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 stimulated 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-
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 lypophilized Bacille Calmette-Guérin, BCG (Fundação Ataúlo de Paiva, RJ, Brazil) as adjuvant. The first dose corresponded to 0.6 ml of Leishvaccine (360 μg of protein) plus 0.4 ml of physiological saline (NaCl 0.9% in distilled water) containing 400 μg of BGC dry-weight as adjuvant. The second dose corresponded to 0.6 ml of Leishvaccine (360 μg of protein) plus 0.3 ml of physiological saline (NaCl 0.9% in distilled water) containing 300 μg of BGC dry-weight as adjuvant. The third dose corresponded to 0.6 ml of Leishvaccine (360 μg of protein) plus 0.2 ml of physiological saline (NaCl 0.9% in distilled water) containing 200 μg of BGC 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 deacetylated) 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 promastigotes forms

L. chagasi promastigotes 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 × 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 polycarbonate tube and centrifuged at 1000 \times g for 10 min at 4–8 °C. The supernatant was discarded and the pellet washed twice (1000 \times g, 10 min, 4–8 °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 promastigotes.

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 m disposable syringe sterile filters of under aseptics 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 g/ml. Antigen preparation was stored in 100 \mu l aliquots at −70 °C prior the use in the in vivo 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 \times 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 suspension 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 \times 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 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 \times 10^-5 M) and 10% FBS. Aliquots of 100 \mu l of the PBMC suspension (1 \times 10^6 cells/well) were added to quadruplicate wells following the addition of 100 \mu l of RPMI 1640 (control cultures, CC) or 100 \mu l of soluble L. chagasi antigen (SLA) at final concentration of 25 pg/ml (stimulated cultures, SLA). Cultures were submitted to incubation in 5% CO2 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 g/ml and cultures submitted to an additional period of 4 h of incubation in 5% CO2 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-Aacetate). For this purpose, a 500 \mu l aliquot of whole blood was incubated in the presence of 500 \mu 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 g/ml and BFA at final concentration of 10 \mu g/ml. The positive control culture was maintained for 4 h in 5% CO2 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 l of FACS buffer and 200 \mu l aliquots incubated in 5 ml polystyrene tubes (Becton Dickinson, Franklin Lakes, NJ, USA) in the presence of 60 \mu l of previously diluted anti-canine CD4-1:320 (rat IgG2a, clone YK4X302.9) 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 l of PE-labeled anti-bovine cytokines mAb that cross-react with canine cytokines (as reported by the manufacturer), including anti-IFN-γ (clone CC3032) 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 FACSscan instrument (Becton Dickinson, Mountain View, CA) interfaced to an apple G3 FACSStation. The Cell-QuestTM 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-γ*CD4+, IL-4*CD4+, IFN-γ*CD8+ and IL-4*CD8+) within the lymphocyte logical gate.

2.5. Analysis of nitric oxide (NO) synthesis

The concentration of nitrite (NO2−) released in the supernatant of in vitro PBMC cultures was measured using the Griess reaction [28]. Briefly, a 100 \mu l aliquot of cell-free culture supernatant was mixed with 100 \mu l of Griess reagent (1% sulphanilamide, 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 μM. 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 (μM). 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 μl aliquots of the pre-fixed parasite suspension (5.0 × 10⁵ parasites/well) were incubated at 37 °C for 30 min in the presence of 50 μl 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 μl of PBS-3% FBS (1000 × g for 10 min, 4 °C). The parasites were re-incubated in the dark, for 30 min at 37 °C in the presence of 50 μl 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 Bollections laboratories Inc. (Montgomery, TX, EUA). After incubation parasites were submitted to two washing procedure using 150 μl of PBS-3% FBS (1000 × g, 10 min, 4 °C), fixed with 200 μl 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 × SSC dot plots. The relative FITC fluorescence intensity of the selected promastigotes was then quantified on single color fluorescence-1 (FITC/FL1) histograms. The histogram distributions, obtained for the internal control of unspecific binding of the second step reagents, were used to set a 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-
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 Leishvaccine and Leishmune® vaccinated dogs.
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 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 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 (Fig. 3B). Distinct cut-offs were employed for IFN-γCD4+ (A), IL-4CD4+ T-cells (B), IFN-γCD8+ (C) and IL-4CD8+ T-cells (D).

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-IgG2 reactivity was detected in Leishmune® vaccinated dogs (Fig. 5C). Further analyses demonstrated a positive correlation between FC-AFPA-IgG and FC-AFPA-IgG2 in Leishvaccine immunized dogs (Fig. 5D), whereas higher median FC-AFPA-IgG2 reactivity was detected in Leishmune® vaccinated dogs (Fig. 5C). 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 (M) 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.

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 changed 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 features 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 immunobiologics. In this context, the CD4+ T-cells derived INF-γ could represents 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 poten-
tiate the establishment of the mixed cytokine profile observed
following vaccination. On the other hand, the saponin-based adju-
vant 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 criti-
cal 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 charac-
terized 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 unvac-
cinated dogs. This type-2 related cytokine profile has been already
reported in unvaccinated dogs [47]. Together, our cytokine find-
ings highlighted that both immunobiologicals display potential applica-
ability 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 t-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 pro-
tective 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 sup-
port 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 alter-
ations in the in vitro NO synthesis by monocytes from Leishvaccine
immunized dogs, even in the presence of MLA. 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 immunopro-
phylaxis with Leishvaccine and Leishmune® [38,49]. Our findings
demonstrated a positive correlation between the frequency of SLA-induced IL-4+CD8+ T-cells and
IgG2. Moreover, our findings demonstrated a positive correla-
tion 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 trig-
erg ing 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 effective-
ness in the management of CVL.

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