



T-cell-derived cytokines, nitric oxide production by peripheral blood monocytes and seric anti-*Leishmania (Leishmania) chagasi* IgG subclass patterns following immunization against canine visceral leishmaniasis using Leishvaccine and Leishmune[®]

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ABSTRACT

It is generally accepted that distinct cytokine expression by the cellular immune response plays a critical role during the outcome of experimental as well as natural canine visceral Leishmaniasis (CVL). Despite the fact that immunoprophylaxis of CVL has become an important control strategy and protective immunity has been reported upon immunization with whole as well as purified *Leishmania* antigens, the cytokine profile of T-cells triggered by anti-CVL vaccines still remain to be determined. Herein, we have developed a cross-sectional analysis of German Shepherd dogs submitted to vaccination protocols with Leishvaccine ($n = 6$) and Leishmune[®] ($n = 6$). Our data identified distinct immunological profiles elicited by Leishvaccine and Leishmune[®], with the Leishvaccine triggering a mixed, IFN- γ and IL-4, cytokine pattern in addition to high levels of anti-*Leishmania* IgG1, whereas the Leishmune[®] induced an immunological pattern characterized by enhanced levels of IFN- γ , NO and anti-*Leishmania chagasi* IgG2. It was important to notice that despite the distinct immunological patterns triggered by Leishvaccine and Leishmune[®], the ability of both immunobiologicals to activate T-cell-derived IFN- γ synthesis further suggesting their immunogenic potential against CVL. These findings added support to our hypothesis that both antigenic composition (whole antigen in Leishvaccine versus purified antigen in Leishmune[®]) as well as the adjuvant nature (BGC and saponin) used for the vaccine formulation may count for the distinct activation pattern observed.

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1. Introduction

Visceral leishmaniasis (VL) affects 500,000 people worldwide with a dramatic increase in the number of reported cases during the last few years [1–3]. This disease is considered as a zoonoses or antroponoses, depending on the studied region. In Brazil, likewise in the New World and in the Mediterranean area, VL is a

canidae zoonoses and therefore the current strategy for managing the disease control is based on three major actions, including (1) the systematic treatment of human cases, (2) vector control and (3) the elimination of seropositive dogs [4]. The establishment of immunoprophylactic tools to control the canine visceral leishmaniasis (CVL) represent an important issue to upgrade the strategy for managing the leishmaniasis control [5,6]. In this context, considerable effort has been dedicated in the development of vaccines against CVL, able to modify the immune repertoire and advances have been already reported [7]. Several promising vaccine have been proposed as anti-CVL vaccines and include live/killed *Leish-*

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mania parasites (first-generation), purified *Leishmania* antigens or live recombinant bacteria expressing *Leishmania* antigens (second generation) as well as antigen-encoding DNA plasmids (third generations) [8–11].

Recently, the Leishmune[®], the pioneer second generation vaccine, composed of purified fraction named fucose mannose ligand (FML) isolated from *Leishmania donovani* promastigotes plus saponin as adjuvant, have been licensed in Brazil and become commercially available [8–11]. It is important to mention that the Leishmune[®] is the only anti-CVL vaccine registered in the world, currently in use in Brazil, with protective potential supported by the literature [8,10]. This formulation proved to be safe, protective and highly immunogenic for dogs [8–13] as well as able to block CVL transmission [14].

It is a generally accepted that a protective vaccine against *Leishmania* infection should be able to trigger a specific cellular immune response, besides a type-1 cytokine pattern, which may play a critical role in resistance to the disease onset [15]. In the murine model for *Leishmania* infection, IFN- γ and IL-4 has been already reported as important hallmarks of these two poles of immune response, as they are associated with protection and susceptibility to *Leishmania* infection, respectively [16–18]. In CVL, although the general cytokine expression during the outcome of experimental infection has been already reported [19–22], little is known about the cellular immune response triggered by immunoprotective vaccines [23], especially regarding the Leishvaccine and Leishmune[®]. Recently, we have demonstrated that Leishvaccine and Leishmune[®] elicited distinct profiles of innate and adaptive immune response [24]. While Leishvaccine promoted an early recruitment of neutrophils and eosinophils with late involvement of monocytes, Leishmune[®] induced early and persistent recruitment of neutrophils and monocytes, without the enrollment of eosinophils. Regarding the adaptive immunity, Leishvaccine sponsored a mixed profile, associated with phenotypic changes in both CD4⁺ and CD8⁺ T-lymphocytes as well as on B-cells. In contrast, Leishmune[®] was associated with phenotypic changes in T-lymphocytes, particularly in CD8⁺ T-cells. We have then hypothesized that the ability of both immunobiologicals to enroll macrophages and CD8⁺ T-cells could be considered priority vaccines with a high-quality immunogenic potential against CVL [24]. Aiming to further focus on the immune response triggered by Leishvaccine and Leishmune[®] herein, we have further characterized the cellular and humoral immune response of dogs submitted to vaccination protocols with Leishvaccine and Leishmune[®]. The main goal of the current work was to investigate, in a cross-sectional study, the cytokine profile of circulating T-cell subsets, besides the level of nitric oxide synthesis by peripheral blood monocytes as well as the anti-*Leishmania* IgG subclass patterns after Leishvaccine and Leishmune[®] vaccination.

2. Materials and methods

2.1. Animals, vaccination protocols and blood sampling

Twelve healthy German Shepherd dogs, 8 males and 4 females, age ranging from 18 to 60 months old, maintained at the kennel of Polícia Militar de Minas Gerais, Brazil during the entire experimental procedures, were included in this investigation. The dogs included in this study were selected based on their negative serological results in the enzyme-linked immunosorbent assay (ELISA, Biomanguinhos, FIOCRUZ, RJ, Brazil) used as a 'reference standard' test for the diagnosis of CVL. All animals were treated for intestinal helminth infections and immunized against parvovirus, leptospirosis, distemper, parainfluenza and hepatitis and maintained in quarantine prior their inclusion in the study. During the entire experimental procedures, the animals received drinking water and a balanced feed given *ad libitum*.

The selected dogs were divided into three groups named: Group 1: "Unvaccinated" dogs, Group 2: "Leishvaccine" immunized dogs and Group 3: "Leishmune[®]" vaccinated dogs. The group "Unvaccinated" consisted of 8 animals, 4 males and 4 females (Dog#1, Dog#2, Dog#3, Dog#4, Dog#5, Dog#6, Dog#7 and Dog#8). The group "Leishvaccine" was consisted of 6 animals, 4 males and 2 females, 4 of them also evaluated before vaccination (Dog#1, Dog#2, Dog#3, Dog#4) and 2 evaluated only after vaccination (Dog#9 and Dog#10). The group "Leishmune[®]" was consisted of 6 animals, 4 males and 2 females, 4 of them also evaluated before vaccination (Dog#5, Dog#6, Dog#7, Dog#8) and 2 evaluated only after vaccination (Dog#11 and Dog#12).

Dogs in the Leishvaccine group were immunized throughout a complete vaccination regimen that included three subcutaneous doses of the vaccine with an interval of 21 days between each. Leishvaccine was prepared likewise described by Mayrink et al. [25], but consisted of *Leishmania (Leishmania) amazonensis* (strain IFLA/BR/1967/PH8) antigenic preparation using non-live lyophilized Bacille Calmette-Guérin, BCG (Fundação Ataulfo de Paiva, RJ, Brazil) as adjuvant. The first dose corresponded to 0.6 ml of Leishvaccine (360 μ g of protein) plus 0.4 ml of physiologic saline (NaCl 0.9% in distilled water) containing 400 μ g of BCG dry-weight as adjuvant. The second dose corresponded to 0.6 ml of Leishvaccine (360 μ g of protein) plus 0.3 ml of physiologic saline (NaCl 0.9% in distilled water) containing 300 μ g of BCG dry-weight as adjuvant. The third dose corresponded to 0.6 ml of Leishvaccine (360 μ g of protein) plus 0.2 ml of physiologic saline (NaCl 0.9% in distilled water) containing 200 μ g of BCG dry-weight. Dogs in the Leishmune[®] group were submitted to a complete vaccination regimen as recommended by the manufacturer (FortDodge[®], Campinas, SP, Brazil), which included three subcutaneous doses of 1.0 ml of vaccine with an interval of 21 days between each dose. Leishmune[®] is composed of 1.5 mg lyophilized FML antigen plus 0.5 mg of Quillaja saponaria saponins (QS21 and deacylated) of Riedel de Haen reconstituted in 1 ml NaCl 0.9% sterile saline solution and administered subcutaneously. The FML-vaccine, Leishmune[®], is registered as a Patent: INPI number: PI1100173-9 (18.3.97), Federal University of Rio de Janeiro, Brazil.

Peripheral blood samples were collected from the radial vein into two vacutainer tubes, one containing sodium heparin and another one without any anticoagulant (BD Pharmingen, San Diego, CA, USA). The whole blood and sera samples were collected from unvaccinated dogs and also 40 days after the last immunization dose of dogs submitted to vaccination protocols with Leishvaccine and Leishmune[®]. Whole blood samples were maintained at room temperature until processing. The sera samples were stored at -20°C and thawed immediately before the flow immunofluorometric analysis of anti-*Leishmania chagasi* antibodies.

All procedures in this study were according to the guidelines set by the Brazilian Animal Experimental College (COBEA). This study was approved by the Ethical Committee for the use of Experimental Animals of the Universidade Federal de Minas Gerais, Brazil (CETEA).

2.2. Preparation of soluble *L. chagasi* antigen and fixed *L. chagasi* promastigote forms

L. chagasi promastigote forms (MHOM/BR/1972/BH46) were grown in liver infusion tryptose medium (LIT), supplemented with 10% of fetal bovine serum (FBS) [26] at 24°C temperature. Stationary-phase parasites (8 days of growth) were transferred to 50 ml polypropylene tubes (Falcon, Becton Dickinson, San Diego) and submitted to differential centrifugation ($100 \times g$, 10 min, room temperature) to remove remaining clusters of parasites contaminants in the pellet. Prior to recover the single-cell parasite

suspension, the supernatant was left to rest for 10 min at room temperature. The supernatant consisting of single-cell parasite suspension was transferred to another 50 ml polypropylene tube and spin down at high speed ($1000 \times g$) for 10 min at $4-8^\circ\text{C}$. The supernatant was discarded and the pellet washed twice ($1000 \times g$, 10 min, $4-8^\circ\text{C}$) with phosphate buffered saline, PBS (0.15 M, pH 7.2). After the wash steps, the single-cell *L. chagasi* promastigote suspension was then used in the preparation of soluble *L. chagasi* antigen or fixed suspension of *L. chagasi* promastigote.

The soluble *L. chagasi* antigen was prepared from the frozen (-70°C) dry pellet obtained from the *L. chagasi* single-cell promastigote suspension. The frozen pellet was thawed, resuspended into equal volume of cold PBS and submitted to three ultra-sound cycles of 1 min at 40 W on ice bath (Sonifier Cell Disruptor® - Branson Sonic Power Co., EUA). The sonicated material was centrifuged at $50,000 \times g$ for 1 h and 30 min at 4°C . The supernatant was transferred to dialysis tubes and dialyzed through PBS for 24 h, and submitted to three PBS changes every 6 h. The dialyzed soluble antigen was filtered in $0.22 \mu\text{m}$ disposable syringe sterile filters of under aseptic conditions. One aliquot was taken for protein quantification by the method described by Lowry et al. [27]. Final protein concentration was adjusted to $1000 \mu\text{g/ml}$. Antigen preparation was stored in $100 \mu\text{l}$ aliquots at -70°C prior the use in the *in vitro* cultures of peripheral blood mononuclear cells.

The suspension of fixed *L. chagasi* promastigotes were prepared by resuspending the dry pellet obtained from the *L. chagasi* single-cell promastigote suspension in five times higher volume of fix solution consisting of equal volume of cold PBS plus FACS FIX solution (10 g/l paraformaldehyde, 10.2 g/l sodium cacodylate, 6.63 g/l sodium chloride, pH 7.2, all from Sigma, St. Louis, MO, USA). Following overnight incubation at 4°C , the fixed promastigotes were washed once in PBS, counted in Neubauer chamber and stored at 4°C . The suspension adjusted to 1.0×10^7 promastigotes/ml in PBS 3% FBS prior use in the flow cytometric analysis of anti-fixed *L. chagasi* promastigotes antibodies by flow cytometry—FC-AFPA.

2.3. Isolation and *in vitro* culture of peripheral blood mononuclear cells

Peripheral blood mononuclear cells (PBMC) were isolated from heparinized blood samples previously diluted in equal volume of RPMI 1640 (Gibco, Grand Island, NY, USA) that had been layered onto 15 ml of Ficoll-Hypaque density gradient (Histopaque® 1.077; Sigma Chemical Co.) and centrifuged at $700 \times g$ for 40 min at room temperature. The PBMC cushion was collected in the interface between the top plasma layer and the Ficoll-Hypaque column. Cells were then washed twice with RPMI 1640 ($600 \times g$, for 10 min, at room temperature), counted and resuspended in RPMI 1640 at 1×10^7 cells/ml.

The culture assays were performed in 24-well flat-bottomed tissue culture plates (Corning, New York, NY, USA), each well containing $800 \mu\text{l}$ of cell culture medium comprising of RPMI 1640 supplemented with streptomycin (100 mg/ml), penicillin (100 U/ml), L-glutamine (2 mM), β -mercaptoethanol (5×10^{-5} M) and 10% FBS. Aliquots of $100 \mu\text{l}$ of the PBMC suspension (1×10^6 cells/well) were added to quadruplicate wells following the addition of $100 \mu\text{l}$ of RPMI 1640 (control cultures, CC) or $100 \mu\text{l}$ of soluble *L. chagasi* antigen (SLA) at final concentration of $25 \mu\text{g/ml}$ (stimulated cultures, SLA). Cultures were submitted to incubation in 5% CO_2 humidified incubator, at 37°C , for 5 days. Brefeldin A, BFA (Sigma, St. Louis, MO, USA) was added to each well at final concentration of $10 \mu\text{g/ml}$ and cultures submitted to an additional period of 4 h of incubation in 5% CO_2 humidified incubator, at 37°C .

Short-term whole blood cultures were performed to evaluate the assay and sample viability (positive control cultures, PMA -Phorbol 12-Myristate 13-Acetate). For this purpose, a $500 \mu\text{l}$

aliquot of whole blood was incubated in the presence of $500 \mu\text{l}$ of RPMI-1640 plus PMA (Sigma, St. Louis, MO, USA) at a final concentration of 25 ng/ml, ionomycin (Sigma, St. Louis, MO, USA) at $1 \mu\text{g/ml}$ and BFA at final concentration of $10 \mu\text{g/ml}$. The positive control culture was maintained for 4 h in 5% CO_2 humidified incubator at 37°C . At the end of incubation periods, all cultures were treated with EDTA diluted in PBS (Sigma, St. Louis, MO, USA) at a final concentration of 2 mM for 15 min, at room temperature.

2.4. Immunostaining for cell surface markers and intracellular cytokines

The EDTA-treated cell cultures were washed once with FACS buffer prepared as PBS supplemented with 0.5% of bovine serum albumin-BSA (Sigma, St. Louis, MO, USA) by centrifugation at $600 \times g$ for 7 min at room temperature. Cell pellet was resuspended with $400 \mu\text{l}$ of FACS buffer and $200 \mu\text{l}$ aliquots incubated in 5 ml polystyrene tubes (Becton Dickinson, Franking Lakes, NJ, USA) in the presence of $60 \mu\text{l}$ of previously diluted anti-canine CD4-1:320 (rat IgG2a, clone YKIX302.9) or anti-canine CD8-1:40 (rat IgG1, clone YCATE55.9) monoclonal antibodies (mAb), both labeled with fluorescein isothiocyanate dye (FITC) and purchased from SEROTEC (Oxford, UK). Following incubation at room temperature, for 30 min, in the dark, the membrane-stained samples were treated with 3 ml of FACS Lysing/fix Solution (BD Biosciences, San Jose, CA, USA), immediately vortexed and re-incubated for 10 min at room temperature. After one wash procedure with FACS buffer, the membrane-stained lymphocytes were then permeabilized by incubation for 15 min with 3 ml of FACS perm-buffer (FACS buffer supplemented with 0.5% of saponin). After one wash procedure with FACS buffer, cells were incubated in the dark, for 30 min at room temperature, in the presence of $50 \mu\text{l}$ of PE-labeled anti-bovine cytokines mAb that cross-react with canine cytokines (as reported by the manufacturer), including anti-IFN- γ (clone CC302) and anti-IL-4 (clone CC303), both purchased from SEROTEC (Oxford, UK). After intracellular staining, the cells were washed once with FACS perm-buffer, followed by one wash step with FACS buffer and finally fixed in FACS FIX Solution. FITC and PE-labeled isotypic controls were included in each batch of experiments.

Flow cytometric measurements were performed on a FACScan instrument (Becton Dickinson, Mountain View, CA) interfaced to an apple G3 FACStation. The Cell-Quest™ software package provided by the manufacturer (Franklin Lakes, NJ, USA) was used for data acquisition and analysis. A total of 30,000 events were acquired for each preparation.

The analysis of the cytokine profile of CD4^+ and CD8^+ T-cell subsets was performed by first establishing a scattering gate on the lymphocyte population, using laser forward scatter (FSC) versus laser side scatter (SSC) dot plot distributions, followed by quantification of cytokine expressing cells on FL1/FITC versus FL2/PE dot plots combinations. The cytokine⁺ T-cell subsets were identified into the upper-right quadrant on dual color graphs with the FL1/FITC axis representing immunostaining with the anti-cell surface marker FITC-labeled mAb (CD4 or CD8) and FL2/PE corresponding to the immunostaining with the anti-cytokine PE-labeled mAb (IFN- γ or IL-4). The results were expressed as the percentage of double labeled cells (IFN- $\gamma^+\text{CD4}^+$, IL-4⁺ CD4^+ , IFN- $\gamma^+\text{CD8}^+$ and IL-4⁺ CD8^+) within the lymphocyte logical gate.

2.5. Analysis of nitric oxide (NO) synthesis

The concentration of nitrite (NO_2^-) released in the supernatant of *in vitro* PBMC cultures was measured using the Griess reaction [28]. Briefly, a $100 \mu\text{l}$ aliquot of cell-free culture supernatant was mixed with $100 \mu\text{l}$ of Griess reagent (1% sulphonylamide, 0.1% naphthylethylene-diamide-dihydrochloride and 2.5% phos-

phoric acid, all from Sigma, St. Louis, MO, USA). Following 10 min of incubation at room temperature, in the dark, the absorbance was measured at 540 nm, using a microplate reader. Each sample was assayed in duplicate and the concentration of nitrite was determined by interpolation from a standard curve constructed using sodium nitrite solutions of known concentration in the range 0–100 μ M. To discount the interference of nitrites already present in the culture medium, data was calculated taking into account the blank for each experiment, assayed by using the medium employed for the *in vitro* PBMC cultures. The results were first expressed as nitrite concentration (μ M). The nitrite level was then divided by the number of monocytes added to each *in vitro* PBMC culture in order to yield the nitric oxide index (nitrite/monocytes).

2.6. Detection of anti-fixed *L. chagasi* promastigotes antibodies by flow cytometry–FC-AFPA

The FC-AFPA procedure was carried out as proposed by de Andrade et al. [29] for the detection of anti-fixed promastigote *L. chagasi* immunoglobulins. Briefly, in 96-well U-bottom plates (LINBRO, ICN Biomedicals, Inc. Aurora, OH), 50 μ l aliquots of the pre-fixed parasite suspension (5.0×10^5 parasites/well) were incubated at 37 °C for 30 min in the presence of 50 μ l of pre-diluted serum samples in PBS-3% FBS (1:256–1:8,192 for IgG; 1:256–1:4096 for IgG1 and 1:256–1:16,384 for IgG2). Following the incubation, parasites were washed twice with 150 μ l of PBS-3% FBS (1000 \times g, 10 min, 4 °C). The parasites were re-incubated in the dark, for 30 min at 37 °C in the presence of 50 μ l of FITC-labeled sheep (putatively anti-IgG and anti-IgG2) and goat (putatively anti-IgG1) polyclonal antibodies anti-canine second step reagents previously diluted in PBS-3% FBS (anti-IgG 1:1000-cat. A40-105F, anti-IgG1 1:500-cat. A40-120F and anti-IgG2 1:1000-cat. A40-121F), all purchased from Bethyl laboratories Inc. (Montgomery, TX, EUA). After incubation parasites were submitted to two washing procedure using 150 μ l of PBS-3% FBS (1000 \times g, 10 min, 4 °C), fixed with 200 μ l of FACS FIX solution and maintained for at least 30 min, at 4 °C in the dark, prior the flow cytometric data acquisition. An internal control of the reaction, in which the parasites were incubated in the absence of dog serum, but in the presence of the FITC-labeled secondary reagents, was included in all set of experiments to monitor unspecific bindings. In all batches of FC-AFPA, positive and negative controls were also run to certify the test performance. Flow cytometric acquisition was performed using a FACScalibur® flow cytometer (BD Pharmingen) considering a total of 5000 events per tube. CELLQuest® software package (Franklin Lakes, NJ, USA) was used for data acquisition and analysis. Data analysis was performed by first gating the promastigote forms based on their size and granularity properties on FSC \times SSC dot plots. The relative FITC fluorescence intensity of the selected promastigotes was then quantified on single color fluorescen-1 (FITC/FL1) histograms. The histogram distributions, obtained for the internal control of unspecific binding of the second step reagents, were used to a set up a marker to confine at least 98% of promastigotes into a region of negative fluorescence intensity. Once established, the marker was used to determine the percentage of positive fluorescent parasites (PPFP) for each tested sample.

2.7. Statistical analysis

Statistical analysis was performed using the GraphPad Prism 4.03 software package (San Diego, CA, USA). Data analysis was carried out by ANOVA followed by Tukey's multiple comparison test to evaluate the cytokine and NO profiles between unvaccinated and vaccinated dogs 40 days after the end of the vaccination protocols. Analyses of immunoglobulin reactivity as well as the association between immunoglobulin and cytokine profiles were carried out

by Spearman correlation test. In all cases, the differences were considered significant when the probabilities of equality, *p*-values were <0.05.

3. Results

3.1. Regardless of the distinct cytokine patterns observed in CD4⁺ and CD8⁺ T-cell subsets, both Leishvaccine and Leishmune® vaccines elicited high levels of T-cell-derived IFN- γ

Aiming to further focus on the impact of Leishvaccine and Leishmune® vaccines on the T-cell cytokine pattern, herein we have characterized for the first time, the frequency of both IFN- γ ⁺ and IL-4⁺ T-cells as well as their major subsets (CD4⁺ and CD8⁺) within PBMC collected from unvaccinated and Leishvaccine and Leishmune® vaccinated dogs. For this purpose, PBMC were isolated and submitted to *in vitro* cultures in the absence (control, CC) or presence of soluble *L. chagasi* antigens (SLA). The frequency of both IFN- γ ⁺ and IL-4⁺ T-cell subsets observed in the CC and SLA cultures are presented in Fig. 1. Our data demonstrated that both Leishvaccine and Leishmune® induced increased levels of IFN- γ ⁺ T-cells as compared with unvaccinated dogs, in both CC and SLA cultures (Fig. 1A). Further analysis highlighted that this phenotype was selectively observed in CD4⁺ T-cells from Leishvaccine and Leishmune® vaccinated dogs, in CC and SLA cultures (Fig. 1B). No changes were observed in the IFN- γ synthesis by CD8⁺ T-cells (Fig. 1C). It was interesting to notice that Leishmune® showed no significant levels of IL-4, which corroborate the previous finding about protection in the field studies.

On the other hand, higher levels of IL-4⁺ T-cells were observed selectively in the SLA cultures of PBMC from dogs immunized with Leishvaccine as compared to both unvaccinated and Leishmune® vaccinated dogs (Fig. 1D). Further analysis demonstrated that this phenotype were mainly due to IL-4⁺CD8⁺ T-cells (Fig. 1F). Increased percentages of IL-4⁺CD4⁺ T-cells was also observed in Leishvaccine group as compared to unvaccinated dogs, however, selectively in the CC culture (Fig. 1E).

Additional analyses showed significant increase in the IFN- γ ⁺/IL-4⁺ T-cells ratio in CC cultures of PBMC from both Leishvaccine and Leishmune® vaccinated dogs as compared to unvaccinated dogs. However, increased IFN- γ ⁺/IL-4⁺ T-cells ratio was selectively observed in Leishmune® vaccinated dogs when analyzing the SLA cultures. Interestingly, the IFN- γ ⁺/IL-4⁺ T-cells ratio observed in the SLA cultures for Leishmune® vaccinated dogs was higher than that observed in the CC culture of PBMC obtained from these animals (Fig. 1G).

The cytokine patterns of T-cell subsets from Leishvaccine and Leishmune® vaccinated dogs are illustrated by flow cytometry dot plots representatives of CC (Fig. 1H) and SLA cultures (Fig. 1I). The analysis of cytokine⁺ T-cell subsets in the PMA-induced cultures confirmed the cell viability of all samples, as demonstrated by high levels of IFN- γ ⁺ and IL-4⁺ cells within both, CD4⁺ and CD8⁺ T-cell subsets (Fig. 1J).

3.2. Despite triggering distinct cytokine profiles, both Leishvaccine and Leishmune® were able to shift the overall cytokine balance toward a type-1 immune response

Taking the general hypothesis that a fine balance between IFN- γ and IL-4 profile is more relevant that a shift toward a polarized cytokine pattern, we have further characterize for each animal within the Leishvaccine and Leishmune® vaccinated dogs, the overall balance of IFN- γ and IL-4 derived from CD4⁺ and CD8⁺ T-cell subsets. This strategy allows the characterization of the resultant cytokine profile from T-cells driven by the vaccination interven-

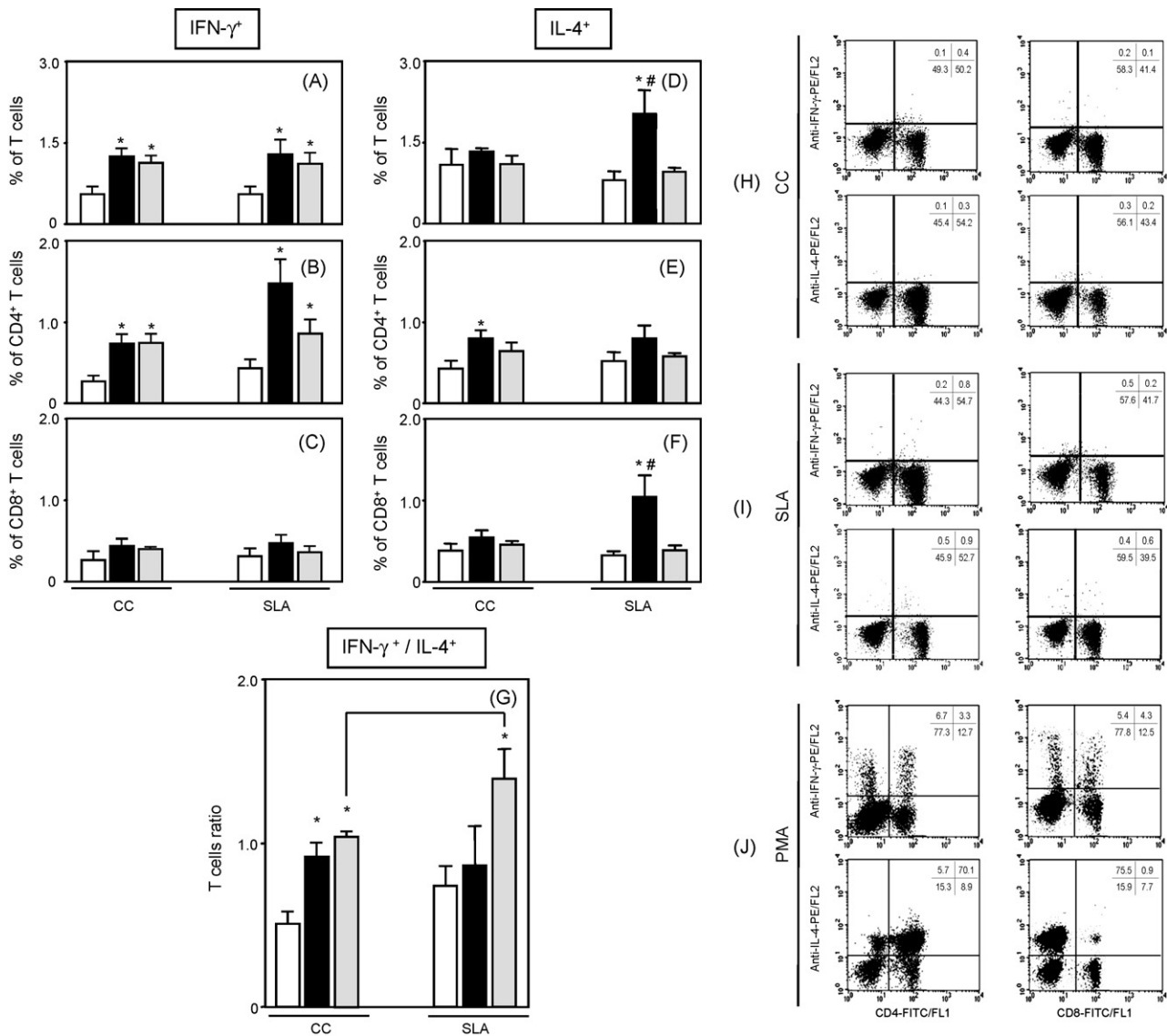


Fig. 1. IFN- γ and IL-4 profile of T-cell subsets from German Shepherd dogs following Leishvaccine (■) and Leishmune® (□) vaccination regimens as compared to unvaccinated controls (□). Peripheral blood mononuclear cells (PBMC) were cultured *in vitro* in the absence (CC) as well as in the presence of soluble *L. chagasi* antigen (SLA). IFN- γ + (A–C) and IL-4+ T-cell subsets (D–F) as well as IFN- γ + / IL-4+ T-cell ratio (G) were analyzed after flow cytometric immunostaining for cell surface markers and intracellular cytokines. The results are expressed as mean percentage of cytokine+ cells \pm standard error. Significant differences at $p < 0.05$ are indicated by * for comparisons between Leishvaccine and Leishmune® vaccinated dogs and unvaccinated dogs and by # for comparisons between CC and SLA cultures. Representative dot plots illustrating the intracellular cytokine profile of T-cell subsets in the CC, SLA and PMA-stimulated positive control cultures are provided (H–J, respectively).

tion. To establish the overall cytokine balance data obtained by flow cytometry the results were further transformed as proposed by Vitelli-Avelar et al. [30]. This new strategy consisted of a four step platform that includes: 1st, the establishment of “low” and “high” cytokine-producers based on global median of cytokine+ T-cell subsets calculated from the whole range of values obtained for the whole study population (all dogs) including unvaccinated dogs, Leishvaccine and Leishmune® vaccinated dogs (Fig. 2); 2nd, the construction of color diagrams for each group of animals (Unvaccinated, Leishvaccine and Leishmune® vaccinated dogs) showing the “cytokine pattern” of “low” and “high” cytokine-producers within CD4+ and CD8+ T-cell subsets (Fig. 3A); 3rd, the compilation of the “cytokine balance” defined as predominant low cytokine-producers, inflammatory, regulatory or mixed cytokine-producers within T-cells (Fig. 3B) and 4th, the assemble of the “overall cytokine balance” as the proportion of high inflammatory, regulatory or mixed cytokine producers within T-cells (Fig. 3C).

The analysis of the “cytokine pattern” of T-cell subsets, based on the three major classes of cytokine-producers named as “low” cytokine-producers, “high” IFN- γ -producers and “high” IL-4-producers, observed in color diagram, demonstrated that among unvaccinated dogs there is a predominance of “low” cytokine-producers within all T-cell subsets as compared to Leishvaccine and Leishmune® vaccinated dogs, with four animals (Dog#2, 3, 5 and 8) showing a general profile of low cytokine-producers (Figs. 2 and 3A). On the other hand, the Leishvaccine and Leishmune® vaccinated dogs showed predominance of high cytokine-producers, with only one animal from the Leishmune group (Dog#7) displaying a general profile of low cytokine-producer (Fig. 2A).

Taking the “cytokine balance” within CD4+ and CD8+ T-cell subsets, our data re-emphasized the predominant pattern of low-cytokine producers within the unvaccinated dogs and pointed out the existence of a distinct cytokine balance between Leishvac-

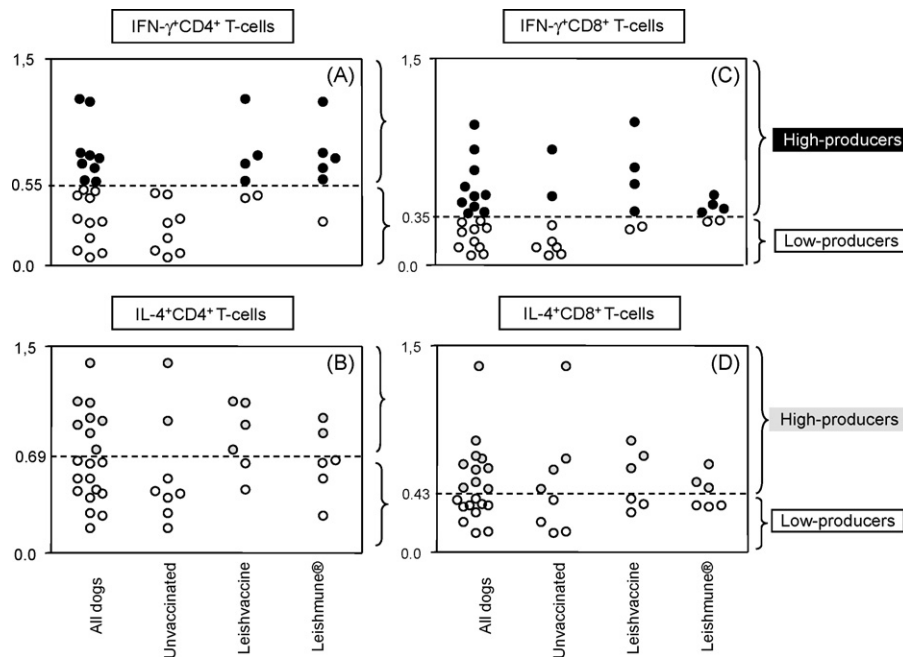


Fig. 2. Representative scatter graphs employed to identify “low” cytokine-producers (○), “high” IFN- γ -producers (●) and “high” IL-4-producers (⊙) amongst peripheral blood lymphocytes from unvaccinated and Leishvaccine and Leishmune[®] German Shepherd dogs following *in vitro* stimulation with soluble *L. chagasi* antigen. “Low” and “high” cytokine-producers were defined for each lymphocyte subset based on global median cut-off edge (---) obtained for the whole study population (all dogs). Distinct cut-offs were employed for IFN- γ +CD4⁺ (A), IL-4+CD4⁺ T-cells (B), IFN- γ +CD8⁺ (C) and IL-4+CD8⁺ T-cells (D).

cine and Leishmune[®] vaccinated dogs with distinct predominance of animals displaying inflammatory or mixed cytokine profiling (Fig. 3B).

The analysis of the “overall cytokine balance” demonstrated that 50% of the unvaccinated group is confined within low cytokine-producers (Fig. 3C, bar chart) and that amongst the high cytokine-producers, most animals (75%) presented a predominance of IL-4 cytokine profile (Fig. 3C, pie chart). On the other hand, Leishvaccine induced the pattern of high cytokine-producers in all vaccinated dogs (Fig. 3C, bar chart). In fact, the Leishvaccine drove a predominant IFN- γ profile (50%) or a mixed IFN- γ \approx IL-4 pattern (16.7%) in 66.7% of the vaccinated dogs, suggesting a protective pattern (Fig. 3C, pie chart). The Leishmune[®] was capable to shift the overall cytokine profile toward high cytokine-producers in most (87.5%) vaccinated dogs (Fig. 3, bar chart). Interestingly, the analysis of Leishmune[®] vaccinated dogs within the high cytokine-producers demonstrated a shift toward a predominant IFN- γ profile (83.3%) or a mixed IFN- γ \approx IL-4 pattern (16.7%) representing a protective pattern (Fig. 3C, pie chart).

3.3. Leishmune[®] vaccination induced high nitric oxide index (nitrite/monocytes) following *in vitro* PBMC cultures even in the presence of soluble *L. chagasi* antigens

The induction of nitric oxide (NO) is one of the major effector mechanisms leading the *Leishmania* elimination by activated phagocytes [23,31,32]. Previous report has pointed out that canine macrophages from killed *Leishmania infantum* vaccinated dogs are able to perform *Leishmania* killing throughout NO-dependent mechanisms [23]. In attempt to characterize the ability of Leishvaccine and Leishmune[®] to induce the NO production by peripheral blood monocytes, we have investigated the levels of nitrite in the supernatants of *in vitro* PBMC cultures following Leishvaccine and Leishmune[®] vaccination. Despite no differences in the levels of nitrite observed in both CC and SLA cultures (data not shown), the analyses of nitric oxide index (nitrite/monocytes), taking in account the number of monocytes added into each *in vitro* PBMC culture,

pointed out that Leishmune[®] vaccines displayed in the CC cultures, higher mean nitric oxide index as compared to unvaccinated and Leishvaccine immunized dogs. Interestingly, monocytes from Leishmune[®] vaccines also presented higher ability to produce NO as compared to unvaccinated dogs, even in the presence of soluble *L. chagasi* antigens (Fig. 4).

3.4. Distinct pattern of anti-*L. chagasi* IgG subclasses were observed in Leishvaccine and Leishmune[®] vaccinated dogs

Aiming to further characterize the immune response triggered by Leishvaccine and Leishmune[®] immunization, we have accessed the profile of seric anti-*L. chagasi* IgG in a broader range of vaccinated dogs, including 24 dogs, categorized into two subgroups referred as Leishvaccine and Leishmune[®]. Data analysis demonstrated that all vaccinated dogs seroconvert after immunization as confirmed by seropositivity in indirect immunofluorescence assay test (IFAT). Data analysis did not demonstrate any significant differences in the mean IFAT titers between Leishvaccine and Leishmune[®] vaccinated dogs (Fig. 5A). Additional analysis of anti-*L. chagasi* IgG was performed by semi-quantitative flow cytometric detection of anti-fixed *L. chagasi* promastigotes FC-AFPA-IgG, IgG1 and IgG2 [29]. Our data demonstrated that despite no changes in the FC-AFPA-IgG profiling (Fig. 5B), the median FC-AFPA-IgG1 and IgG2 PFP values suggested distinct pattern of sero-reactivity between Leishvaccine and Leishmune[®] vaccinated dogs (Fig. 5C and D). In fact, higher median FC-AFPA-IgG1 reactivity was observed in Leishvaccine group (Fig. 5C), whereas higher median FC-AFPA-IgG2 reactivity was detected in Leishmune[®] vaccinated dogs (Fig. 5D). Further analyses demonstrated a positive correlation between FC-AFPA-IgG with both FC-AFPA-IgG1 and FC-AFPA-IgG2 in Leishvaccine immunized dogs (Fig. 5E and F), while in Leishmune[®] vaccinated dogs a positive correlation was observed selectively between FC-AFPA-IgG and FC-AFPA-IgG2 (Fig. 5G and H).

The dynamic interplay of IL-4 in regulating the production of IgG subclasses has been recently documented in humans [33], suggest-

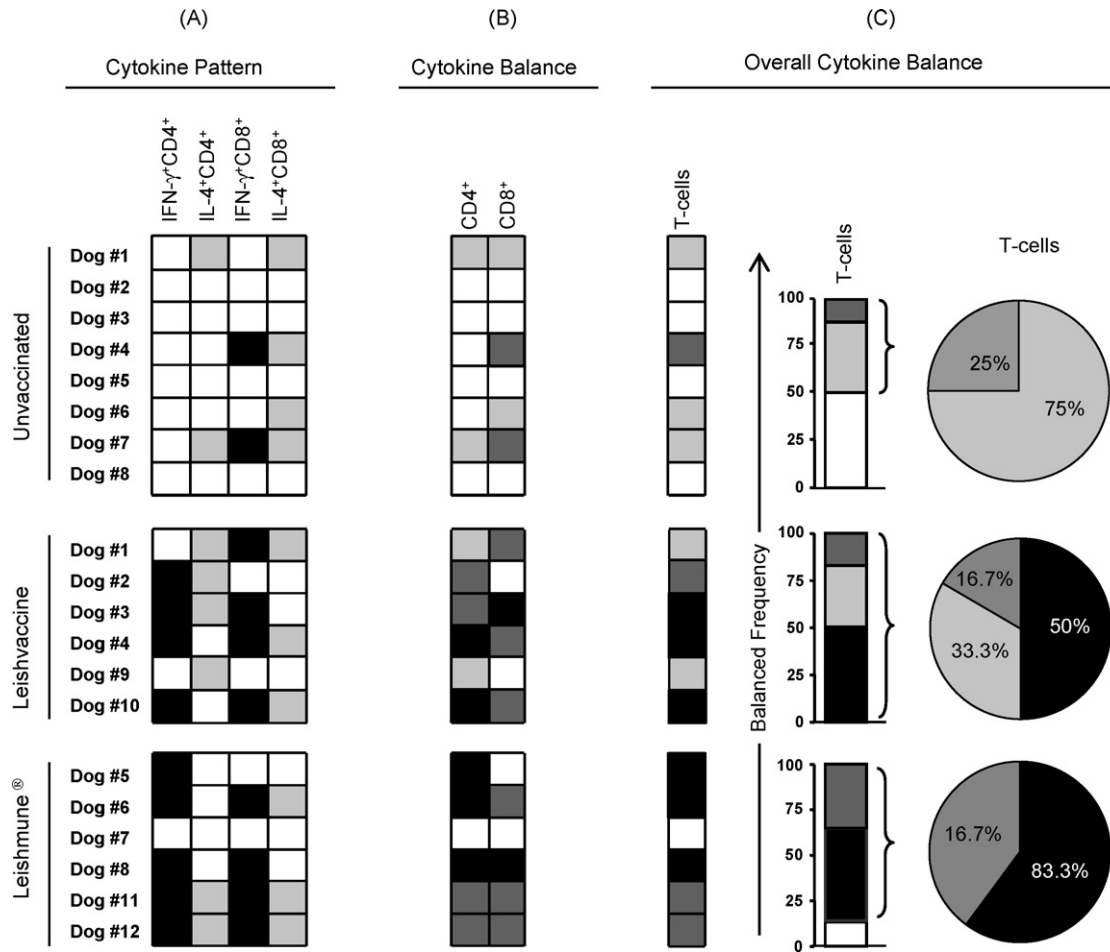


Fig. 3. Cytokine profile of peripheral blood T-lymphocytes from unvaccinated and Leishvaccine and Leishmune[®] German Shepherd dogs following *in vitro* stimulation with soluble *L. chagasi* antigen. Color diagrams were used to represent the cytokine pattern (A) and the cytokine balance within T-cell subsets (B) besides the overall cytokine balance with T-cells (C), highlighting the predominance of “low” cytokine-producers (□), “high” IFN- γ -producers (■), “high” IL-4⁺-producers (▨) or “high” mixed cytokine-producers (▩). Pie charts represent the percentage of animals displaying a given T-cells overall cytokine balance selectively amongst the “high” cytokine-producers.

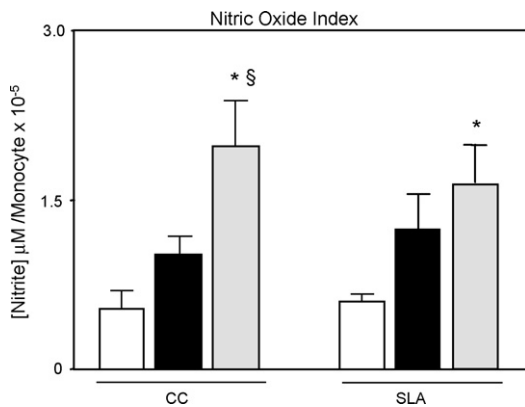


Fig. 4. Production of nitric oxide (NO) by peripheral blood monocytes from German Shepherd dogs following Leishvaccine (■) and Leishmune[®] (▨) vaccination regimens as compared to unvaccinated controls (□). As an indirect measurement of NO production, the Griess reaction was used to determine the nitrite levels in the supernatants of *in vitro* PBMC cultures performed in the absence (CC) as well as in the presence of soluble *L. chagasi* antigen (SLA). The results are expressed as nitric oxide index that represent the nitrite level (μM) divided by the number of monocytes added to each *in vitro* PBMC culture in order to yield the nitric oxide index (nitrite/monocytes). Significant differences at $p < 0.05$ are indicated by * or § for comparisons between Leishmune[®] and unvaccinated or Leishvaccine immunized dogs, respectively.

ing a temporal role for this cytokine in humoral immune responses to specific pathogens. Aiming to investigate whether the IL-4 versus IgG interplay take place during Leishvaccine and Leishmune vaccination, we have performed a correlation analysis between the IgG reactivity, detected at sera dilution 1:256 and the frequency of IL-4⁺ T-cells, focusing specifically on CD8⁺ T-cells, since they were pointed out as an important source of IL-4 in Leishvaccine immunized dogs, specially following *in vitro* SLA stimulation of PBMC cultures (Fig. 1F). Our findings demonstrated that a positive correlation could be observed between the frequency of SLA-induced IL-4⁺CD8⁺ T-cells and the reactivity of anti-*L. chagasi* IgG, selectively in the dogs immunized with Leishvaccine (Fig. 5I). No significant correlations were found between the anti-*Leishmania* IgG profiling (including IgG, IgG1 and IgG2) and the frequency of IL-4⁺CD4⁺ T-cells (Fig. 5J) neither the frequency of SLA-induced IFN- γ ⁺ T-cell subsets (data not shown).

4. Discussion

Despite the large amount of immunological data derived from clinical studies of CVL [34–36] there are still limited data available regarding the immune response triggered by anti-CVL vaccines [8–14,24,37–40]. The present work attempted to perform a phenotypic/functional analysis of canine peripheral blood cells to understand the immunological mechanisms related to immunogenicity elicited by Leishvaccine and Leishmune[®], focusing on three

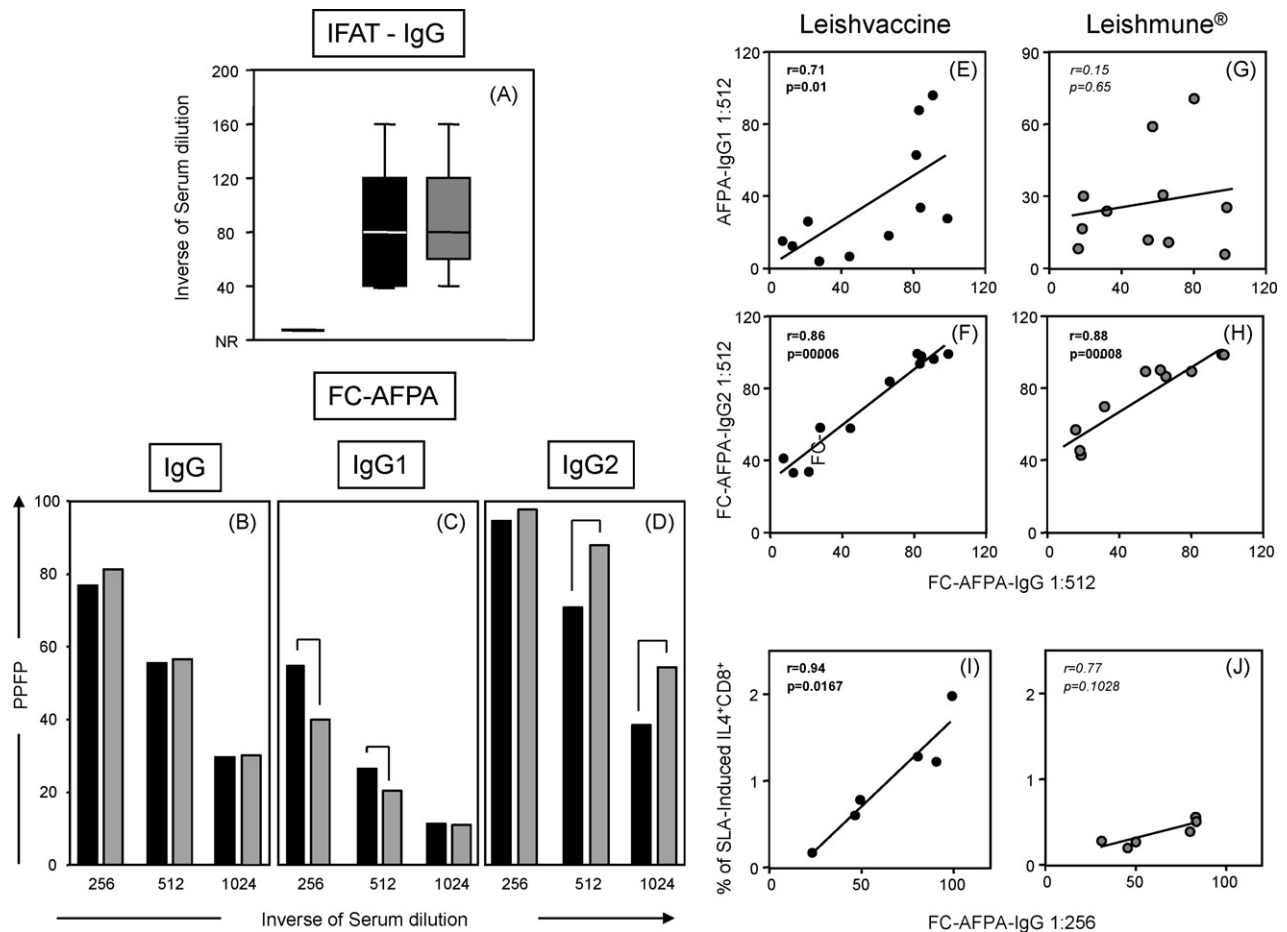


Fig. 5. Profile of seric anti-*Leishmania* IgG in German Shepherd dogs following Leishvaccine (■) and Leishmune[®] (□) vaccination regimens as compared to unvaccinated controls (□). IgG reactivity was detected by indirect immunofluorescence assay test/IFAT (A) as well as semi-quantitative (1:256–1:10,024 sera dilutions) flow cytometric detection of anti-fixed *L. chagasi* promastigotes/FC-AFPA-IgG, IgG1 and IgG2 (B–D). The results are expressed as median of IgG reactivity defined as inverse of serum dilution for IFAT and as the percentage of positive fluorescent parasites (PPFP) for FC-AFPA-IgG, IgG1 and IgG2. The data demonstrated that despite no changes in the IFAT profile, the median FC-AFPA-IgG1 and IgG2 PPFP values suggested distinct patterns of IgG reactivity between Leishvaccine and Leishmune[®] vaccinated dogs highlighted by connecting lines. Correlation analyses further re-emphasize the distinct association between the FC-AFPA-IgG reactivity and the FC-AFPA-IgG1 and FC-AFPA-IgG2 at serum dilution 1:512 observed in Leishvaccine (E and F) and Leishmune[®] vaccinated dogs (G and H). Additional correlation analysis point toward a positive correlation between the FC-AFPA-IgG reactivity at serum dilution 1:256 and the % of SLA-induced IL-4⁺CD8⁺ T-cells.

major aspects of the immune response, including: T-cell-derived cytokines, nitric oxide production by peripheral blood monocytes and seric anti-*L. (Leishmania) chagasi* IgG subclass patterns.

Cumulative studies show that the protective mechanisms are mainly associated with a specific type-1 immune response, specially linked with high IFN- γ secretion, whereas the role of type-2 immune response in the susceptibility to *Leishmania* infection dogs still remains to be elucidated [15,33,36,41–44]. In this context, a critical matter for screening and development of anti-leishmanial vaccines in CVL is to define *Leishmania* antigens and adjuvant systems that elicit a favorable and sustained type-1 cytokine environment *in vivo*. Herein, our data showed that despite the intrinsic differences in the immune response, intervention with either Leishvaccine or Leishmune[®] vaccines was accompanied by high levels of IFN- γ ⁺ T-cells, mainly due to an increase of the percentage of IFN- γ ⁺CD4⁺ T-cell subset, with an expressive increase observed upon SLA-stimuli, suggesting that both vaccines have the promising ability to elicit the establishment of anti-*Leishmania* immune mechanisms. Additionally, Leishvaccine immunized dogs showed increased levels of IL-4⁺ T-cells, with involvement of both CD4⁺ and CD8⁺ T-cell subsets. Consistent with this distinct cytokine profiling, we have previously reported that while Leishvaccine is able to trigger a mixed immunological profile involving changes in CD4⁺ and CD8⁺ T-cells as well as in the B-cell compartment,

Leishmune[®] elicited selective changes on the cellular adaptive immune response, mainly linked to the activation CD8⁺T-cell [24]. Since CD4⁺ T-cells are the major source of IFN- γ and the activation of CD8⁺ T-cells is an important phenotypic feature observed in the peripheral blood of Leishvaccine and Leishmune[®] vaccines, we hypothesize that the cross-talk between CD4⁺ and CD8⁺T-cells may underlay the protective mechanism triggered by these immunobiologicals. In this context, the CD4⁺ T-cells derived IFN- γ could represent the link to activate the CD8⁺ T-cell functions, already reported to play a crucial role in controlling the primary *Leishmania* infection in concurrence with the early shift in the type-2 toward a type-1 immune response [45,46]. It is possible that the multiple interactions induced by the whole crude *L. amazonensis* antigens (carbohydrate, protein, lipid and nucleic acid) would be the basis for the mixed IFN- γ and IL-4 cytokine pattern following immunization with Leishvaccine, characterizing the multiplicity of interactions with a wide range of cell surface receptors distributed on several cell subsets. On the other hand, the purified nature of FML antigen (glycoprotein) support the more selective IFN- γ linked cytokine profile observed in Leishmune[®] vaccines, with selective involvement of T-cells. We have previously suggested that besides the molecular nature of the antigens the adjuvants could also affect the properties of cell-mediated immunity [24]. We hypothesized that the BCG-based adjuvant in association with the complex whole crude *L.*

amazonensis antigen formulation used in Leishvaccine could potentiate the establishment of the mixed cytokine profile observed following vaccination. On the other hand, the saponin-based adjuvant used in Leishmune® would prompt a more selective cellular immune response as previously reported [37].

There is a current consensus amongst immunologists that rather than a shift toward a polarized cytokine pattern, the fine balance between type-1 and type-2 cytokines derived from distinct cell sources, may be more relevant for directing immune-mediated mechanisms that drive the disease outcome, as well as critical for determining the success of immunoprophylactic tools. We have recently proposed a novel strategy to assess the overall cytokine profile of circulating leukocytes, since the conventional strategies may not capture the global cytokine imprint, and does not reflect the panoramic cytokine profile of the wide range of circulating T-cell subsets [30]. Using this approach, we have characterized the T-cell overall cytokine profile triggered by Leishvaccine and Leishmune® vaccination regimens. Our results highlight that while a resultant type-1 cytokine pattern was observed in 50% of dogs immunized with Leishvaccine, a predominance of type-1 cytokine profile was observed in over 80% of Leishmune® vaccines. A predominant type-2 cytokine pattern was the hallmark of unvaccinated dogs. This type-2 related cytokine profile has been already reported in unvaccinated dogs [47]. Together, our cytokine findings highlighted that both immunobiologicals display potential applicability to drive a type-2 cytokine pattern observed in unvaccinated dogs toward a type-1, presumably protective against CVL.

It has been suggested that the IFN- γ -induced L-arginine nitric oxide (NO) pathway mediated by monocytes/macrophages is one of the major effector mechanism involved in the protective immune response in dogs infected with *Leishmania* [19,20]. Our data showed that Leishmune® vaccinated dogs presented increased levels of NO producing monocytes. Besides with the higher levels of IFN- γ CD4⁺ T-cells, the Leishmune® immunized dogs may possess the immunological events needed to establish a anti-*Leishmania* protective immune response. Pinelli et al. [20] have reported that recombinant canine IFN- γ alone is sufficient to induce *in vitro* NO production by canine macrophages. These findings re-emphasize that Leishmune® has the ability to activate phagocytes and support its high-quality immunogenic potential against CVL. Despite NO production and anti-leishmanial activity have been already reported in macrophages from dogs immunized with whole killed *L. infantum* promastigotes antigen [23], our data did not show alterations in the *in vitro* NO synthesis by monocytes from Leishvaccine immunized dogs, even in the presence of SLA. It is possible that the high levels of IL-4⁺ T-cells observed in these animals may count for the impaired NO production by circulating monocytes.

The humoral immune response has been usually associated with worsening outcome of CVL [48]. However, it has been proposed that differential responses of IgG subclasses can be in fact, indicative of dichotomous antibody response following anti-CVL immunoprophylaxis with Leishvaccine and Leishmune® [38,49]. Our results re-emphasize the existence of this dichotomous anti-*Leishmania* immunoglobulin profile, demonstrating that the Leishvaccine tend to induce higher levels of anti-*Leishmania* IgG1, whereas higher levels anti-*Leishmania* IgG2 are more likely to be observed in the Leishmune® vaccinated dogs. These findings are in agreement and re-enforce those previously reported by Fujiwara et al. [38] and de Oliveira Mendes et al. [49]. Additional analysis demonstrated the existence of a positive correlation between the reactivity of anti-*Leishmania* IgG with both IgG1 and IgG2 in Leishvaccine immunized dogs. On the other hand, Leishmune® vaccinated dogs, a positive correlation was observed selectively between anti-*Leishmania* IgG and IgG2. Moreover, our findings demonstrated a positive correlation between the frequency of SLA-induced IL-4+CD8⁺ T-cells and the reactivity of anti-*L. chagasi* IgG, selectively in the dogs immu-

nized with Leishvaccine, re-enforcing that the IL-4 triggered by this vaccine may also interplay a role interfering on the anti-IgG subclasses profile.

Altogether, our data pointed out to distinct immunological profiles elicited by Leishvaccine and Leishmune®, with the first triggering a mixed (IFN- γ and IL-4) cytokine pattern besides upper levels of anti-*Leishmania* IgG1, whereas the former induced an immunological pattern characterized by enhanced levels of IFN- γ , NO and anti-*Leishmania* IgG2. It is important to notice that the ability of both immunobiologicals to activate T-cell-derived IFN- γ synthesis suggested their high-quality immunogenic potential against CVL. These findings added support to further investigation focusing on perspectives of rational improvement of the antigenic composition as well as the adjuvant nature used for these vaccines formulation that might impact their immunoprophylactic effectiveness in the management of CVL.

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