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B cell profile, B-cell activating factor concentration and IgG levels in human cutaneous and mucosal leishmaniasis

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B-cell alterations in human leishmaniasis

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Data availability: The data that supports the findings of this study are available from the corresponding author upon reasonable request.

Author contributions: C.P. designed and performed the experiments and wrote the manuscript; A.B. obtained samples of lesions, made the diagnosis of leishmaniasis and performed the identification of *Leishmania* species; M.F.G.B. obtained samples of lesions and conducted clinical follow-up of patients; A.G.G.P. performed cultures of parasites and identification of *Leishmania* species; J.P., N.B., M.C.A. and M.E.C.E. performed ELISA and CBA analysis; S.A.L. and M.M.E.B. participated in the design, discussion and edition of the final version of the manuscript.

Abstract

Aims: The aim of this study was to evaluate characteristics of B cells in human tegumentary leishmaniasis (TL) analyzing cutaneous leishmaniasis (CL), most prevalent form and mucosal leishmaniasis (ML), aggressive form characterized by the destruction of the oral-nasal-pharyngeal cavities.

Methods and results: By flow cytometry analysis we found decreased percentages of nonclass-switched memory B cells in TL with the degree of the loss related to clinical severity. Using commercial ELISA, we reported high levels of B-cell activating factor (BAFF) and IgG preferentially in aggressive CL and markedly in ML together with decreased BAFF receptors in the latter. We also found lower levels of BAFF after clinical recovery suggesting a relation between BAFF and disease activity.

ML history of therapeutic failure presented high levels of BAFF accompanied by detectable concentrations of IFN- γ and IL-6 (assayed by commercial ELISA and cytometric bead arrays respectively), cytokines involved in exaggerated inflammatory responses and tissue damage in TL.

Conclusion: We demonstrate B cell disturbances in TL with the degree of the alterations related to clinical severity. We suggest a relation between excess of BAFF and disease activity and point towards a possible implication of BAFF in the inflammatory phenomenon of ML.

Keywords

B-cell activating factor; Cutaneous leishmaniasis; Mucocutaneous leishmaniasis; B lymphocytes; Hypergammaglobulinemia; Interferon-gamma; Interleukin-6

1. Introduction

B cells have the ability to produce antibodies but are also involved in antigen presentation, cytokine production and activation of T cells ¹. Most reports analyzing tegumentary *Leishmania* strains suggest dual roles for B cells, contributing to disease susceptibility or enhancing protection ^{2, 3}, depending on the experimental model and the parasite strain used. Although trypanosomatids are able to hinder and skew B cell functions to favor their own survival ¹, the role of the humoral response in human tegumentary leishmaniasis (TL) is currently still scant. The peripheral memory B cells can be characterized by their surface expression of IgD and CD27 ⁴. B cell responses are also guided by B cell activating factor (BAFF), crucial regulator of peripheral B cell homeostasis ^{5, 6}, whose overproduction leads to detrimental effects in autoimmune disorders ^{7, 8}. Less is known about BAFF in microbial diseases, with elevated levels associated with disease progression, expansion of atypical memory B cells and inefficient antibody response in plasmodium and HIV infections ^{9, 10, 11, 12}. Increased BAFF has also been reported in visceral leishmaniasis ¹³ but its participation in TL remains unknown.

To describe the characteristics of B cells in TL, we analyzed plasma BAFF and total IgG, BAFF receptors, B cell phenotype and the presence of autoantibodies in cutaneous leishmaniasis (CL), the most prevalent form, in comparison to mucosal leishmaniasis (ML), chronic and aggressive form characterized by the destruction of the oral-nasal-pharyngeal cavities ^{14, 15}. Our aim was to determine if B cell disturbances are frequent in TL and if the degree of the alterations is related to distinct clinical forms.

2. Material and methods

2.1. Study groups

The study population comprised patients from the Northwest of Argentina diagnosed by clinical evaluation, identification of amastigotes, culture of parasites from lesion samples and detection of *Leishmania* DNA by using k-DNA-based PCR¹⁶.

Table 1 shows the clinical characteristics of the patients and the control group (HS). The analytical approaches were performed irrespectively of the condition for anti- *T. cruzi* antibodies after verifying that there were not significant differences between CL vs. *T. cruzi*-seropositive CL and ML vs. *T. cruzi*-seropositive ML groups.

2.2. *Leishmania* spp. identification

DNA extraction (from samples obtained from lesions and culture-isolated parasites) and polymorphism Specific-PCR were performed as previously described¹⁷.

2.4. Blood sampling procedure

Plasma sample separation and isolation of peripheral blood mononuclear cells were performed as previously described¹⁸.

2.5. Measurement of anti-*T. cruzi* antibodies

Testing for anti-*T. cruzi* antibodies was carried out using a commercial kit (recombinant ELISA v.3.0, Wiener, Argentina).

2.6. B cell phenotype

FITC, PE or PerCP labeled anti-CD19, anti-CD27, anti-IgD, anti-BAFF-R, and anti-TACI monoclonal antibodies (Becton Dickinson, San José, CA, USA) were used. The cells were evaluated using a FACScan flow cytometer and FCS 3 Express, De Novo Software (Los Angeles, CA, USA).

2.7. ELISA and cytometric bead arrays (CBA)

The plasma concentrations of BAFF (Adipogen, San Diego, USA), total IgG (Bethyl, Texas, USA), IgG dsDNA antibodies (IMTEC, Wiesbaden, Germany) and IFN- γ (BD Biosciences, San Diego, USA) were determined by commercial ELISA. IL-6 was measured by CBA (BD

Biosciences, San Diego, USA) using a FACScalibur flow cytometer and the BD CBA Software, version 1.4.

2.8. Statistical analysis

A level of $p < 0.05$ was accepted as being statistically significant. The normality of the variable distribution was assessed by the D'Agostino & Pearson omnibus test. Continuous variables between two groups were compared with the Mann-Whitney U-test. A nonparametric Kruskal-Wallis test with Dunn's post-test was used to compare differences among three or more groups. The data were analyzed with GraphPad Prism 6 (GraphPad Software, San Diego, CA).

3. Results

3.1. Identification of the causative agent of CL and ML

Leishmania spp. could be identified in 23 CL and in 11 ML samples (Table 1). In the rest of the samples, we obtained negative PS-PCR results after DNA extraction due to a low number of parasites in the chronic lesions. *L. (V.) braziliensis* was found in the majority of the CL and ML cases, followed by *L. (L.) amazonensis*.

3.2. Peripheral B cell subsets in TL

Gating on CD19⁺ B cells, we analyzed naïve (IgD⁺CD27⁻); nonclass-switched (IgD⁺CD27⁺), class-switched (IgD⁻CD27⁺) and double-negative (IgD⁻CD27⁻) B cells. We compared CL with single lesions (CL-S); CL with multiple lesions (CL-M) and ML. Both CL groups and more markedly ML patients showed significantly lower percentages of nonclass-switched memory B cells in comparison with the HS group (Fig. 1B). The gating strategy is shown in Figure 2.

3.3. Plasma concentrations of BAFF and total IgG in active TL

We next evaluated crucial factors and found elevated levels of BAFF and total IgG in CL-M and in ML (Fig. 3A and 3B) compared with the HS group. When patients were split into those with active lesions and those who showed complete clinical recovery (Table 1) BAFF levels were significantly lower in the groups with clinical improvement in contrast to CL-M and more markedly ML (Fig. 3A).

3.4. BAFF levels and proinflammatory cytokines in ML

We assessed proinflammatory cytokines in ML because IFN- γ contributes to tissue damage and IL-6 to active disease in aggressive forms of leishmaniasis ¹⁹. We compared the concentration of BAFF in 12 ML patients with detectable IFN- γ [39.75 (20-112.8) pg/ ml; median (range)] and in 8 ML patients with undetectable IFN- γ (Figure 4A). Besides, we compared the levels of BAFF in 11 ML patients with IL-6 median concentration of 5.32 (1.25–33.91) pg/ ml and in 5 ML patients with absence of IL-6 (Figure 4B). We found higher levels of BAFF within the groups with detectable cytokines. Although aggressive nasopharyngeal lesions characterize the entire ML cohort, the majority of the ML patients with history of therapeutic failure (Table 1) showed simultaneous detection of BAFF, IFN- γ and IL-6 (Figure 4).

3.5. Expression of the BAFF receptors in patients with ML

We measured the expression of BAFF-receptor (BAFF-R) and TACI on CD27⁺CD19⁺ memory B cells from the CL-S, CL-M and ML groups.

Both receptors were diminished only in ML compared with the HS group (Fig. 5A and 5B).

3.6. Detection of anti-double-stranded DNA antibodies (dsDNA antibodies) in TL

To evaluate the production of autoantibodies, we assessed dsDNA antibodies, markers of Systemic lupus erythematosus (SLE) also detected in infections as chronic hepatitis C ²⁰. We assayed 22 CL and 22 ML samples, with the absence of these antibodies in all but one of the tested samples (Table 2).

4. Discussion

In the present work we found that *L. (V.) braziliensis* is the predominant causative agent of CL and ML in the Northwest of Argentina according to previous data from the group ²¹.

We report that nonclass-switched memory B cells, essential in secondary immune responses ⁴, are decreased in CL and markedly diminished in ML, suggesting a relation between the degree of the loss and clinical severity. This subset was also diminished in malaria ²², rheumatoid arthritis ²³ and HIV infections ²⁴.

. We then analyzed BAFF, a member of the trimeric TNF family crucial for B cell function ²⁵. We report that in TL high levels of BAFF and of total IgG are more frequently found in aggressive clinical forms, named CL with multiple lesions and ML. BAFF demonstrated an impact over the nonclass-switched memory B subset ²⁶, so it could be affecting the memory B cell composition in TL. In addition, only ML showed diminished BAFF receptors on memory B cells, in agreement with autoimmune diseases whereas the lower expression of BAFF receptors was associated with more severe disease ^{27, 28}. In contrast, recovered CL and ML patients showed lower BAFF levels than their active disease counterparts. These observations suggest that in TL an excess of BAFF might be related to disease activity, as previously reported for autoimmune diseases ²⁷ and malaria ²⁹. The induction of BAFF could be favored by antigen persistence, as other protozoan antigens have shown the ability to induce this molecule in experimental conditions ³⁰. An exaggerated inflammatory response directed by IFN- γ is mainly responsible for the tissue damage in TL ^{31,32} and although less is known about IL-6 ³³, increased levels were found in active visceral leishmaniasis ¹⁹. We report that in the majority of ML cases with history of therapeutic failure BAFF was accompanied by the detection of IFN- γ and IL-6. These findings suggest that BAFF could contribute to the inflammatory phenomenon produced in ML.

The absence of antibodies to DNA used as unique markers of autoimmunity ³⁴ in almost all of the studied patients suggest that in TL, the activation of B cells might not necessarily be linked to autoantibodies production, although assessment of a wider set of antibodies is needed to validate this idea.

Our work demonstrates that the degree of the B cell alterations is related to clinical severity in TL involving changes in peripheral B cell subsets, BAFF and IgG levels and expression of BAFF receptors. We suggest a relation between BAFF and disease activity in TL and indicate a possible implication of this molecule in the inflammatory phenomenon of ML.

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Ethical Approval: All procedures performed in this study involving human participants were in accordance with the ethical standards of the institutional research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards. Informed consent was obtained from all individuals participants included in the study.

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Figure legends

Fig. 1. Subtypes of peripheral memory B cells in cutaneous leishmaniasis (CL) and mucosal leishmaniasis (ML). The CD19⁺ B cell population was gated and analyzed for CD27 versus IgD in (A) naïve B cells, IgD⁺CD27⁻; (B) nonclass-switched memory B cells, IgD⁺CD27⁺; (C) class-switched memory B cells, IgD⁻CD27⁺; and (D) double-negative B cells, IgD⁻CD27⁻. Four clinical groups were compared for each set of markers: healthy subjects, HS, white circles (n = 15); CL with single lesions, CL-S (n = 13) and CL with multiple lesions, CL-M (n = 6), light grey circles; ML, dark grey circles (n = 19). The CL and ML cases with detectable antibodies for *T. cruzi* are represented by half black circles. **p* < 0.05, ***p* < 0.01, *****p* < 0.0001 using the Kruskal-Wallis test and Dunn's post-test. Scatter dot plots, line and error bars show the median and the range.

Fig. 2. Gating strategy for the analysis of the CD19⁺ B cell peripheral population. (A) Lymphocytes were gated by forward and side light scatter, and subsequently (B) the CD19⁺ B cells were selected. (C) Representative dot plots from a healthy subject (HS), a cutaneous leishmaniasis (CL) patient and a mucosal leishmaniasis patient (ML) are shown. The graphs show IgD and CD27 staining of the gated CD19⁺ B cells.

Fig. 3. Plasma concentrations of B-cell activating factor (BAFF) and total IgG in cutaneous (CL) and mucosal (ML) leishmaniasis. The amount of soluble factors was compared in six clinical groups: healthy subjects, HS, white circles (n = 13-24); CL with single lesions, CL-S (n = 19-22) and CL with multiple lesions, CL-M (n = 8-11), light grey circles; patients recovered from CL (R-CL, light grey squares, n = 7); ML, dark grey circles (n = 28-32) and patients recovered from ML (R-ML, dark grey squares, n = 7). CL and ML patients with detectable antibodies for *T. cruzi* are represented by half black circles or squares. (A) Plasma BAFF concentration; (B) total IgG concentration. Comparison among the HS, CL-S, CL-M and ML groups by Kruskal-Wallis test and Dunn's post-test, ***p* < 0.01, ****p* < 0.001. Comparison of CL-M vs. R-CL and ML vs. R-ML by Mann Whitney U-test, **p* = 0.0022 and ****p* = 0.0003, respectively. Scatter dot plots, line and error bars show the median and range.

Fig. 4. Plasma levels of B-cell activating factor (BAFF) and proinflammatory cytokines in mucosal leishmaniasis (ML). (A) BAFF levels in ML patients with detectable concentration of IFN- γ (ML IFN γ POS, n = 12) compared with ML patients with undetectable levels of IFN- γ (ML

IFN γ NEG, n = 8), Unpaired *t* test, **p* < 0.05; (B) BAFF levels in ML patients with detectable concentration of IL-6 (ML IL6 POS, n = 11) compared with ML patients with undetectable levels of IL-6 (ML IL6 NEG, n = 5), Mann Whitney U-test, **p* < 0,05. ML patients are represented by dark grey circles; ML patients with history of therapeutic failure are represented by light grey squares.

Fig. 5. Expression of the BAFF receptors on CD27⁺CD19⁺ B lymphocytes from patients with cutaneous (CL) and mucosal (ML) leishmaniasis. Four clinical groups were compared: healthy subjects, HS, white circles (n = 9-10); CL with single lesions, CL-S (n = 6-9) and CL with multiple lesions (n = 3-4) light grey circles; ML, dark grey circles (n = 8-12). CL and ML patients with detectable antibodies for *T. cruzi* are represented by half black circles. (A) Expression of BAFF-receptor (BAFF-R); (B) Expression of transmembrane activator and calcium-modulator and cyclophilin ligand interactor (TACI). **p* < 0.05; ***p* < 0.01 using Kruskal-Wallis test and Dunn's post-test. Scatter dot plots, line and error bars show the median and range.

Table 1. Clinical features of patients with cutaneous and mucosal leishmaniasis

Clinical form	No. patients	Age (years)*	Male gender	<i>L. (V.) braziliensis</i>	<i>L. (L.) amazonensis</i>	Anti- <i>T. cruzi</i> antibodies	Lesion age*	History of therapeutic failure	Time of complete recovery*
Cutaneous leishmaniasis									
Single active lesion	25	30 (13-79)	14/25 (60 %)	13/15 (87 %)	2/15 (13 %)	5/25 (20 %)	3 (0.5–48) months	4/25 (16 %)	NA
Multiple active lesions	12	38 (19-80)	8/12 (67 %)	7/8 (87.5 %)	1/8 (12.5 %)	3/12 (25 %)	5 (2–36) months	2/12 (17 %)	NA
Clinical recovery	7	34 (15-40)	5/7 (70 %)	NA	NA	1/7 (14 %)	NA	NA	2 (2-36) months
Mucosal leishmaniasis									
Mucosal active lesions	40	48 (13-75)	32/40 (80 %)	9/11 (82 %)	2/11 (18 %)	17/40 (42.5 %)	3 (0.17-30) years	19/40 (48%)	NA
Clinical recovery	7	39 (13-75)	5/7 (71 %)	NA	NA	4/7 (57 %)	NA	NA	10 (10-22) months
Control group									
Healthy subjects	25	34 (22-68)	16/25 (64 %)	NA	NA	0/25 (0 %)	NA	NA	NA

Note: NA, not applicable; *Median (Min-Max)

Table 2. Assessment of double-stranded DNA (dsDNA) antibodies in tegumentary leishmaniasis

Patient groups	dsDNA antibodies	
	No. of subjects with negative findings (< 25 WHO-IU/ml)	No. of subjects with positive findings (> 40 WHO-IU/ml)
CL	17	0
CL with detectable antibodies for <i>T. cruzi</i>	5	0
ML	13	1
ML with detectable antibodies for <i>T. cruzi</i>	9	0
Total	44	1

Note: CL, cutaneous leishmaniasis; ML, mucosal leishmaniasis









