



The effects of increased heme oxygenase-1 on the lymphoproliferative response in dogs with visceral leishmaniasis



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ABSTRACT

Canine visceral leishmaniasis (CVL) is known to affect the cellular immunity of infected dogs, through impairing lymphoproliferation and microbicidal mechanisms. This study examined heme oxygenase-1 (HO-1) and its metabolites, oxidative stress and IL-10 levels in CVL and investigated correlations between these parameters. Additionally, the effects of HO-1 inhibition on the lymphoproliferative response and cytokine production in lymph node cells (LNCs) from infected dogs were evaluated. Forty-four dogs, 24 controls and 20 dogs with CVL were selected. Plasma and splenic levels of HO-1, haptoglobin, soluble CD163 receptor, ferritin and IL-10 were determined using capture ELISA. The HO-1 levels and relative gene expression in peripheral blood and bone marrow mononuclear cells were also determined. LNCs proliferation was evaluated with an HO-1 activator and with an HO-1 inhibitor, in the presence of the *Leishmania infantum* soluble antigen (SAgL), using flow cytometry. HO-1, IL-2, IFN-gamma and IL-10 were also determined in these cultures using capture ELISA. Infected dogs presented oxidative stress and increased HO-1 levels and relative gene expression, with correlation between oxidative stress and HO-1. The substances from heme metabolism and IL-10 were also elevated in the plasma and spleens of infected dogs. IL-10 and HO-1 levels were positively correlated with one another. Inhibition of HO-1 increased LNCs proliferation and decreased IL-10 and IL-2 production in the presence of SAgL. The increased HO-1 metabolism observed in CVL is probably associated with oxidative stress and increased IL-10, which could be one of the mechanisms responsible for inhibition of the lymphoproliferative response in sick dogs.

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Abbreviations: ALP, alkaline phosphatase; ALT, alanine aminotransferase; AST, aspartate aminotransferase; BMNC, bone marrow mononuclear cells; CBC, complete blood count; CFSE, carboxyfluorescein diacetate succinimidyl ester; CK, creatine kinase; CO, carbon monoxide; ConA, concanavalin-A; CoPP, cobaltic protoporphyrin IX chloride; CPDA-1, citrate-phosphate-dextrose-adenine-1; CVL, canine visceral leishmaniasis; EDTA, ethylenediaminetetraacetic acid; ELISA, enzyme-linked immunosorbent assay; GGT, gamma-glutamyl transferase; HO-1, heme oxygenase-1; IL, interleukin; LDH, lactate dehydrogenase; LNCs, lymph node cells; MCHC, mean corpuscular hemoglobin concentration; MCV, mean corpuscular volume; PBMC, peripheral blood mononuclear cells; PBS, phosphate buffer solution; PCV, packed cell volume; PHA-M, phytohemagglutinin-M; qPCR, real-time polymerase chain reaction; qRT-PCR, real-time reverse transcription-polymerase chain reaction; RBC, red blood cells; ROS, reactive oxygen species; RT, reverse transcription; SAgL, soluble antigen of *Leishmania infantum*; sCD163, soluble CD163 receptor; SnMSP, Sn(IV) mesoporphyrin IX dichloride; TAC, total antioxidant capacity; TBARS, thiobarbituric acid reactive substances; TOC, total oxidant capacity; TPP, total plasma protein; UPC, urinary protein/creatinine ratio; VL, visceral leishmaniasis; WBC, white blood cells.

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1. Introduction

Visceral leishmaniasis (VL) is a parasitic zoonosis that is widely distributed throughout the world and is among the six most important tropical diseases according to the World Health Organization (WHO, 2015). Over recent years, increases in the incidence of VL have been reported in Brazil (Alvar et al., 2012). The increased incidence of human cases appears to be associated with increases in canine infections (Coura-Vital et al., 2011; Nunes et al., 2010) because dogs are the primary source of this disease in urban areas (Gramiccia and Gradoni, 2005).

This dog-susceptible disease is characterized by skin lesions, generalized lymphadenopathy, progressive weight loss with muscle atrophy and weakness, exercise intolerance, decreased appetite, lethargy, splenomegaly, polyuria and polydipsia, eye lesions, epistaxis, onychogryphosis, vomiting and diarrhea (Solano-Gallego et al., 2009).

It has been accepted that susceptibility is associated with an exacerbated humoral immune response (Th2) and a reduced cellular immune response (Th1) (Alvar et al., 2004; Baneth et al., 2008). Increased production of IL-10, a major Th2 cytokine, is associated with progression of the canine infection (Boggiatto et al., 2010), while higher expression of IFN- γ , a Th1 cytokine, is correlated with a lower blood parasite load (Solano-Gallego et al., 2016).

Recently, it has been suggested that oxidative stress in infected dogs presenting clinical signs of canine visceral leishmaniasis (CVL) is associated with increased production of reactive oxygen species (ROS), which may be a critical mechanism of the pathogenesis of this disease (Almeida et al., 2013b; Bildik et al., 2004; Heidarpour et al., 2012). This increased ROS production has been demonstrated in human macrophages (Luz et al., 2012) and canine neutrophils (Almeida et al., 2013a,b).

The mechanisms associated with CVL susceptibility are not well-understood. Information regarding CVL disease pathogenicity is fundamental for future treatments for dogs. Involvement of heme oxygenase-1 (HO-1) in human VL susceptibility has been reported (Das et al., 2013; Luz et al., 2012). HO-1 is responsible for degrading the heme in hemoglobin into iron, carbon monoxide and biliverdin, which is rapidly converted into bilirubin (Tenhun et al., 1969). Under physiological conditions, hemoglobin from erythrocyte destruction binds with haptoglobin and this complex is then internalized by macrophages via the CD163 receptor. Inside the macrophage, heme is metabolized by HO-1 and iron stimulates ferritin production (Kristiansen et al., 2001).

Human patients with clinical VL have been found to present increased HO-1 levels, which then decreased after disease treatment. This clinical finding demonstrates the involvement of HO-1 in VL susceptibility (Das et al., 2013; Luz et al., 2012). Furthermore, both the lipophosphoglycan and promastigote parasite forms of *Leishmania* spp. are responsible for increasing HO-1 expression in macrophages, resulting in the persistence of the infection in humans (Luz et al., 2012). Because heme is necessary for complete NADPH oxidase activation, the underlying mechanism causing persistence of this macrophage infection is possibly associated with heme degradation caused by HO-1, which prevents complete activation of NADPH oxidase. Thus, ROS production by infected macrophages is impaired, and this allows the parasite to survive inside the host cell (Pham et al., 2005).

HO-1 also presents immunoregulatory functions, mainly those associated with lymphocytes, by inhibiting activation and proliferation of T lymphocytes and stimulating apoptosis in these cells (Pae et al., 2004). To date, the involvement of HO-1 in the lymphoproliferative response of dogs with CVL has not been studied. However, it is known that infected dogs with clinical signs of CVL exhibit low lymphocyte proliferative responses (Pinelli et al., 1994). The mechanisms involved in this impaired cell response may be related to

increased T lymphocyte apoptosis (Chiku et al., 2016; Lima et al., 2012; Perosso et al., 2014). Therefore, gaining an understanding of the mechanisms that improve the lymphoproliferative response in sick dogs is critical for development of treatments and elimination of this parasite in dogs.

For the present study, it was assumed that HO-1 production is activated under stress conditions, that oxidative stress occurs in CVL and that anemia is a common finding in relation to this condition. Hence, this study aimed to evaluate HO-1 levels and relative gene expression, along with HO-1 metabolites, in infected dogs and to determine the correlations between HO-1, oxidative stress markers, IL-10 levels and parasite load. Additionally, we evaluated the regulatory role of HO-1 relating to the antigen-specific proliferative response of lymph node cells from infected dogs and determined the effects of HO-1 activation and inhibition on the lymphoproliferative response and on IL-10, IL-2 and IFN- γ production.

2. Material and methods

2.1. Ethics approval

This study was approved by the Ethics Committee for Animal Experimental Research (Comitê de Ética em Pesquisa Experimental Animal, COBEA) and the Ethics Committee for Animal Use (Comitê de Ética no Uso Animal, CEUA) of the School of Veterinary Medicine (Faculdade de Medicina Veterinária, FMVA) at the Araçatuba Campus of São Paulo State University (Universidade Estadual Paulista “Júlio de Mesquita Filho”, Unesp), under procedural number FOA-00532-2013. The participation of dogs in the control group was authorized by their keepers, who provided free and informed consent.

2.2. Dog selection

The control group consisted of 24 dogs of various breeds (10 males and 14 females; average age of 4 ± 2 years), which were recruited in the city of Araçatuba. The keepers of these dogs authorized their participation in the experiment. All the dogs were clinically healthy with normal physical examinations. These dogs presented normal hematological, biochemical and urinary profiles, and negative indirect ELISA reactions (Lima et al., 2003) for the *L. infantum* antigen obtained within the last three months before the experiment. There was no *Leishmania* DNA amplification (Perosso et al., 2014) in the spleen or bone marrow samples taken at the time of the experiment.

The infected group consisted of 20 dogs (10 males and 10 females; average age of 5 ± 2 years) of varying breeds that were naturally infected with *Leishmania* spp. These dogs were acquired from the Zoonosis Control Center of Araçatuba. General clinical examinations, complete blood counts, plasma biochemistry analyses and urinalyses were performed on these dogs to stage the disease in accordance with the LeishVet Consensus (Solano-Gallego et al., 2009). All the animals were reactive to total crude antigen of *L. infantum* through indirect ELISA (Lima et al., 2003) and presented positive *Leishmania* DNA amplification (Perosso et al., 2014) in spleen and bone marrow samples.

2.3. Sample collection and laboratory analysis

Prior to blood collection, the dogs were kept fasting for 8–12 h. Twenty-one milliliters of blood were extracted via jugular venipuncture; 20 mL was placed in heparinized tubes (BD Vacutainer®, Franklin Lakes, NJ, USA) and 1 mL of ethylenediaminetetraacetic acid (EDTA)-treated blood was used to determine the complete blood count (CBC).

Heparinized blood was used to obtain plasma and peripheral blood mononuclear cells (PBMC) were isolated using a gradient density of 1.077 g/mL (Histopaque-1077[®], Sigma-Aldrich Co., St. Louis, USA), in accordance with the manufacturer's recommendations.

The CBC, plasma biochemistry, urinalysis and urinary protein/creatinine ratio (UPC) were determined using methods described previously (Almeida et al., 2013b).

Bone marrow obtained from the right iliac crest using a Jamshidi needle was placed with citrate-phosphate-dextrose-adenine-1 (CPDA-1) anticoagulant (JP Ind. Pharmaceutical, Ribeirão Preto, SP, Brazil) (four parts of bone marrow to one part of anticoagulant). Isolation of bone marrow mononuclear cells (BMMC) followed the same protocol as for PBMC.

Partial splenectomy was performed by means of surgical excision, following the protocol described by Lima et al. (2012).

2.4. Oxidative stress markers

Total antioxidant capacity (TAC) was determined colorimetrically using a reduction method involving the cation ABTS (2,2'-azino-bis-[3-ethylbenzothiazoline-6-sulfonic acid]), following methods published previously (Erel, 2004).

Total oxidant capacity (TOC) was determined colorimetrically using the xylenol orange method, following methods described previously (Erel, 2005).

Lipid peroxidation was determined using thiobarbituric acid reactive substances (TBARS) in an automated plate reader (Readwell Touch, Robonik Private, Ltd., Navi Mumbai, Maharashtra, India) at 545 nm, in accordance with previously described methods (Hunter et al., 1985). Lipid peroxidation was obtained after sample absorbance interpolation using a standard curve, with concentrations ranging from 0 to 100 μmol of malondialdehyde/L. All reagents were obtained from Sigma-Aldrich (Sigma-Aldrich Co., St. Louis, USA).

2.5. Lymphoproliferation assay

The lymphoproliferation assay was performed on 15 infected dogs using lymph node cells (LNCs). After each dog had been sacrificed, its right popliteal lymph node was removed and macerated to obtain LNCs ($50 \times 10^6/\text{mL}$ in RPMI 1640 medium without calf bovine serum). LNCs were stained with the dye carboxyfluorescein diacetate succinimidyl ester (2.5 μM) (CFSE, CellTrace[™] CFSE Cell Proliferation Kit, Invitrogen Molecular Probes, Eugene, OR, USA) for 10 min at 37 °C, in accordance with the manufacturer's recommendations and with the previously published protocol (Lyons and Parish, 1994).

Stained LNCs were cultured in sterile 24-well plates ($10 \times 10^6/\text{mL}$) in RPMI 1640 medium (Sigma-Aldrich Co., St. Louis, USA) alone (control) or with the HO-1 activator cobaltic protoporphyrin IX chloride (50 μM) (CoPP), the HO-1 inhibitor Sn(IV) mesoporphyrin IX dichloride (50 μM) (SnMsP) (Santa Cruz Biotechnology, Dallas, TX, USA) or 20 μg of protein/mL of soluble antigen from *L. infantum* (SAGL) (MHOM/BR/00/MERO2). Cultures of two combined treatments (CoPP + SAGL and SnMsP + SAGL) were also performed. The mitogens concanavalin-A (ConA, 5 mg/mL) and phytohemagglutinin-M (PHA-M, 5 $\mu\text{L}/\text{mL}$) were used as positive controls. CFSE-unmarked LNCs were used to verify CFSE staining.

LNCs were cultured for 6 days at 37 °C and at 5% CO₂. Ten thousand events were acquired in a flow cytometer (BD C5 Accuri Flow Cytometer, Ann Arbor, MI, USA). The analysis was performed using BD Accuri C6 Software (v. 1.0.264.21, Ann Arbor, MI, USA). Cell populations with a size and complexity similar to those of the lymphocyte population were selected.

The assays were repeated in duplicate under the same conditions without CFSE staining. Supernatants were used to determine IL-2, IFN-gamma and IL-10 levels. Pellet cells were used to determine HO-1 levels and relative expression.

SAGL production was carried out using a protocol described previously (Scott et al., 1987) with the following modifications: *L. infantum* promastigotes ($10^8/\text{mL}$) in a protease inhibitor solution (Complete, EDTA-free, Roche, Indianapolis, IN, USA) were sonicated and the suspension was centrifuged at 20,000g for 30 min at 4 °C. The supernatant was removed and centrifuged again at 20,000g for 4 h at 4 °C. The supernatant from the second centrifugation was collected, dialyzed and sterilized, and its protein content was measured using the bicinchoninic acid method (BCA Protein Assay Kit[™], Thermo Scientific, Rockford, IL, USA), in accordance with the manufacturer's recommendations, and was stored at –80 °C for later use.

2.6. Substances in HO-1 metabolic pathway and interleukins

Ferritin, sCD163 and haptoglobin levels were determined in 50- μL samples of plasma or spleen homogenate that were prepared in accordance with methods described previously (Corrêa et al., 2007), using capture ELISA with canine-specific reagents (Novateinbio, Woburn, MA, USA) following the manufacturer's recommendations.

HO-1 levels were determined in 50 μL of plasma, LNCs, PBMC or BMMC homogenate using capture ELISA with canine-specific reagents (Novateinbio, Woburn, MA, USA), following the manufacturer's recommendations. Homogenates were prepared using 5×10^6 cells resuspended in 120 μL of 0.1 M phosphate buffer solution (PBS, pH 7.2), following sonication (2×20 s, 5 W, 4 °C). Further sample dilution was not necessary. The values were normalized using the protein levels that had been determined spectrophotometrically using the bicinchoninic acid method (BCA Protein Assay Kit[™], Thermo Scientific, Rockford, IL, USA).

IL-2, IL-10 and IFN-gamma levels were determined using capture ELISA and commercial reagents (Canine IL-10 DuoSet, Canine IL-2 DuoSet and Canine IFN-gamma DuoSet; R&D Systems, Minneapolis, MN, USA), in accordance with the manufacturer's recommendations.

Plasma-free heme was determined biochemically using a commercial reagent (QuantiChrom Heme[™] Assay Kit Bioassay Systems, Hayward, CA, USA), following the manufacturer's instructions.

2.7. HO-1 relative gene expression

To evaluate the relative gene expression of HO-1 (*Hmox1*), RNA was extracted from 25 mg of spleen tissue or from 5×10^6 cells from the proliferation assay, BMMC and PBMC that had been stored at –80 °C, using commercial reagents (RNeasy[®] Mini Kit, Qiagen, Valencia, CA, USA), following the manufacturer's instructions. RNA eluted in nuclease-free water was quantified spectrophotometrically (NanoDrop ND-1000, Nano-Drop Technologies[®], Wilmington, DE, USA) with a 260/280 absorbance ratio varying from 2.0 to 2.2. Samples were stored at –80 °C until use in the reverse transcription reaction.

Reverse transcription (RT) was performed using a commercial reagent (QuantiTect[®] Reverse Transcription Kit, Qiagen, Valencia, CA, USA), in accordance with the manufacturer's specifications, from 100 ng of RNA from spleen samples and 300 ng from LNCs, PBMC and BMMC, all at a final volume of 20 μL . To produce a standard curve, two samples of each experimental group from each tissue were randomly selected and RT was performed using 1 μg of RNA. The samples from each tissue were then pooled. All the

cDNA was stored at -80°C and pooled cDNA was diluted (1:5) immediately before sample amplification.

The primers (Sigma-Aldrich Co., St. Louis, USA) for each gene are described in Table 1.

Real-time reverse transcription-polymerase chain reactions (qRT-PCR) were performed in a thermocycler (CFX96™ Real-Time System, Bio-Rad, Hercules, CA, USA), using a commercial reagent (QuantiNova™ SYBR® Green PCR Kit, Qiagen, Valencia, CA, USA) and $2\ \mu\text{L}$ of cDNA from the sample or a standard curve following the manufacturer's instructions.

In triplicate, the samples and standard curve were incubated at 95°C for 2 min and then subjected to 40 cycles of 95°C for 15 s and 60°C for 30 s, followed by a dissociation curve. The efficiency values, correlation coefficients and slopes ranged from 97.4 to 101.7%, 0.994–0.999 and -3.386 to -3.283 , respectively. Nuclease-free water (Sigma-Aldrich Co., St. Louis, USA) was used as the negative control for the reaction.

HO-1 relative gene expression was determined after interpolating the cycle threshold (Ct) of the samples on the standard curve. The values were normalized using the geometric mean of the reference gene expression (beta-actin and HPRT-1) and are represented as the relative gene expression.

2.8. Parasite load

Parasite load was determined in infected dogs from 25 mg of spleen and $200\ \mu\text{L}$ of bone marrow. The DNA extraction and quantification, along with the real-time polymerase chain reaction (qPCR), were performed as previously described (Perosso et al., 2014).

2.9. Statistics

Variables were tested for normality (Shapiro-Wilk test) and homoscedasticity (Bartlett test). Unpaired *t* or Mann-Whitney tests were then used to compare the controls and infected dogs. To assess the effects of treatment on the lymphoproliferative response, an analysis was performed using the Friedman test with Dunn's post-test after testing for normality. Correlations were determined using the Spearman correlation coefficient. Tests were performed using field-specific software (GraphPad Prism v.6.00 for Windows, GraphPad Software, La Jolla, CA, USA; www.graphpad.com). The results were considered to be statistically significant when $P < 0.05$.

3. Results

3.1. Characterization of infected dogs

According to the staging criteria proposed by the LeishVet Consensus (Solano-Gallego et al., 2009), the infected dogs were predominantly in the stage II (moderate) and stage III (severe) disease states. They presented clinical signs such as lymphadenomegaly, skin lesions (exfoliative and seborrheic dermatitis, ulcerations and onychogryphosis), anorexia, weight loss and epistaxis. The main laboratory findings in the dogs in these stages of CVL were mild non-regenerative normocytic and hypochromic anemia, hyperproteinemia due to hyperglobulinemia, hypoalbuminemia, normal biochemical renal function (creatinine $<1.4\ \text{mg/dL}$) and marked proteinuria (UPC >0.5 for stage II and UPC >1.0 for stage III). The infected dogs also presented high levels of antibodies against total crude *L. infantum* antigen in the indirect ELISA reaction (Table 2).

Other hematological disorders in the dogs with CVL included basophilia and neutrophilia, with or without associated leukocytosis, lymphopenia and eosinopenia (Table 2). Regarding the biochemical profile, the infected dogs presented increased AST, CK

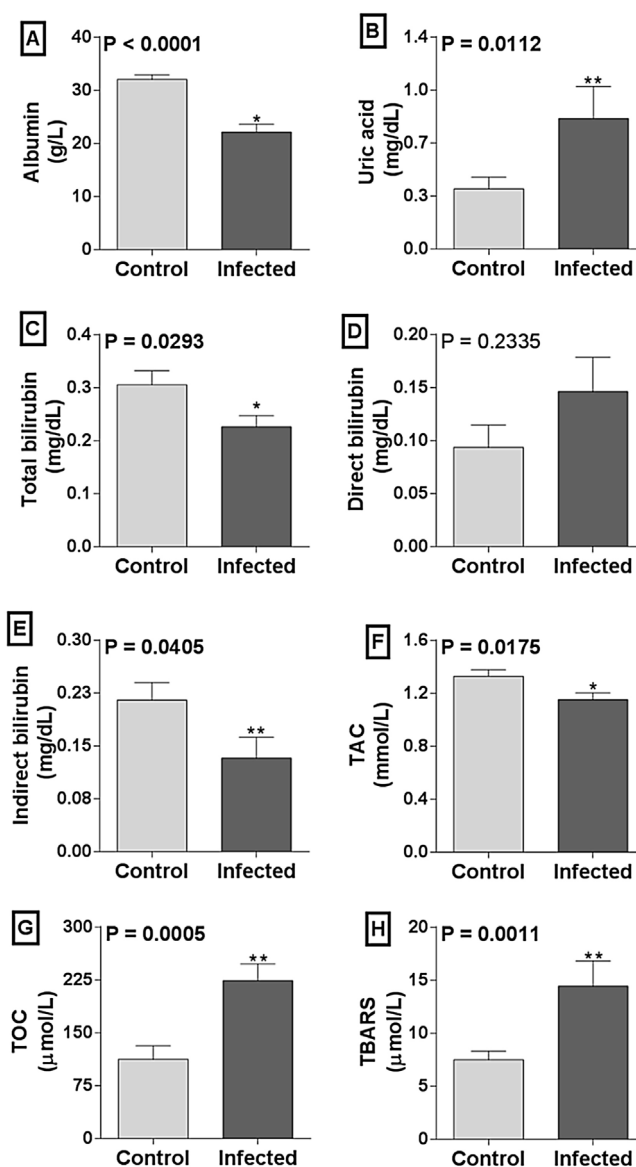


Fig. 1. Oxidative stress markers: albumin (A), uric acid (B) total bilirubin (C), direct bilirubin (D), indirect bilirubin (E), total antioxidant capacity (TAC, F), total oxidant capacity (TOC, G) and plasma lipid peroxidation measured using thiobarbituric acid reactive substances (TBARS, H) in control dogs (Control, $n = 24$) and dogs with naturally occurring canine visceral leishmaniasis dogs (Infected, $n = 20$). The graphs show the mean and standard error of the mean. Statistically significant differences are indicated by * (unpaired *t*-test) or ** (Mann-Whitney test).

and LDH activity and decreased albumin, total and indirect bilirubin and calcium levels, compared with the uninfected dogs (Table 2).

The control dogs did not have any clinical or laboratory abnormalities (Table 2).

3.2. Oxidative stress in CVL

Confirming previous reports that oxidative stress occurs in CVL (Almeida et al., 2013b), we demonstrated that marked oxidative stress was present in the infected dogs. This can be seen through the reduced levels of antioxidants such as albumin (Fig. 1A), total bilirubin (Fig. 1C), indirect bilirubin (Fig. 1E) and TAC (Fig. 1F), as well as through the increased levels of oxidants measured via TOC (Fig. 1G) and lipid peroxidation (Fig. 1H) in the infected dogs. Antioxidant uric acid (Fig. 1B) was elevated in dogs with CVL. There were no differences in the direct bilirubin content (Fig. 1D).

Table 1
Sequences of forward (F) and reverse (R) primers for HO-1, beta-actin and HPRT-1 genes.

Gene	Primer	Sequence (5' → 3')	Product (bp)	GenBank access	Reference
HO-1	F	GCGTCGACTTCTTCACCTTC	195	AY563546.1	Primer 3 Plus ^a
	R	GGTCCTCAGTGTCTTGCTC			
Beta-actin	F	CCAGCAAGGATGAAGATCAAG	100	AF021873	Peters et al. (2007)
	R	TCTGCTGGAAGGTGGACAG			
HPRT-1	F	CACTGGGAAAACAATGCAGA	123	AY283372	Peters et al. (2007)
	R	ACAAAGTCAGGTTTATAGCCAACA			

^a Available from: www.primer3plus.com.

Table 2
Hematological, biochemical, urinary and serological parameters (mean ± SD) of control dogs (Control, n = 24) and dogs with naturally occurring canine visceral leishmaniasis (Infected, n = 20).

Parameter	Control	Infected	P-value	Reference range
Hematological parameters				
PCV (%)	46.70 ± 5.60	28.45 ± 8.25*	<0.0001 [‡]	37–55
RBC (10 ¹² /L)	6.44 ± 0.88	4.44 ± 1.24 [†]	<0.0001 [‡]	5.5–8.5
Hemoglobin (g/dL)	15.13 ± 2.07	8.66 ± 2.71 [†]	<0.0001 [‡]	12–18
MCV (fL)	72.64 ± 2.80	63.85 ± 4.52*	<0.0001 [‡]	60–77
MCHC (%)	32.37 ± 1.66	30.29 ± 1.55*	0.0001 [†]	32–36
WBC (× 10 ⁹ /L)	10.83 ± 1.85	15.00 ± 9.92	0.5995 [†]	6.0–17.0
Neutrophils (× 10 ⁶ /L)	5973 ± 1786	12,740 ± 9506*	0.0022 [†]	3000–11,500
Lymphocytes (× 10 ⁶ /L)	3177 ± 916.3	1305 ± 1200*	<0.0001 [†]	1000–4800
Monocytes (× 10 ⁶ /L)	608.4 ± 435.2	707.7 ± 646.0	0.7586 [†]	150–1350
Eosinophils (× 10 ⁶ /L)	1029 ± 870.5	210.5 ± 280.4 [†]	<0.0001 [‡]	150–1250
Basophils (× 10 ⁶ /L)	0 ± 0	57.0 ± 92.59*	0.0051 [‡]	Rare
Platelet (× 10 ⁹ /L)	289.8 ± 79.61	237.6 ± 119.9	0.0953 [‡]	160–430
TPP (g/dL)	7.01 ± 0.98	9.74 ± 1.76 [†]	<0.0001 [‡]	6.0–8.0
Biochemical profile				
Total protein (g/L)	61.5 ± 10.6	86.7 ± 11.9*	<0.0001 [‡]	54–71
Albumin (g/L)	32.06 ± 3.98	22.12 ± 6.69*	<0.0001 [‡]	26–33
Globulin (g/L)	30.69 ± 10.17	63.51 ± 10.18*	<0.0001 [‡]	27–44
ALT (IU/L)	34.05 ± 10.48	33.86 ± 22.69	0.2684 [†]	21–102
AST (IU/L)	23.63 ± 5.59	58.05 ± 27.07*	<0.0001 [‡]	23–66
ALP (IU/L)	60.0 ± 40.4	86.7 ± 75.5	0.6285 [†]	20–156
GGT (IU/L)	3.15 ± 1.0	2.74 ± 1.65	0.0910 [†]	1.2–6.4
Total bilirubin (mg/dL)	0.30 ± 0.12	0.22 ± 0.09*	0.0293 [†]	0.1–0.5
Direct bilirubin (mg/dL)	0.09 ± 0.10	0.14 ± 0.13	0.1766 [†]	0.06–0.12
Indirect bilirubin (mg/dL)	0.21 ± 0.12	0.13 ± 0.12*	0.0405 [†]	0.01–0.49
Cholesterol (mg/dL)	163.6 ± 47.84	183.5 ± 64.56	0.4732 [†]	135–270
Triglycerides (mg/dL)	62.1 ± 23.7	68.2 ± 24.7	0.3403 [†]	20–112
Glucose (mg/dL)	83.9 ± 13.7	90.2 ± 16.0	0.1689 [‡]	68–118
Urea (mg/dL)	35.6 ± 9.5	35.4 ± 24.4	0.0534 [†]	10–50
Creatinine (mg/dL)	0.99 ± 0.18	0.99 ± 0.28	0.1827 [†]	0.5–1.5
CK (IU/L)	96.96 ± 59.55	320.8 ± 254.8*	<0.0001 [‡]	1.5–28.4
LDH (IU/L)	51.9 ± 38.2	155.0 ± 110.2*	<0.0001 [‡]	45–233
Calcium (mg/dL)	9.94 ± 1.03	8.92 ± 0.62*	0.0004 [‡]	9.0–11.3
Phosphorus (mg/dL)	4.66 ± 1.67	5.56 ± 1.69	0.0746 [†]	2.6–6.2
Urinary				
Density	1.043 ± 0.01	1.033 ± 0.01	0.1163 [‡]	>1.035
UPC	0.09 ± 0.07	1.66 ± 1.84*	<0.0001 [‡]	<0.5
Indirect ELISA Serology				
Optical density	0.052 ± 0.03	0.935 ± 0.211 [†]	<0.0001 [‡]	<0.270

Canine reference ranges are in accordance with Rizzi et al. (2010) for hematology, Kaneko et al. (2008) for biochemistry, Osborne et al. (1995) for urinalysis and Lima et al. (2003) for indirect ELISA serology.

Abbreviations: ALP, alkaline phosphatase; ALT, alanine aminotransferase; AST, aspartate aminotransferase; CK, creatine kinase; GGT, gamma-glutamyl transferase; LDH, lactate dehydrogenase; MCHC, mean corpuscular hemoglobin concentration; MCV, mean corpuscular volume; PCV, packed cell volume; RBC, red blood cells; TPP, total plasma protein; UPC, urinary protein/creatinine ratio; WBC, white blood cells.

* Statistically significant difference using unpaired *t*-test (†) or Mann-Whitney test (‡).

3.3. Increased HO-1 levels and relative expression are tissue-dependent

Increased HO-1 levels and relative expression in human VL patients have been correlated with disease susceptibility (Das et al., 2013; Luz et al., 2012). Until now, the role of HO-1 had not been investigated in relation to CVL. With the aim of ascertaining the involvement of HO-1 in the disease pathogeny of susceptible dogs,

we determined the levels and relative gene expression of HO-1 in infected dogs and compared them with healthy control dogs.

The infected dogs presented increased HO-1 levels in the plasma, spleen and PBMC; there was no difference in BMBC (Fig. 2A). Increased relative expression of HO-1 was observed in the PBMC and BMBC of infected dogs, with no significant differences in spleen tissue between infected and healthy dogs (Fig. 2B).

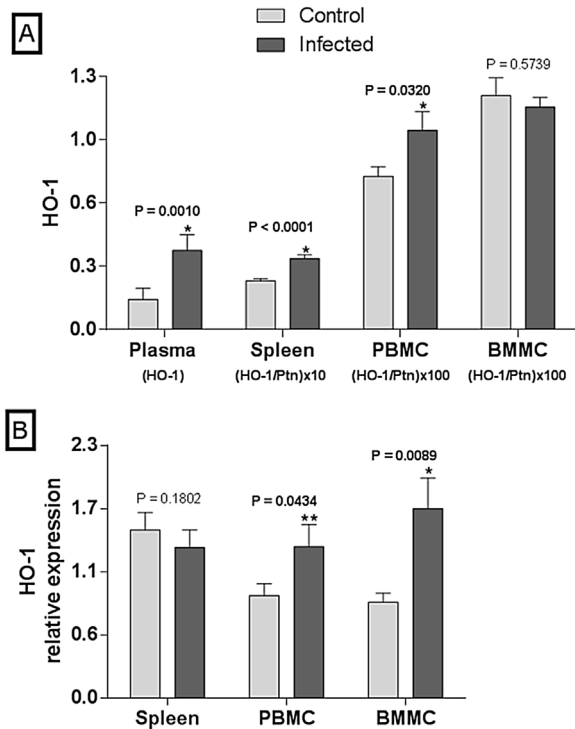


Fig. 2. Levels of heme oxygenase-1 (HO-1) determined in plasma, spleen homogenate, peripheral blood (PBMC) or bone marrow (BMNC) mononuclear cells normalized to protein levels in control dogs (Control, plasma HO-1 n=24 and other variables n=10) and dogs with naturally occurring canine visceral leishmaniasis (Infected, plasma HO-1 n=20 and other variables n=10) (A). Relative expression of HO-1 normalized using geometric mean of standard expression of the genes beta-actin and HPRT-1 in spleen, PBMC and BMNC of control dogs (Control, n=10) and dogs with naturally occurring canine visceral leishmaniasis (Infected, n=10) (B). The graphs show the mean and standard error of the mean. Statistically significant differences are indicated by * (unpaired t-test) or ** (Mann-Whitney test).

3.4. Increased levels of heme metabolism substances and IL-10 in CVL

In addition to determining HO-1, we also assessed the IL-10 and HO-1 enzymatic metabolic pathway in infected dogs by determining the levels of plasma-free heme and the levels of ferritin, haptoglobin and sCD163 in the plasma and spleen of dogs. Similarly to HO-1, the plasma and spleen levels of IL-10, ferritin, sCD163 and haptoglobin were elevated in the infected dogs (Figs. 3 and 4). The plasma-free heme levels were not altered by the infection (Fig. 3A).

3.5. HO-1 correlated with oxidative stress markers and IL-10, but not with the parasite load

With the aim of ascertaining the correlation between HO-1, oxidative stress and parasite load, these parameters were correlated with HO-1 levels and relative gene expression. We observed that the HO-1 levels in bone marrow had a positive correlation with plasma TOC and that the HO-1 levels in the spleen had a positive correlation with lipid peroxidation, while the spleen HO-1 levels had negative correlations with the plasma antioxidants albumin, total bilirubin, indirect bilirubin and TAC. The HO-1 levels in bone marrow also correlated negatively with albumin, total bilirubin and indirect bilirubin (Table 3).

The plasma and spleen levels of IL-10 correlated positively with plasma HO-1 levels, while spleen IL-10 also correlated positively with spleen HO-1 levels and BMNC HO-1 relative gene expression. There was no correlation between the parasite load and HO-1 in the infected dogs (Table 3).

3.6. Inhibition of HO-1 increases lymphoproliferation induced by antigens of *L. infantum* in LNCs from infected dogs

Because of the increased HO-1 metabolism in CVL, we aimed to determine whether inhibition of HO-1 might improve the

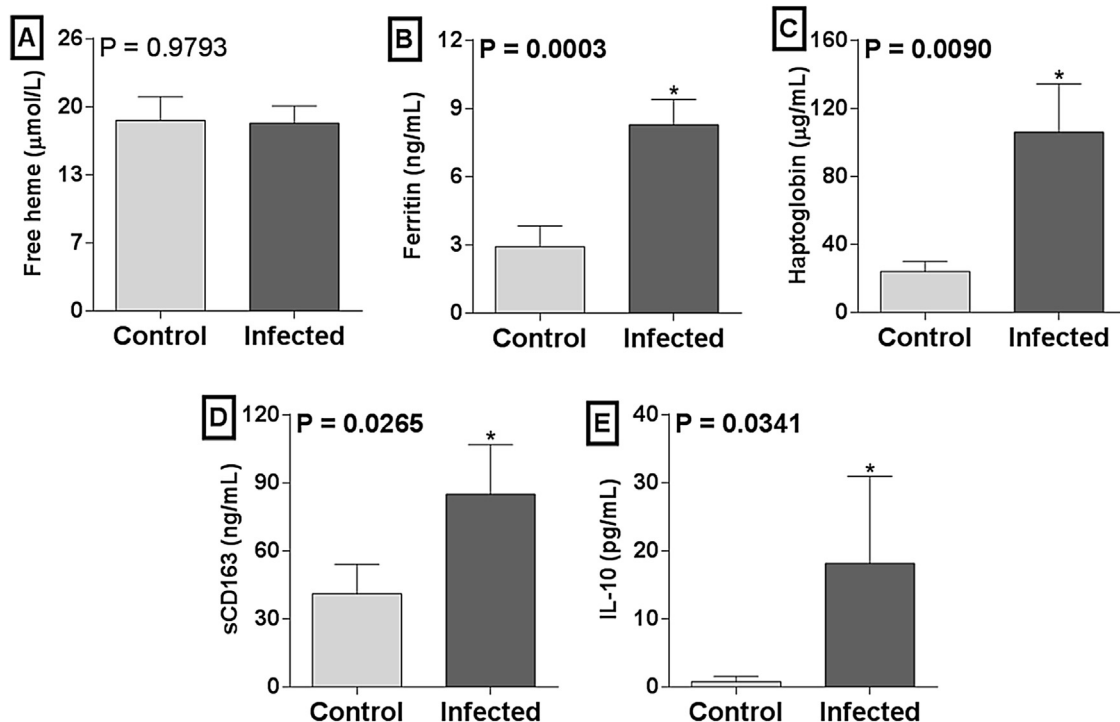


Fig. 3. Plasma levels of free heme (A), ferritin (B), haptoglobin (C), soluble CD163 receptors (sCD163, D) and IL-10 (E) in control dogs (Control, n=24) and dogs with naturally occurring canine visceral leishmaniasis (Infected, n=20). The graphs show the mean and standard error of the mean. Statistically significant differences are indicated by * (Mann-Whitney test).

Table 3

Spearman correlation of heme oxygenase-1 (HO-1) levels and relative expression with oxidative stress markers and IL-10, in controls and in dogs with naturally occurring visceral leishmaniasis, and correlation of HO-1 levels and expression with parasite load in infected dogs.

Parameter	Plasma HO-1 (ELISA)	PBMC HO-1 (ELISA)	BMMC HO-1 (ELISA)	Spleen HO-1 (ELISA)	PBMC HO-1 expression	BMMC HO-1 expression	Spleen HO-1 expression
Albumin	P = 0.0509	P = 0.5137	P = 0.1451	P = 0.0026 r = -0.6361	P = 0.1043	P = 0.0016 r = -0.6571	P = 0.5801
Uric acid	P = 0.9069	P = 0.7984	P = 0.9233	P = 0.0675	P = 0.8888	P = 0.0414 r = 0.4597	P = 0.5264
Total bilirubin	P = 0.5177	P = 0.4543	P = 0.2077	P = 0.0084 r = -0.5724	P = 0.7924	P = 0.0103 r = -0.5596	P = 0.2741
Direct bilirubin	P = 0.3531	P = 0.9399	P = 0.6278	P = 0.1356	P = 0.0334 r = 0.3414	P = 0.0160 r = 0.5308	P = 0.3502
Indirect bilirubin	P = 0.8162	P = 0.2383	P = 0.1884	P = 0.0022 r = -0.6431	P = 0.2619	P = 0.0072 r = -0.5814	P = 0.1609
TAC	P = 0.4631	P = 0.2708	P = 0.8723	P = 0.0099 r = -0.5620	P = 0.8988	P = 0.6195	P = 0.8846
TOC	P = 0.5378	P = 0.4413	P = 0.0020 r = 0.6632	P = 0.3695	P = 0.0852	P = 0.6224	P = 0.2928
TBARS	P = 0.0730	P = 0.4641	P = 0.2809	P = 0.0227 r = 0.5064	P = 0.3083	P = 0.2426	P = 0.0608
Plasma IL-10	P = 0.0275 r = 0.4921	P = 0.1739	P = 0.5126	P = 0.1988	P = 0.4696	P = 0.2529	P = 0.2014
Spleen IL-10	P = 0.0218 r = 0.5093	P = 0.4263	P = 0.8250	P = 0.0001 r = 0.7564	P = 0.4523	P = 0.0405 r = 0.4617	P = 0.3851
Spleen parasite load	P = 0.3923	P = 0.3363	P = 0.3869	P = 0.3129	P = 0.7723	P = 0.7850	P = 0.7330
Bone marrow parasite load	P = 0.6244	P = 0.463	P = 0.8916	P = 0.0806	P = 0.1777	P = 0.6821	P = 0.8382

Abbreviations: BMMC, bone marrow mononuclear cells; HO-1, heme oxygenase-1; PBMC, peripheral blood mononuclear cells; TAC, total antioxidant capacity; TBARS, lipid peroxidation determined via thiobarbituric acid reactive substances; TOC, total oxidant capacity.

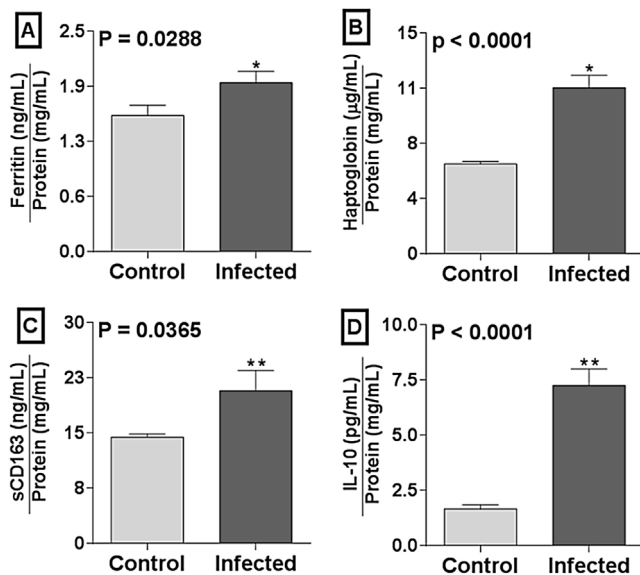


Fig. 4. Spleen homogenate levels of ferritin (A), haptoglobin (B), soluble CD163 receptor (sCD163, C) and IL-10 (D) normalized to protein levels in control dogs (Control, n = 10) and dogs with naturally occurring canine visceral leishmaniasis (Infected, n = 10). The graphs show the mean and standard error of the mean. Statistically significant differences are indicated by * (unpaired *t*-test) or ** (Mann-Whitney test).

proliferative response of LNCs in infected dogs. We observed that HO-1 inhibition using SnMsP significantly increased the LNCs proliferation from infected dogs both at the baseline and in the presence of SAgL, while HO-1 activation using CoPP decreased the antigen-specific lymphoproliferation (Fig. 5A).

HO-1 activation with CoPP and inhibition with SnMsP, respectively, significantly increased and decreased the enzyme levels (Fig. 5B). However, when evaluating gene expression, only activation with CoPP increased HO-1 relative expression significantly,

while inhibition with SnMsP did not alter HO-1 relative expression (Fig. 5C).

3.7. The increase in LNCs proliferation involves decreased production of IL-10 and IL-2 through HO-1 inhibition

After we had determined that HO-1 inhibition improved the antigen-specific lymphoproliferative response in infected dogs and that activation impaired it, we evaluated IL-10 because of its regulatory function in the lymphocyte proliferative response in VL (Strauss-Ayali et al., 2005). We also evaluated IL-2 because of its effect on the activation and proliferation of T lymphocytes (Santana and Rosenstein, 2003) and IFN-gamma because of its production during lymphocyte activation and association with disease resistance in canine infection (Pinelli et al., 1999a,b). We observed that HO-1 inhibition using SnMsP, verified through its reduced levels (Fig. 5B), occurred with decreased production of IL-2 (Fig. 6B) and IL-10 (Fig. 6C) in the LNCs culture supernatant from infected dogs.

The stimulation of LNCs from infected dogs with SAgL increased lymphoproliferation (Fig. 5A) and also decreased IL-2 production (Fig. 6B). Interestingly, both activation and inhibition of HO-1 caused a significant reduction in IL-2 levels, with greater decreases observed in the presence of the SnMsP HO-1 inhibitor (Fig. 6B), which also improved the proliferation rate (Fig. 5A).

The production of IFN-gamma did not consistently change with HO-1 inhibition or activation. IFN-gamma production was reduced in the presence of the HO-1 activator without SAgL, while an increasing trend (not statistically significant) was observed in the presence of SAgL (Fig. 6A).

The ConA mitogen induced the greatest cell proliferation rate (Fig. 5A). This occurred with inhibition of HO-1 relative expression (Fig. 5C) and with higher IFN-gamma, IL-2 and IL-10 production (Fig. 6).

A negative correlation was observed between HO-1 relative expression and the LNCs proliferation rate in the presence of SAgL ($P = 0.0213$, $r = -0.4185$). This indicated that HO-1 relative expression increased when cell proliferation decreased.

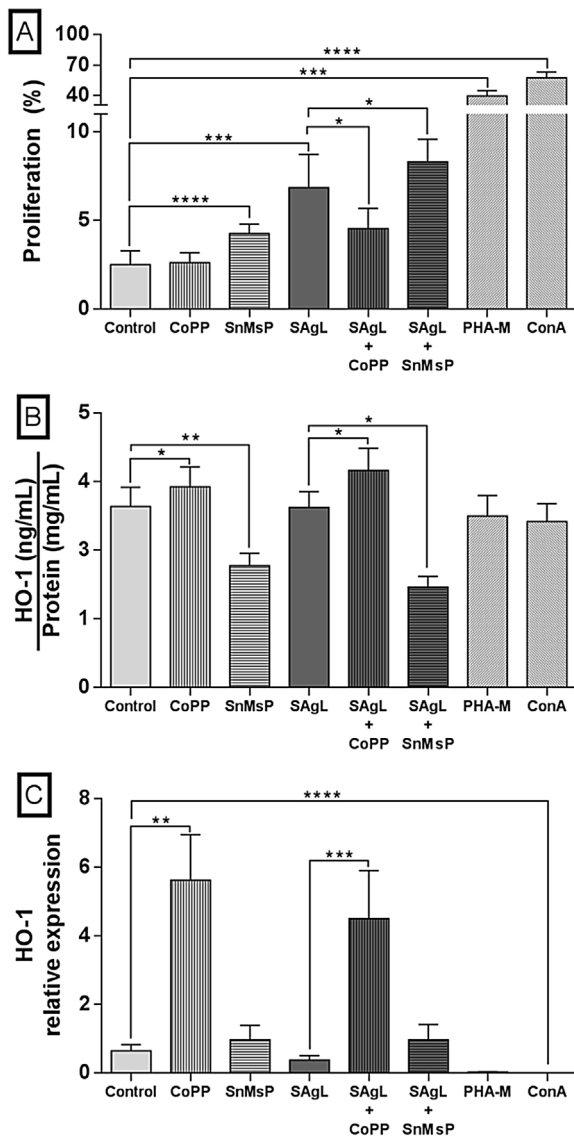


Fig. 5. Proliferation rate of lymph node cells from dogs with naturally occurring visceral leishmaniasis ($n = 15$) stained with CFSE that were cultured for six days without any treatment (control) in the presence of the mitogens phytohemagglutinin-M (PHA-M) and concanavalin-A (ConA), and the soluble antigen of *Leishmania infantum* (SAgL) with or without associated cobaltic protoporphyrin IX chloride (CoPP) and Sn(IV) mesoporphyrin IX dichloride (SnMsP) (A). Heme oxygenase-1 (HO-1) levels measured by means of capture ELISA and normalized to protein levels from lymph cells without CFSE labeling, which had been subjected to the same treatments as described previously (B). HO-1 levels measured by means of capture ELISA and normalized to protein levels from lymph cells without CFSE labeling, which had been subjected to the same treatments as described previously (B). HO-1 relative expression determined by means of qRT-PCR and normalized according to the geometric mean of the standard expression of the genes beta-actin and HPRT-1 from lymph cells without CFSE labeling that had been subjected to the same treatments as described previously (C). The graphs show the mean and standard error of the mean. Statistics: Friedman test with Dunn's multiple comparison. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, **** $P < 0.0001$.

4. Discussion

We demonstrated that dogs with CVL present oxidative stress associated with increased levels and relative gene expression of HO-1. Positive correlations were observed between HO-1 and oxidizing substances, and negative correlations were observed between HO-1 and anti-oxidant substances. Our data also revealed that increases in HO-1 levels correlated with increased IL-10 production. The antigen-specific lymphoproliferative responses from lymph node cells of infected dogs decreased with HO-1 activation and increased with HO-1 inhibition. This increase coincided

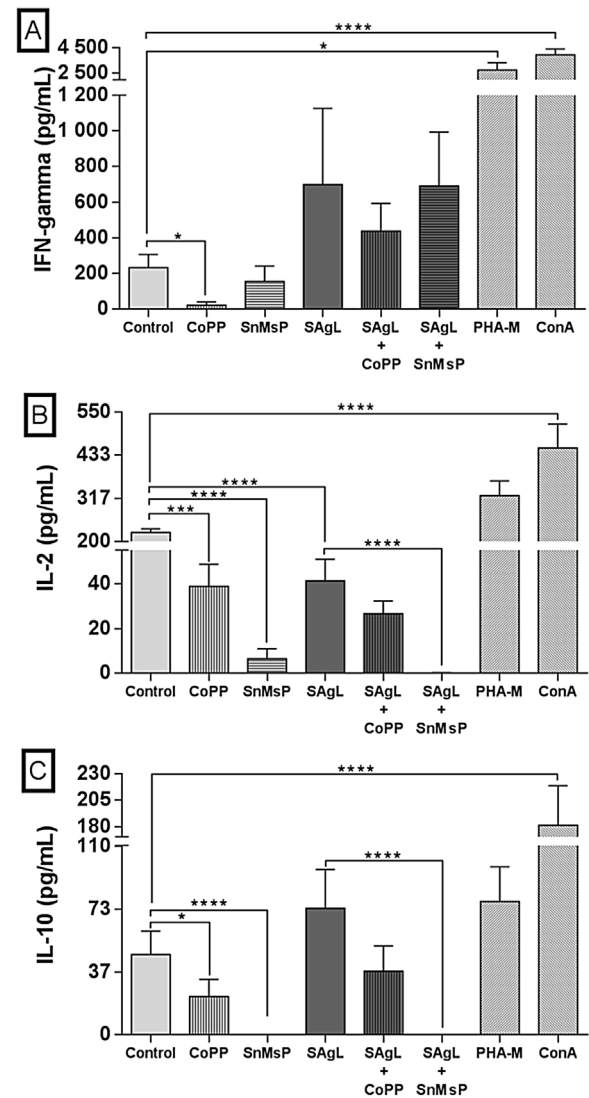


Fig. 6. Interferon-gamma (IFN-gamma, A), IL-2 (B) and IL-10 (C) measured from the supernatant of lymph node cells that were cultured for six days from dogs with naturally occurring visceral leishmaniasis ($n = 15$) without any treatment (control) and in the presence of the mitogens phytohemagglutinin-M (PHA-M) and concanavalin-A (ConA), and the soluble antigen of *Leishmania infantum* (SAgL) with or without associated cobaltic protoporphyrin IX chloride (CoPP) and Sn(IV) mesoporphyrin IX dichloride (SnMsP). The graphs show the mean and standard error of the mean. Statistics: Friedman test with Dunn's multiple comparison. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, **** $P < 0.0001$.

with reductions in IL-10 and IL-2 levels in the cell culture supernatant. These data demonstrate that the increased HO-1 in CVL is likely to be one of the mechanisms responsible for reducing cellular immunity and impaired lymphoproliferative responses in infected dogs.

The oxidative stress observed in CVL was due to reduced antioxidant levels and increased oxidant levels. This resulted in enhanced lipid peroxidation in infected dogs, which is consistent with our previous report (Almeida et al., 2013b). Although the levels of uric acid (an antioxidant) were higher in infected dogs, this was not enough to improve oxidative stress because TAC remained low in these animals.

The positive correlation between HO-1 and oxidants and the negative correlation between HO-1 and antioxidants is the first evidence of the relationship between oxidative stress and increased HO-1 levels and relative gene expression in dogs. However, it has been shown that HO-1 is activated by stress in other species

(Vile et al., 1994). These findings suggest that higher oxidant production and lipid peroxidation reduce the antioxidant levels and increase the HO-1 production by blood and bone marrow monocytes/macrophages in infected dogs. A previous report demonstrated the possibility of development of oxidative stress in humans with VL caused by *L. donovani* infection, which revealed increased HO-1 levels, increased sorbitol dehydrogenase activity and reduced glutathione peroxidase activity. These alterations were reversed after treatment (Das et al., 2013). There was no correlation between HO-1 and the parasite load in the spleens and bone marrow of infected dogs, thus suggesting that oxidative stress has a more important role in HO-1 increases in infected dogs than the number of parasites.

Although HO-1 exhibits antioxidant effects and is increased in CVL, oxidative stress still remained evident in infected dogs, which may have contributed to anemia in these dogs (Britti et al., 2008). Additionally, plasma HO-1 levels correlated negatively with RBC ($r = -0.3595$, $P = 0.0210$), hemoglobin ($r = -0.4728$, $P = 0.0021$) and PCV ($r = -0.4369$, $P = 0.0043$) (data not shown). These correlations point towards an association between increased HO-1 levels and anemia in infected dogs because HO-1 is responsible for metabolizing the heme in hemoglobin (Tenhunen et al., 1969). It is important to highlight that HO-1 is a microsomal enzyme and possibly presents no plasma activity (Tenhunen et al., 1969; Yoshinaga et al., 1982). Moreover, our methodology to determine HO-1 plasma levels using capture ELISA only determined the plasma protein content. However, increased plasma levels of HO-1 are possibly caused by higher synthesis of this enzyme and/or injury in tissues responsible for heme degradation such as the liver or spleen (Tenhunen et al., 1969).

The clinical signs and laboratory abnormalities were similar to those in previous studies (Almeida et al., 2013a,b; Ikeda-Garcia et al., 2008). Additionally, we observed that blood basophil counts were also increased, which had not been reported previously, in other studies examining CVL in infected dogs. We observed increased activity of the muscle injury enzymes AST, CK and LDH in the plasma of infected dogs, compared with uninfected dogs, which might predict muscle injury, as has been shown in previous studies on dogs with VL (Paciello et al., 2009; Vamvakidis et al., 2000).

However, with the exception of CK activity, it is important to emphasize that the other enzymes were within the reference range for dogs. Controls and infected dogs presented CK activity beyond the reference range interval, possibly due to the short reference interval proposed by Kaneko et al. (2008), since other studies have proposed longer intervals for healthy dogs (Lucas et al., 2015; Vamvakidis et al., 2000).

This study was the first to report increased HO-1 levels in the plasma, spleen and PBMC of infected dogs, hence suggesting that the HO-1 enzyme is involved in the pathogenesis of CVL. Studies evaluating HO-1 in VL are still scarce and most involve in vitro studies with macrophages (El Fadili et al., 2008; Luz et al., 2012; Pham et al., 2005). In vivo studies have demonstrated elevated HO-1 serum levels in symptomatic human patients infected with *L. infantum* (Luz et al., 2012) and *L. donovani* (Das et al., 2013) and have shown that these values were reduced after treatment with miltefosine (Das et al., 2013) and meglumine antimoniate (Luz et al., 2012). Similarly to results in humans, increased HO-1 levels in dogs presenting with clinical signs of VL also seem to be related to increased disease susceptibility. However, studies examining HO-1 in resistant asymptomatic dogs are required to support this hypothesis.

HO-1 relative expression was only increased in PBMC and BMCM in dogs with VL. There were no differences in spleen tissue. A similar study in symptomatic human patients infected with *L. donovani* showed increased HO-1 gene expression in PBMC and in the splenic aspirate of those patients, with no alterations in bone marrow

aspirate (Das et al., 2013). The discrepancies between these results and ours may be attributable to differences in the hosts, *Leishmania* spp. strains and disease stages.

In addition to the HO-1 levels and relative expression, we evaluated part of the HO-1 metabolic pathway in the plasma and spleen. We observed increased levels of all substances, except for plasma-free heme. When hemoglobin is released from erythrocytes, it forms a stable complex with haptoglobin, which is recognized and internalized by splenic macrophages through the CD163 receptor of these cells (Tenhunen et al., 1969). Free iron, in turn, induces synthesis of ferritin, an important antioxidant that is responsible for transporting this substance (Balla et al., 1992). The increased levels of these metabolites in the plasma and spleens of infected dogs reinforce the increased hemoglobin metabolism in infected animals, as demonstrated by the negative correlations between HO-1 and RBC, hemoglobin and PCV.

The unaltered free heme levels in infected dogs may be attributable to differences in hemoglobin metabolism. The heme is possibly degraded directly inside macrophages (extravascular hemolysis) in CVL, thereby avoiding formation of free heme. Increases in the levels of this metabolite have been described in intravascular hemolytic diseases, such as malaria (Andrade et al., 2010; Sinha et al., 2008). Both ferritin and haptoglobin are acute-phase inflammatory proteins (Ceron et al., 2005) and increased serum levels of these substances have already been demonstrated in dogs with clinical VL (Martinez-Subiela et al., 2014; Martínez-Subiela et al., 2002; Martinez-Subiela and Ceron, 2005). This indicates that the increase in these protein levels could also be a consequence of the inflammatory process.

Similarly to our results, increased levels of sCD163 have been demonstrated in the serum of human patients with LV (Schaer et al., 2005), with no previous reports in CVL. This receptor is restricted to cells of the macrophage lineage and demonstrates macrophage activation caused by infection (Law et al., 1993), which could also contribute to greater ferritin production (Ravelli, 2002). The role of the CD163 receptor in macrophage activation has not been fully elucidated. This receptor may exert pro-inflammatory activity, through inducing secretion of proinflammatory cytokines (Van den Heuvel et al., 1999), or it may exert anti-inflammatory activity, in which cytokines such as IL-10 can induce CD163 expression (Sulahian et al., 2000). These results support our findings, given that we observed increased concentrations of IL-10 and sCD163 in infected dogs. Thus, the inflammation and oxidative stress that occur in CVL appear to contribute to the increases in HO-1 levels and relative gene expression, as well as to increased macrophage activation and hemoglobin metabolism, thereby leading to increased concentrations of haptoglobin, ferritin and sCD163.

Since we observed increased HO-1 metabolism in susceptible dogs with VL, we assessed whether inhibition of HO-1 could improve the proliferative response of lymph node cells from infected dogs. Additionally, we identified the cytokines involved in this process, given that it has been determined that infected dogs have impaired cellular immunity (Baneth et al., 2008). We observed reductions in the antigen-specific LNCs proliferative response through activation of HO-1 and increased proliferation of these cells with reduced HO-1 levels. Negative correlations between HO-1 relative expression and the antigen-specific lymphoproliferation rate were also observed.

This is the first evidence that HO-1 inhibition with SnMsP improves the antigen-specific proliferative response in lymph node cells of infected dogs. Previous studies in humans have shown that SnMsP increases unspecific proliferation of T CD3+ cells (Bunse et al., 2015; Burt et al., 2010) through inducing expression of costimulatory molecules on monocytes, such as CD86 (B7-2), and optimizing the antigen presentation process (Burt et al., 2010). In CVL, it is already known that the proliferative response of

T lymphocytes is reduced due to suppression of B7 molecules in the macrophages of infected dogs (Pinelli et al., 1999a,b). Thus, one of the possible mechanisms activated through HO-1 inhibition could involve restoration of these molecules, thereby optimizing cell activation and proliferation.

IL-10, an important anti-inflammatory interleukin, was increased in the plasma and spleens of infected dogs. This increase correlated positively with HO-1 levels and relative gene expression, and its synthesis was suppressed by HO-1 inhibition in LNCs cultures from infected dogs. These findings suggest that there is a relationship between HO-1 and IL-10 levels, in which increased levels of this cytokine appear to stimulate HO-1 synthesis, or vice versa, thus contributing to the impaired lymphoproliferative response of sick dogs. This positive correlation between HO-1 and IL-10 has been demonstrated in humans with VL (Luz et al., 2012) and is consistent with other studies that have claimed that IL-10 induces HO-1 gene expression in human monocytes that are stimulated with lipopolysaccharide (Petit-Bertron et al., 2003) and in murine macrophages (Lee and Chau, 2002). Furthermore, activation of HO-1 activity causes increased production of carbon monoxide (CO), which also increases the levels of anti-inflammatory cytokines such as IL-10 (Brouard et al., 2000), thus confirming our results from infected dogs and cultured LNCs from these animals.

We observed that HO-1 inhibition with SnMsP in the presence of *L. infantum* antigen significantly reduced IL-2 levels and increased LNCs proliferation. Although it has been accepted that increased IL-2 is required for full activation and proliferation of CD4+ T lymphocytes (Santana and Rosenstein, 2003), other studies have demonstrated that this cytokine has an immunoregulatory role (Bachmann and Oxenius, 2007). High IL-2 levels would be crucial for activation of Treg lymphocytes (Brunkow et al., 2001; Fontenot et al., 2003) and could induce cell death in lymphocytes via FAS/FAS-L after repeated stimulation (Krammer, 2000). However further studies to determine IL-2 function in CVL antigen-specific proliferation are needed.

Use of SnMsP was effective in improving the proliferative response of LNCs, but did not inhibit HO-1 relative expression. Other studies have shown increasing HO-1 relative expression due to SnMsP (Burt et al., 2010) and due to other inhibitors, such as zinc protoporphyrin IX (El Fadili et al., 2008). This suggests that the decreased HO-1 levels resulted from inhibition at a post-transcriptional level.

Given that dogs with CVL exhibit increased HO-1 metabolic pathway activity and that the HO-1 enzyme has immunoregulatory effects on cell proliferation (Listopad et al., 2007), we demonstrated that HO-1 inhibition improves cellular immunity in infected animals. Thus, new therapeutic approaches for CVL should consider using pharmacological HO-1 inhibitors along with antioxidants, to reduce oxidative stress under these conditions. Evaluation of whether oxidative stress is the only factor that is responsible for HO-1 activation in infected dogs should be considered in other studies, since HO-1 is known to be activated by stress conditions.

5. Conclusion

Increased HO-1 metabolism is associated with oxidative stress and IL-10 and appears to be one of the mechanisms involved in inhibition of cellular immunity in dogs with visceral leishmaniasis, thereby impairing the lymphoproliferative response of sick dogs.

Conflict of interest

The authors declare that there were no financial or commercial conflicts of interest.

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