Altering in cytokines and haematological parameters during the acute and convalescent phases of *Plasmodium falciparum* and *Plasmodium vivax* infections

Rodrigo Nunes Rodrigues-da-Silva¹, Josué da Costa Lima-Junior¹, Bruna de Paula Fonseca e Fonseca², Paulo Renato Zuquim Antas³, Arlete Baldez³, Fabio Luiz Storer⁴, Fátima Santos⁵, Dalma Maria Banic⁶, Joseli de Oliveira-Ferreira¹/*

¹Laboratório de Imunoparasitologia ¹Laboratório de Imunologia Clínica ¹Laboratório de Simulídeos e Oncocercose, Instituto Oswaldo Cruz ¹Laboratório de Tecnologia Diagnóstica, Bio-Manguinhos-Fiocruz, Rio de Janeiro, RJ, Brasil ¹Agência de Vigilância em Saúde, Secretaria de Estado da Saúde, Porto Velho, RO, Brasil ³Faculdade São Lucas, Porto Velho, RO, Brasil ³Odebrecht Energia/Usina Hidrelétrica Santo Antônio, Porto Velho, RO, Brasil

Haematological and cytokine alterations in malaria are a broad and controversial subject in the literature. However, few studies have simultaneously evaluated various cytokines in a single patient group during the acute and convalescent phases of infection. The aim of this study was to sequentially characterise alterations in haematological patterns and circulating plasma cytokine and chemokine levels in patients infected with *Plasmodium vivax* or *Plasmodium falciparum* from a Brazilian endemic area during the acute and convalescent phases of infection. During the acute phase, thrombocytopenia, eosinopenia, lymphopenia and an increased number of band cells were observed in the majority of the patients. During the convalescent phase, the haematologic parameters returned to normal. During the acute phase, *P. vivax* and *P. falciparum* patients had significantly higher interleukin (IL)-6, IL-8, IL-17, interferon-γ, tumour necrosis factor (TNF)-α, macrophage inflammatory protein-1β and granulocyte-colony stimulating factor levels than controls and maintained high levels during the convalescent phase. IL-10 was detected at high concentrations during the acute phase, but returned to normal levels during the convalescent phase. Plasma IL-10 concentration was positively correlated with parasitaemia in *P. vivax* and *P. falciparum*-infected patients. The same was true for the TNF-α concentration in *P. falciparum*-infected patients. Finally, the haematological and cytokine profiles were similar between uncomplicated *P. falciparum* and *P. vivax* infections.

Key words: cytokines - chemokines - platelets - *P. falciparum* - *P. vivax*

Malaria remains a major health problem worldwide, with 300-500 million cases annually and nearly one million deaths (Murray et al. 2012). Although *Plasmodium falciparum* malaria represents the majority of these cases and is responsible for almost all of the associated mortality, *Plasmodium vivax* malaria has a wider geographic distribution and is responsible for high morbidity worldwide. Despite this widespread prevalence, *P. vivax* has long been overshadowed by the burden caused by *P. falciparum*. In Brazil, *P. vivax* accounts for more than 70% of all malaria cases. The infections are chronic, can produce profound anaemia, can be incapacitating for days or weeks and have the added complication of recurrent clinical episodes due to the developmental reactivation of hypnozoites, the dormant liver stage form (Mendis et al. 2001, Sina 2002).

It is well documented that disease severity depends strongly on the previous immunological experience of the host (Schofield & Mueller 2006). Therefore, in areas of high malaria transmission, infants and young children are more frequently affected, whereas in malaria-endemic areas, where transmission is unstable, adults are the most commonly affected population (Schofield & Mueller 2006). Although sterile immunity is most likely never achieved, individuals can develop essentially complete protection from severe illness and death after continuous exposure (Langhorne et al. 2008). Clinical malaria infection causes a range of symptoms from asymptomatic infection to severe disease complication. Although different theories have been proposed to explain the disease in humans, malaria pathogenesis remains controversial (Miller et al. 2002, Weatherall et al. 2002).

Although there is an extensive body of literature describing variations in haematological parameters and immune cytokine responses during malaria infection, their link to disease manifestation is still a subject of much debate. The subject of haematological changes is controversial in the malaria field, although anaemia and thrombocytopenia are the most prominent alterations during both *P. falciparum* and *P. vivax* infections (Agarwal et al. 1983, Lacerda et al. 2011). Severe malaria has long been associated with high circulating levels of inflammatory cytokines such as tumour necrosis factor (TNF)-α, interleukin (IL)-1 and IL-6. Studies have demonstrated a link between TNF-α, IL-6, IL-10 and the severity of the disease in human malaria (Akanmori et
al. 2000) and high plasma levels of these cytokines are found at diagnosis, even in uncomplicated malaria cases (Baptista et al. 1997). Anti-inflammatory cytokines have also been found to have important roles in the immune response against *Plasmodium*. IL-10 has an important role as an immunoregulator during *P. falciparum* infection, neutralising the effects of other cytokines produced by T-helper (Th1) and CD8 cells (Couper et al. 2008, Langhorne et al. 2008). In recent years, increasing evidence has suggested that regulatory T cells are key anti-inflammatory cells that are critical to limiting the inflammatory response (Hansen & Schofield 2010). Additionally, IL-10 and granulocyte-colony stimulating factor (G-CSF) have been found to be elevated and correlated with parasitaemia in asymptomatic pregnant women in Ghana (Wilson et al. 2010), suggesting that these cytokines may act to reduce symptoms. Paradoxically, a significant number of cytokines and chemokines have been associated with severe disease, in particular IL-1β (Brown et al. 1999), IL-2 (Ramharter et al. 2003), IL-6 (Kern et al. 1989, el-Nashar et al. 2002), interferon (IFN)-γ (Day et al. 1999, Ramharter et al. 2003), TNF-α (Kern et al. 1989, Kwiatkowski et al. 1990, el-Nashar et al. 2002, Ramharter et al. 2003), IL-4 (Kumaratilake & Ferrante 1992, Eisenhut 2010), IL-10 (Day et al. 1999, Othoro et al. 1999, Ramharter et al. 2003) and macrophage inflammatory protein (MIP)-1β (Ochiel et al. 2005), whereas low levels of regulatory cytokines, such as TGF-β and IL-10, have been correlated with acute malaria (Peyron et al. 1994, Hansen & Schofield 2010). Studies in Brazil, where malaria is predominantly hypo or mesoendemic and the incidence of complications and mortality due to malaria infections is very low, increased IFN-γ concentrations have been reported in *P. vivax* and *P. falciparum* patients, whereas increased IL-10 levels have been observed only in *P. vivax* patients (Medina et al. 2011). A comparison between asymptomatic individuals with those experiencing severe malaria showed that the IFN-γ/IL-10 ratio was higher in severe cases, whereas IL-10 levels were elevated in asymptomatic individuals (Andrade et al. 2010). Another study in the same area showed a significant increase in the concentrations of TNF-α, IFN-γ, migration inhibitory factor and monocyte chemotactic protein (MCP)-1 in patients with *P. vivax* and *P. falciparum*, whereas IL-10 was observed in only *P. vivax*-infected patients (Fernandes et al. 2008).

A retrospective analysis of a database containing clinical epidemiological data from patients with mild or severe malaria from the state of Rondônia (RO), Brazil revealed that systematic analysis of several inflammatory mediators that are measured simultaneously can characterise the overall immune response pattern of patients infected with *P. vivax*. In the group of patients with severe malaria, *P. vivax* parasitaemia had several positive interactions with inflammatory mediators such as TNF-α, IFN-γ and IL-10 (Mendonça et al. 2013).

Although clinical descriptions of the illness caused by *P. vivax* infection are available, data regarding inflammatory patterns are scarce and few studies have evaluated numerous different cytokines in a single patient group in paired samples using one methodology. A better approach for investigating putative relationships between cytokine production and disease state is to simultaneously profile temporal changes in multiple cytokines. Therefore, in this study, we determined the concentrations of various cytokines and chemokines directly from plasma in an endemic population infected with *P. falciparum* and *P. vivax* as part of an effort to identify a single marker or combination of cytokines and chemokines that could potentially characterise differences between the acute and convalescent phases (pre and post-treatment time points) of infection and to discriminate between *P. falciparum* and *P. vivax* infections.

**SUBJECTS, MATERIALS AND METHODS**

**Subjects and sampling** - The study was carried out in Porto Velho, RO, an unstable malaria-endemic area, where *P. vivax* accounts for more than 75% of all malaria cases (Oliveira-Ferreira et al. 2010). Symptomatic patients diagnosed with malaria infection by a thick blood smear in an outpatient clinic in Porto Velho were asked to participate in the study. A total of 71 patients were enrolled for the study, 47 and 24 of whom were infected with *P. vivax* and *P. falciparum*, respectively. Blood samples were collected by venipuncture from each patient at the day of diagnosis (D0 - in the acute phase) and, after collection, all patients were treated with the regimen recommended by the Brazilian Ministry of Health (MS 2010). Patients returned 15 days later (D15 - in the convalescent stage) for follow-up examinations and paired blood samples were collected from 40 *P. vivax* and 15 *P. falciparum* infected patients. All patients were symptomatic and had clinical symptoms ranging from very mild illness to full-blown paroxysms, but there were no severe or complicated malaria cases. The patients were positive for either *P. falciparum* or *P. vivax* parasites as determined by microscopy using thick and thin blood smears at D0. Asexual blood forms of *P. falciparum* or *P. vivax* were cleared from the peripheral blood of all patients included in the study following therapy and no parasite reappearance was observed during follow-up. The control group (n = 12) was composed of apparently healthy individuals who lived in the same area, but were negative for malaria parasites as determined thick blood smear at D0. Asexual blood forms of *P. falciparum* or *P. vivax* were cleared from the peripheral blood of all patients included in the study following therapy and no parasite reappearance was observed during follow-up. The control group (n = 12) was composed of apparently healthy individuals who lived in the same area, but were negative for malaria parasites as determined thick blood smear and had not reported any malaria episodes for at least one year. Ethical approval for the study was granted by the Oswaldo Cruz Foundation Ethical Committee and by the National Ethical Committee of Brazil and informed consent was given by the patients.

**Laboratory tests** - Thick and thin blood films were stained with Giemsa and the *Plasmodium* species were identified and parasitaemia was determined by microscopic examination at D0 and D15. Parasitaemia levels were estimated by counting the number of parasites (all species and stages) per 200 white blood cells (WBC) on blood films. If fewer than nine parasites were detected, 300 additional leucocytes were counted to obtain more precise results. Complete blood cell counts, including haematologic indices, were performed at D0 and D15 using an automatic haematology analyser (Pentra ABX) and peripheral blood smears were performed for routine dif-
fetal blood cellular quantification. The cell counters provided data on WBC counts and red blood cell (RBC) counts, haemoglobin (Hb) levels, haematocrit and reticulocyte, platelet, lymphocyte, eosinophil, segmented neutrophil, band cell, monocyte and basophil counts. The smear was examined by qualified pathologists. The presence of reticulocytes was evaluated using Brilliant Cresyl blue solution. The patients were considered anaemic when their Hb levels were ≤ 13 g/dL in males and ≤ 12 g/dL of blood in females. Thrombocytopenia was defined as a platelet count < 150 × 10^9/mL.

**Multiplex microsphere cytokine immunoassay** - The levels of 16 cytokines and chemokines were detected in plasma samples using Luminex technology (Luminex Corporation, Austin, TX, USA). Thirteen cytokines [IL-1β, IL-2, IL-4, IL-5, IL-6, IL-7, IL-10, IL-12 p70, IL-17, IFN-γ, TNF-α, G-CSF, granulocyte-macrophage colony-stimulating factor (GM-CSF)] and three chemokines (IL-8, MCP-1 and MIP-1β) were analysed using a BioPlex-Kit assay (Bio-Rad Laboratories, Hercules, CA, USA). The assay was performed according to the manufacturer’s instructions using a BioPlex-kit in combination with the Luminex system. Briefly, 50 μL of standard or test sample along with 50 μL of mixed beads were added into the wells of a pre-wetted 96-well microtitre plate. After 1 h of incubation and washing, 25 μL of detection antibody mixture was added and the samples were incubated for 30ºC min and then washed. Finally, 50 μL of streptavidin-PE was added and after 10ºC min of incubation and washing, the beads were resuspended in 125 μL assay buffer and analysed using a BioPlex suspension array system (Bio-Rad Laboratories) and the Bio-Plex manager software (v.3.0). A minimum of 100 beads per region were analysed. A curve fit was applied to each standard curve according to the manufacturer’s manual and sample concentrations were interpolated from the standard curves. The limit of cytokine detection using this method was 2 pg/mL for all cytokines and chemokines. The median cytokine and chemokine levels in 12 healthy controls were 2 pg/mL for IL-1β, IL-2, IL-4, IL-5, IL-6, IL-7, IL-10, IL-12 p70, IL-10 and GM-CSF, 15.53 pg/mL for IFN-γ, 12.58 mL for TNF-α, 4.3 pg/mL for GCS-F, 495 pg/mL for MCP-1 and 594.5 pg/mL for MIP-1β.

**Statistical analysis** - Survey data were recorded and entered into a database created with Epi Info 2007 (Centers for Disease Control and Prevention, Atlanta, GA, USA). Analyses were performed using Predictive Analytics Software v.17.0 (SPSS Inc, Chicago, IL, USA) and Prism v.5 (GraphPad Software Inc, San Diego, CA, USA). Differences in median haematological parameters and cytokine levels were expressed as medians and interquartile ranges (IR) and compared using Bonferroni’s multiple comparison test. When this test indicated a significant difference (p < 0.05) among pairwise groups, a Mann-Whitney U test was used. To evaluate the significant differences in haematological and cytokine parameters between the acute and convalescent phases from the same patient, non-parametric paired t tests were used. Differences in proportions were evaluated using the chi-squared test. Finally, the correlations between parasitaemia, blood cells and cytokine levels were calculated using Spearman’s rank correlation coefficient and p < 0.05 were considered statistically significant.

**RESULTS**

**Study subjects** - Seventy-one patients infected with malaria were enrolled in the study. The majority of the patients were male. There were no differences in mean age, time of residence in the endemic area and number of past malaria episodes between *P. vivax* (n = 47) and *P. falciparum* (n = 24) infected patients. All patients presented general clinical symptoms such as history of fever and headache at the time of enrolment independent of *Plasmodium* species. The time elapsed between the appearance of the first symptoms and malaria diagnosis was similar between patients with *P. vivax* (3 days) and *P. falciparum* (3.5 days). Although the mean parasitaemia was higher in *P. falciparum*-infected individuals than *P. vivax*-infected individuals, this difference was not statistically significant. All patients were parasitaemia-negative by day 15 of follow up after receiving effective drug treatment. The characteristics of the participants are presented in Table I.

**Haematological results** - To investigate haematological changes during malaria infection, differential haematological parameters during the acute and convalescent phases, expressed as the median (IR), are shown in Table II. The median lymphocyte and platelet counts in *P. falciparum* and *P. vivax* patients during acute disease were lower than in the control subjects and returned to control reference levels during the convalescent stage. In contrast, the median band cell counts were elevated in both *P. vivax* and *P. falciparum*-infected patients during the acute phase (p = 0.0041 and p = 0.0001, respectively) and returned to normal levels during the convalescent stage. We also found that patients with acute *P. vivax* infection had low eosinophil counts (p = 0.013, control values 153/µL) that increased during the convalescent stage. All other haematological values were similar among *P. falciparum*-infected, *P. vivax*-infected and control subjects. Although no differences in Hb values were observed between *P. vivax* and *P. falciparum*-infected patients during the acute phase (p = 0.0041 and p = 0.0001, respectively) and returned to normal levels during the convalescent stage, we also found that patients with acute *P. vivax* infection had low eosinophil counts (p = 0.013, control values 153/µL) that increased during the convalescent stage. All other haematological values were similar among *P. falciparum*-infected, *P. vivax*-infected and control subjects. Although no differences in Hb values were observed between *P. vivax* and *P. falciparum*-infected patients during the acute phase, anaemia was detected in 29.2% of *P. falciparum*-infected, 30% of *P. vivax*-infected and 25% of control subjects. However, there were no differences in the percentage of anaemia between the control subjects and patients with malaria infection, with 38.9% and 40% previously infected by *P. vivax* and 40% previously infected by *P. falciparum* were anaemic.

**Circulating cytokine and chemokine levels during the acute and convalescent phases of a malaria episode** - The data in Fig. 1 compare circulating cytokine and chemokine levels in patients infected with *P. vivax* and *P. falciparum*. First, the cytokines IL-5, IL-7 and GM-CSF were not detectable in most plasma samples and no differences were observed in MCP-1 levels compared with controls. During the acute phase, *P. vivax* and *P. falciparum* patients had significantly higher IL-6, IL-8, IL-17, IFN-γ, TNF-α, MIP-1β and G-CSF plasma...
concentrations than controls. To investigate changes in cytokine levels during infection, we compared cytokine levels in the sera from the same patient during the acute and convalescent phases and plasma levels of IL-6, IL-8, IL-17, IFN-γ, TNF-α, MIP-1β and G-CSF were higher during the convalescent phase. Although P. falciparum and P. vivax malaria patients have similar cytokine profiles during infection, P. falciparum patients presented higher levels of IL-6, IL-8, IL-17, IFN-γ, TNF-α, MIP-1β and G-CSF than P. vivax patients during the convalescent phase. In contrast, only P. vivax patients presented higher levels of TNF-α during the convalescent phase than during the acute phase of infection. IL-10 levels were detected at high concentrations in the majority of P. falciparum and P. vivax malaria patients have similar cytokine profiles during infection, P. falciparum patients presented higher levels of IL-6, IL-8, IL-17, IFN-γ, MIP-1β and G-CSF than P. vivax patients during the convalescent phase. In contrast, only P. vivax patients presented higher levels of TNF-α during the convalescent phase than during the acute phase of infection. IL-10 levels were detected at high concentrations in the majority of P. falciparum and P. vivax malaria patients during the acute phase and returned to completely normal levels during the convalescent phase. The median IL-10 concentration during the acute phase was 1,175 pg/mL (IR = 155-3,135) at D0 and 2 pg/mL (IR = 2-2) at D15 and 1,187 pg/mL (IR = 502.5-3049) at D0 and 2 pg/mL (IR = 2-149.8) at D15 in P. vivax and P. falciparum-infected patients, respectively.

The levels of IL-1β, IL-4 and IL-12 were similar to those of the controls during the acute phase for both P. falciparum and P. vivax patients and were higher during the convalescent phase, with the exception of IL-12, which was higher only in the P. falciparum group. The plasma IL-2 concentration was determined only during the acute phase in P. falciparum patients. Although the median IL-2 levels were higher during the convalescent phase, these differences were not statistically significant.

Relationship between parasitaemia and other variables - Spearman’s correlation coefficient values and p values are shown in Fig. 2. Various combinations of variables were analysed for possible correlational relationships. In P. vivax patients, we found a statistically significant negative correlation between platelet count and parasitaemia during the acute phase (Fig. 2A). The acute phase plasma IL-10 concentration was positively correlated with parasitaemia in P. vivax (Fig. 2C) and P. falciparum-infected patients (Fig. 2D). The same was true for the TNF-α concentration in P. falciparum-infected patients (Fig. 2B). No relationship was found between parasitaemia and the other blood cell counts or the concentration of others cytokines assayed (data not shown).

DISCUSSION

Haematological changes, such as alterations in total and differential WBC counts, are widely used to differentiate between several types of infections and to monitor the course of diseases (Ventura et al. 1999, Lathia & Joshi 2004, Tangpukdee et al. 2008). Malaria-induced changes in the differential white cell counts are very diverse and
contradictory and include leucopenia, lymphopaenia, lymphocytosis, the presence of atypical lymphocytes, monocytosis, neutropenia, neutrophilia, immature neutrophils (band cells), eosinopaenia, eosinophilia and monocytosis, neutropaenia, neutrophilia, immature mononuclear cells (Hviid et al. 1997, Balde et al. 2000). The findings of our study show that increased band cells are common during acute P. falciparum and P. vivax malaria. The decreased lymphocyte levels during malaria infection have been attributed to the reallocation of cells to deep lymphoid organs or by parasite induced apoptosis of human mononuclear cells (Hviid et al. 1997, Balde et al. 2000). Both phenomena most likely contributed to the lymphopenic state in our patients. Fifteen days following treatment, when no parasites were detected in either P. falciparum and P. vivax patients, the lymphocyte counts were similar to those in the control subjects, indicating that this period of time was sufficient for the patients to achieve lymphocyte homeostasis. With respect to eosinopaenia during the acute phase, it has been suggested that malaria either suppresses eosinophil production and release from the bone marrow or enhances the peripheral removal of these cells (Davis et al. 1991, Aubouy et al. 2002, Tangpuksdee et al. 2008). The increased eosinophil counts we observed post-treatment have been observed in previous studies (Kurtzhal et al. 1998, Tangpuksdee et al. 2008). In those studies, the induction of eosinophils was attributed to various factors such as higher release of eosinophils after temporary bone marrow suppression caused by Plasmodium, a direct response to the parasite or a response to antimalarial drugs.

The transitory increase in band cells that was observed in both infections indicates a stronger stimulus for neutrophil production during the acute phase. In this case, early or premature release of neutrophils from the bone marrow occurs, resulting in an increased proportion of younger, less well-differentiated neutrophils into the circulation. Though this alteration is common knowledge in other acute diseases, very few studies evaluating these disturbances have been conducted for this cell type in malaria patients (Hanscheid et al. 2008, Lima-Junior et al. 2011).

Erythrogram abnormalities are also very common in malaria patients and the most prominent alterations are anaemia and thrombocytopaenia (Collins et al. 2003, Ghosh 2007, Araujo et al. 2008, Fernandes et al. 2008, Leowattana et al. 2010, Lacerda et al. 2011). In our study, the absence of marked anaemia in malaria patients may be due to the early diagnosis and prompt treatment, free of charge, provided by the malaria control program in Brazil. However, low platelet counts were observed during acute infection. Although some authors have de-
scribed more intense thrombocytopenia during acute falciparum malaria compared to vivax malaria whereas others have described the opposite, no difference in thrombocytopenia was observed between these types of malaria in our study (Ghosh 2007, Taylor et al. 2008). However, we did observe a negative correlation between platelet counts and parasitaemia during acute *P. vivax* infection only. These inconsistent relationships may reflect differences in epidemiology, the immune status of malaria patients and many others factors (Casals-Pascual et al. 2006). Several causes of thrombocytopenia have been suggested. Disseminated intravascular coagulation...
tion, immune mechanisms, dysmyelopoiesis and hypersplenism are some examples of mechanisms that could be related to platelet reduction in malaria patients (Patel et al. 2004, Wassmer et al. 2008, Bueno et al. 2011).

The role of cytokine signalling during malaria episodes is still far from being understood. The most famous inflammatory marker of severe malaria is TNF-α, which is closely associated with fever, paroxysms, anaemia, cerebral malaria and many other systemic infection symptoms (Karunaweera et al. 2003, Armah et al. 2005). In our study, the cytokine and chemokine profiles in acute *P. vivax* and *P. falciparum* patients were similar. The acute phase was characterised by the presence of pro-inflammatory cytokines (IL-1β, IL-2, IL-6, IL-17, IFN-γ, TNF-α), anti-inflammatory cytokines (IL-4 and IL-10), chemokines (IL-8, MIP-1) and G-CSF in most malaria patients. During the convalescent phase (D15), the levels of all these cytokines increased compared with D0, except for IL-10 levels, which were elevated only during the acute phase and were associated with parasite density. We and others have shown a marked IL-10 response during symptomatic uncomplicated *P. vivax* malaria infection and a significant positive correlation between plasma IL-10 levels and parasite density during *P. vivax* and *P. falciparum* infection. The discordant results regarding cytokine production during the convalescent phase in our study suggests different regulatory mechanisms for early parasite clearance. Whether the differences in serum cytokine levels noted in our study are biologically significant is also unclear. It is tempting to speculate that the phagocytic cells are committed to a more Th1-biased phenotype during drug-induced clearance of parasitaemia and release of parasite metabolites, such as haemozoin, which is a known inducer of pro-inflammatory responses via signalling through Toll-like receptors 9 (Coban et al. 2005). Nonetheless, before the initiation of antimalarial treatment, a notably high IL-10 concentration that markedly decreased with the resolution of parasitaemia was observed and this down regulation of the Th2 response discriminated the successfully treated malaria patients. IL-10 has a number of effects and IL-10 inhibited IL-6, IFN-γ and TNF-α secretion and function in an in vitro malaria model when anti-IL-10 antibody was produced. Another plausible cause for the increase in several cytokine levels between the acute and convalescent phases could be the diminished suppressive effect by IL-10 on other cytokines such as pro-inflammatory cytokines, as well as a reduced systemic inflammatory response due to all haematological alterations returning to references values. IL-10 perturbations appear to have the most significant inhibitory effect on other cytokine concentrations. In this endemic area, we were unable to rule out concomitant intestinal parasite infections with organisms such as helminths, which potentially influenced the results (Hartgers & Yazdanbaksh 2006). An association of IL-10 levels with parasitaemia has also been reported in *Plasmodium knowlesi* and *P. vivax* infections. Moreover, IL-10 levels in both *P. vivax* and *P. knowlesi* patients were elevated, but were not associated with markers of disease severity (Cox-Singh et al. 2011). In contrast, the IFNγ/IL-10 ratio has been suc-
cessfully used as a marker for pathological inflammatory activity in *P. vivax* patients with varying disease severity (Andrade et al. 2010). Several studies report that IL-10 has been implicated in malaria pathophysiology (Dodoo et al. 2002, Corrigan & Rowe 2010, Cox-Singh et al. 2011). In our study, IL-10 predominated in the early anti-inflammatory response in *P. falciparum* and *P. vivax*-infected patients and dropped drastically during the convalescent phase when malaria had been cured in all patients. In conclusion, a complex array of cytokines is released in adult patients with uncomplicated malaria infection with apparent feedback inhibition and cross-regulatory functions. IL-10 appears to be involved during the acute phase of the disease and its decrease correlates with recovery as biological and clinical malaria features disappear. Further studies are required to determine whether these elevated IL-10 levels play a beneficial role by reducing the parasite-induced inflammatory response.

Additionally, there are no reports of cytokine concentrations in humans 15 days after the beginning of treatment; therefore, the data presented here could for the first time indicate a shift from a Th1/Th2 balanced response to a more pronounced Th1-regulated immune response during the first 15 days of uncomplicated malaria treatment in Brazilian endemic areas. Differences in epidemiology, nutritional status, demographic factors and the presence of co-infections are factors that could be related to the ambiguous findings of previous studies.

ACKNOWLEDGEMENTS

To the Secretary of Health of RO and LACEN/RO, for the infrastructure support, and to all individuals who participated in this study, for their cooperation and donation of blood.

REFERENCES


infection - association malaria. malaria in children and malaria. malaria. malaria.


