are statistically different between groups are denoted by asterisks (* $P < 0.05$, as calculated by the Mann-Whitney U-test).

stimulation with PMA and ionomycin in the presence of brefeldin A, were not significantly different between the two groups with different initial clinical presentation (Tables 2, 3; Figure 3). This was apparent for both the

overall production of each cytokine as well as for subsets defined by their characteristic cytokine (co-) expression patterns. Without the addition of mitogenic stimuli, no expression of cytokines was detectable in any cell population studied.

Differences in the frequency of cytokine-expressing *P. falciparum*-specific CD3⁺ cells and of non-specifically stimulated PBMC between healthy children segregated according to their reinfection profile

When participating children with < one reinfection per year ($n = 14$, all from the mild malaria group) were compared to children suffering from > two reinfections annually ($n = 9$, all from the severe malaria group), similar significant differences of cytokine expression profiles were found as shown for the groups with different clinical presentation. This was not surprising as patients with prior severe malaria had acquired many more reinfections per year than the children from the mild malaria group, during the follow-up period of six years. As shown for the groups with the history of either severe or mild malaria at admission, no significant differences with regard to the frequencies of IL-2-, IL-4-, IL-6-, IL-10-, IL-13-, IFN- γ -, and TNF- α - expressing CD4⁺, CD8⁺, TCR γ/δ ⁺ CD3⁺ and CD94⁺CD3⁻ cell populations after non-specific mitogenic stimulation were found when children were segregated according to their reinfection rate.

DISCUSSION

The results of this study point towards a role of IL-2- and TNF- α - expressing *P. falciparum*-specific CD3⁺ cells in the determination of differences in clinical presentation and subsequent reinfection profiles. CD3⁺ cells of the children who initially presented with mild malaria and who experienced less frequent reinfections during the years of follow-up were more responsive to *P. falciparum*-specific stimulation with increased expression of IL-2 and TNF- α when compared to children with initial severe malaria and more frequent reinfections. This was not only true for the overall capacity of IL-2 and TNF- α production but also for those CD3⁺ cells exclusively expressing the respective cytokines (single IL-2 and TNF- α expression). All children were parasite-free and healthy at the time of analyses, thus results were not confounded by disease- or inflammation-associated cellular modulation, which we and others have reported to be present during acute *P. falciparum* malaria [31-34]. In addition, using non-specific mitogenic stimulation with intracellular detection of cytokines, we have shown a clear age-dependency in immune responsiveness of patients with acute malaria [32]. In the present study of age-matched healthy children, no differences in cytokine expression in the CD4⁺, CD8⁺, TCR γ/δ ⁺CD3⁺ and CD94⁺CD3⁻ cell populations were found between both groups when PBMC were non-specifically stimulated. Thus, we suppose, at least in these children and for the subsets studied, that the immunological determinants resulting in different outcomes of infection with respect to clinical severity and reinfection rates are *P. falciparum*-specific. Early type 1 cytokine production during *Plasmodium* infection has been linked to successful control of parasitemia in animal models of malaria [13-20]. This has been attributed mainly to the action of

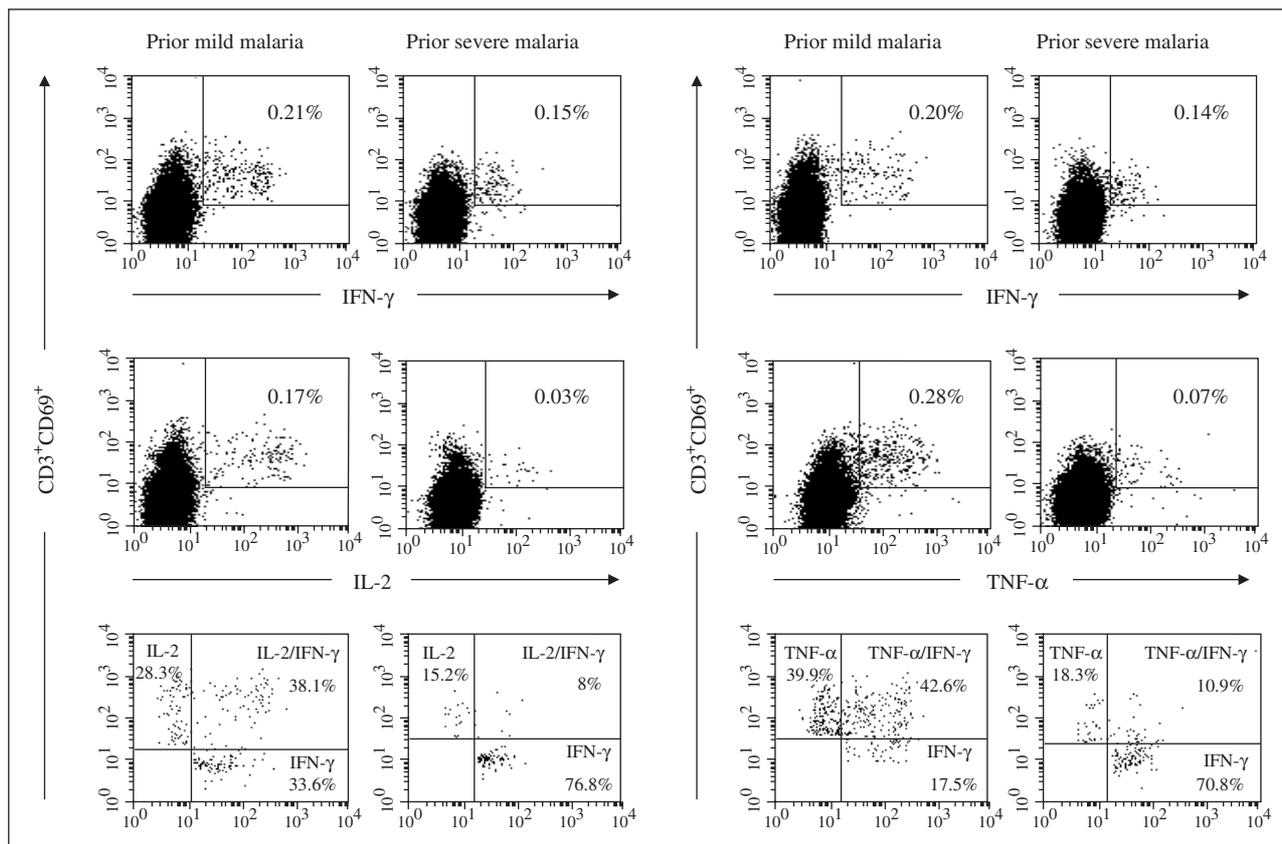


Figure 2

Two-parameter dot plots indicating the frequency of IL-2-, IFN- γ - and TNF- α - expressing *P. falciparum*-specific activated (CD69⁺) CD3⁺ T cells of one representative child with prior mild and one child with prior severe malaria. Note the marked differences in the expression of IL-2 and TNF- α between the two children (second horizontal column). The third horizontal column displays the cytokine distribution pattern of IFN- γ together with IL-2 or TNF- α . PBMC were stimulated with a MACS-enriched *P. falciparum* late stage parasite preparation and the co-stimulatory MAb anti-CD28. Numbers indicate the percentage of CD3⁺ cells positive for the respective cytokine (first and second horizontal column) or CD3⁺ cells co-expressing two cytokines (third horizontal column). Corresponding data and statistics are depicted in Table 1, as well as in Figure 1.

Table 2
Frequency of mitogen-induced cytokine expression within the CD4⁺ and CD8⁺ T cell subsets in healthy children with prior mild (n = 15) or severe malaria (n = 11)¹

Cytokines	% of CD4 ⁺		% of CD8 ⁺	
	Prior mild malaria	Prior severe malaria	Prior mild malaria	Prior severe malaria
IFN- γ only	3.5 \pm 0.3 (1.6-5.9)*	3.2 \pm 0.3 (1.9-5.2)	17.0 \pm 2.1 (8.9-34.3)	17.5 \pm 2.6 (4.7-35.0)
IL-2/IFN- γ	9.9 \pm 0.8 (4.5-15.8)	9.2 \pm 1.0 (5.9-17.4)	8.3 \pm 0.8 (3.7-14.7)	8.9 \pm 1.5 (3.0-18.8)
IL-2 only	38.9 \pm 1.1 (31.1-44.5)	40.3 \pm 1.5 (30.8-47.2)	6.9 \pm 0.5 (4.1-11.6)	6.4 \pm 0.6 (4.2-9.4)
IL-4 only	4.3 \pm 0.4 (2.0-8.0)	4.7 \pm 0.5 (2.0-7.0)	< 1	< 1
IL-4/IFN- γ	1.6 \pm 0.2 (0.5-2.7)	2.0 \pm 0.2 (1.0-3.4)	< 1	< 1
IL-6 only	< 1	< 1	< 1	< 1
IL-6/IFN- γ	< 1	< 1	< 1	< 1
IL-10 only	< 1	< 1	< 1	< 1
IL-10/IFN- γ	< 1	< 1	< 1	< 1
IL-13 only	3.8 \pm 0.3 (2.0-6.3)	4.3 \pm 0.4 (2.3-6.8)	< 1	< 1
IL-13/IFN- γ	< 1	< 1	< 1	< 1
TNF- α only	26.5 \pm 1.6 (16.4-41.2)	28.5 \pm 1.9 (15.9-36.5)	1.2 \pm 0.1 (0.5-2.5)	1.3 \pm 0.2 (0.2-2.6)
TNF- α /IFN- γ	11.9 \pm 0.8 (8.4-18.2)	10.9 \pm 1.1 (6.4-20.1)	16.1 \pm 2.1 (6.1-36.3)	17.8 \pm 2.9 (4.7-38.9)

*Values indicate mean percentages of cytokine-expressing CD4⁺ and CD8⁺ \pm SEM; the respective ranges are given in parentheses. PBMC were stimulated with PMA and ionomycin in the presence of brefeldin A.

¹Differences between groups reached no significance as calculated by the Mann-Whitney U-test.

Table 3
Frequency of mitogen-induced cytokine expression within the CD3⁺TCRγδ⁺ and CD3⁻CD94⁺ cell populations in healthy children with prior mild (n = 15) or severe malaria (n = 11)¹

Cytokines	% of CD3 ⁺ TCRγδ ⁺		% of CD3 ⁻ CD94 ⁺ (NK cells)	
	Prior mild malaria	Prior severe malaria	Prior mild malaria	Prior severe malaria
IFN-γ only	34.7 ± 3.0 (18.6-615)	25.6 ± 4.7 (7.5-590)	47.2 ± 3.7 (13.8-681)	43.2 ± 4.9 (14.9-74.2)
IL-2/IFN-γ	6.7 ± 0.6 (2.5-2.2)	4.9 ± 0.6 (1.3-8.7)	< 1	1.2 ± 0.3 (0.4-4.5)
IL-2 only	7.9 ± 1.1 (2.4-9.0)	9.9 ± 1.3 (4.0-7.2)	< 1	< 1
IL-4 only	< 1	< 1	< 1	< 1
IL-4/IFN-γ	1.3 ± 0.2 (0.3-2.9)	< 1	< 1	< 1
IL-6 only	< 1	< 1	< 1	< 1
IL-6/IFN-γ	< 1	< 1	< 1	< 1
IL-10 only	< 1	< 1	< 1	< 1
IL-10/IFN-γ	< 1	< 1	< 1	< 1
IL-13 only	< 1	< 1	< 1	< 1
IL-13/IFN-γ	< 1	< 1	< 1	< 1
TNF-α only	4.8 ± 0.7 (1.4-10)	4.3 ± 1.1 (1.0-14.9)	8.0 ± 1.0 (3.1-15.0)	8.3 ± 1.6 (1-20.0)
TNF-α/IFN-γ	31.8 ± 3.8 (12.8-69.7)	23.9 ± 4.5 (6.8-48.4)	20.9 ± 2.5 (5.3-39.3)	19.6 ± 3.3 (12.2-45.3)

*Values indicate mean percentages of cytokine-expressing CD3⁺TCRγδ⁺ and CD3⁻CD94⁺ (NK cells) ± SEM; the respective ranges are given in parentheses. PBMC were stimulated with PMA and ionomycin in the presence of brefeldin A.

¹ Differences between groups reached no significance as calculated by the Mann-Whitney U-test.

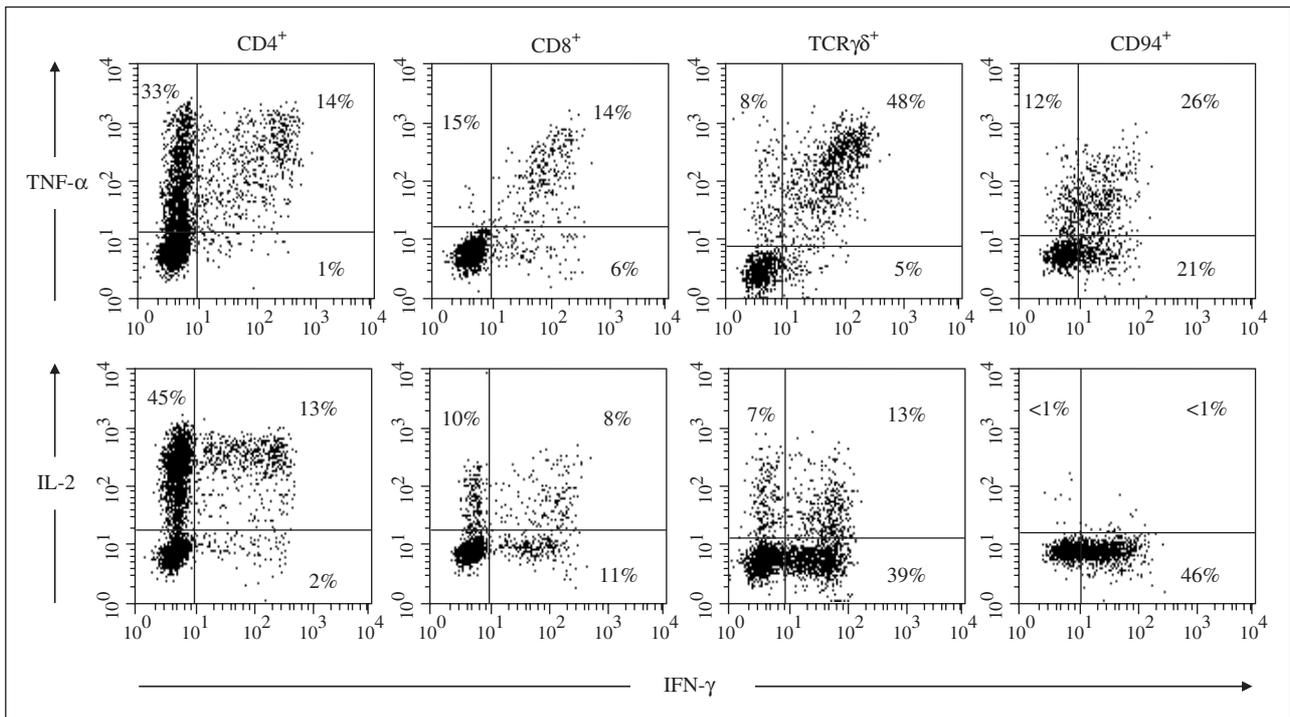


Figure 3

Representative two-parameter dot plots indicating the profile of IL-2-, IFN-γ- and TNF-α- expressing CD4⁺, CD8⁺, TCRγδ⁺CD3⁺ and CD94⁺CD3⁻ cell populations from one child following non-specific stimulation with PMA and ionomycin in the presence of brefeldin A. Similar characterizations were performed for the cytokines IL-4, IL-6, IL-10, and IL-13. The numbers in each quadrant represent the percentage of gated cytokine-expressing cells with the right upper quadrant including cells co-expressing two cytokines. Corresponding data and statistics are depicted in Tables 2 and 3.

IFN-γ and TNF-α. Indeed, IFN-γ was associated with resistance to reinfection in a cohort of children (including those children studied here), when PBMC were stimulated with defined liver-stage and blood-stage antigens [23]. After several years of intense follow-up, only a selection of these children were still available for the present examination, nevertheless much care has been taken to assure comparability of groups with regard to age, socioeconomic

status and parasitological factors such as the estimated annual entomological inoculation rate [2, 35]. As we were unable to find differences in the expression of IFN-γ between groups, we suggest that in older children, other cytokines such as IL-2 and TNF-α might play important roles in the protection against reinfection as well. Some evidence for an age-dependent switch of the type of cytokines involved in antimalarial host defenses comes from a

study of adult Kenyans, where IL-10 and not IFN- γ responses to liver-stage antigen-1 were associated with protection against reinfection [24].

IL-2, a product predominantly of CD4⁺ cells, has originally been described as T cell growth factor [36]. The effects of IL-2 on its cell targets include mitogenesis in type 1 cells, stimulation of cytotoxicity in cytotoxic T lymphocytes and NK cells, as well as generation of lymphokine-activated killer cells. Its role during human malaria has not been very well exploited, as IL-2 is barely detectable in plasma, and most studies focused on the contribution of other cytokines to the pathogenesis of malaria infection. Earlier studies showed a parasite-specific and non-specific defect in IL-2 production during acute *P. falciparum* malaria [37-39], which was thought to provide the basis of malaria-associated cellular immune hyporesponsiveness. By using non-specific stimulation of PBMC from Gabonese patients with uncomplicated malaria, we have found preserved capacity for IL-2 production by CD4⁺ cells, which, however, was downregulated during clearance of parasitemia [31]. In addition, while IL-2 is termed a classical type 1 cytokine together with IFN- γ , we have shown that both cytokines are reciprocally regulated during the course of an acute *P. falciparum* infection. A role for CD4⁺ T cell-derived IL-2 has been shown for the activation of human TCR γ/δ ⁺ cells when stimulated with freeze-thawed schizont extracts [40-42]. In addition, treatment of *P. yoelii*-infected mice with IL-2 preferentially accumulated TCR γ/δ ⁺ cells in their brains and rendered them susceptible to cerebral malaria [43]. Therefore, a link between adaptive and innate responses mediated by IL-2 can be suggested. However, while the role of early IL-2 expression in response to *P. falciparum* infection remains rather speculative until now, more precise data are available with regard to early TNF- α production.

The release of TNF- α in response to parasites has been shown to be dependent upon the presence of CD3⁺ cells and monocytes [44]. TCR γ/δ ⁺CD3⁺ cells have been identified as the major lymphocyte subset involved in this early TNF- α production [28]. Our previously obtained data and those shown here however, ascribe considerable capacity for TNF- α production to other T cell subsets [29]. "Appropriate" and early TNF- α production appears to be protective and necessary to limit parasitemia [21, 22], while high levels in plasma have been linked to cerebral malaria and in the presence of low IL-10 levels, with severe anemia in African children [6, 7, 10, 11, 22, 45]. An increased capacity of TNF- α production might explain the greater capacity of PBMC from children with prior mild malaria to express nitric oxide (NO) synthase and to produce NO [46]. This may allow a more efficient reduction of parasitemia, thus preventing progression to severe disease in this group of children. Anti-inflammatory activities, here represented by both specifically and non-specifically induced IL-10 were not different between groups, neither were the non-specifically induced classical type 2 cytokines IL-4 and IL-13. Most IL-10-expressing CD3⁺ cells also co-expressed IFN- γ , suggesting an immune-regulatory role in response to stimulation with *P. falciparum* [47]. Intracellular IL-6 was not detected in either T cells or within the NK cell subset.

In all, we have detected considerable frequencies of cytokine-expressing, activated T cells in healthy children

upon specific stimulation with late-stage *P. falciparum* parasite preparations. Mean frequencies of responding CD3⁺ T cells were 1/5000 after staining for IL-10, and 1/700 cells expressing TNF- α , which is comparable to the precursor frequencies previously obtained *in vitro* after six days of incubation with *Plasmodium*-infected erythrocytes [48]. Marked differences were shown for the frequency of IL-2- and TNF- α -producing *P. falciparum*-specific activated T cells between children with prior mild malaria and less frequent reinfections on the one hand and children with initial severe malaria and more frequent reinfections on the other, pointing towards a protective role of both cytokines in host defense against malaria. Studies are underway that examine specific cellular cytokine responses both during acute malaria as well as in individuals with different degrees of immunity to *P. falciparum*.

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