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Review/Fórum

Immunomodulating effects of Phlebotomine saliva in *Leishmania* infection: Review

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Abstract. The recognized action of some proteins present in the saliva of the phlebotomine dipteran vectors of Leishmaniasis, as modulators of the hemostatic, inflammatory and immune responses during the colonization of the protozoan in the host organism, created the expectation of formulating a vaccine against this disease. The present study aimed to provide an overview about researchs that deals with the role of phlebotomine saliva in *Leishmania* infections. A bibliographic search was carried out in PubMed, SciELO and LILACS databases. The articles listed address the following aspects of the vector's saliva: intensification of infection; suppression and modulation of the host immune system; chemotaxis on phagocytic cells; induction of innate and adaptive immunity; isolation and characterization of immunoactive molecules and their possible protective function against Leishmaniasis.

Keywords: Lutzomyia; Phlebotomus; protection; sandflies, saliva.

Efeitos imunomoduladores da saliva de flebotomínio na infecção por Leishmania: Revisão

Resumo. A reconhecida ação de algumas proteínas presentes na saliva dos vetores flebotomíneos da Leishmaniose, como moduladores das respostas hemostáticas, inflamatórias e imunes durante a colonização do protozoário no organismo hospedeiro, criou a expectativa de formular uma vacina contra essa doença. O presente estudo teve como objetivo fornecer uma visão geral sobre pesquisas que abordam o papel da saliva de flebotomíneos em infecções por *Leishmania*. Foi realizada uma pesquisa bibliográfica nas bases de dados PubMed, SciELO e LILACS. Os artigos listados abordam os seguintes aspectos da saliva do vetor: intensificação da infecção; supressão e modulação do sistema imune do hospedeiro; quimiotaxia em células fagociticas; indução da imunidade inata e adaptativa; isolamento e caracterização de moléculas imunoativas e sua possível função protetora contra a leishmaniose.

Palavras-chave: Flebótomos; Lutzomyia; Phlebotomus; proteção; saliva.

eishmaniasis, an infection caused by protozoa of the genus *Leishmania*, family Trypanosomatidae and order Kinetoplastida is a zoonotic disease with several clinical manifestations that depend on both the parasite species and the host immune response (VAN DER AUWERA & DUJARDIN 2015). This disease is transmitted by dipterans of the subfamily Phlebotominae of the genera *Lutzomyia* and *Phlebotomus*, found in the New and Old World, respectively (Yong & DUNCAN 1994; GAZZINELLI & DENKERS 2006).

There are some limiting aspects in the treatment of Leishmaniasis, such as its high cost, the difficulty of administration of the drugs (which drugs are usually administrated by endovenous or intramuscularly), long treatment period and drug toxicity. There is also a lack of response to antimonials, the first choice drug for treatment, characterizing resistant forms of the disease (DE PAULA *et al.* 2003).

The difficulties in controlling the disease have boosted studies in the last decades in an effort to uncover mechanisms of infection of Leishmaniasis in order to develop immunization methods and efficient treatments. In this sense, an important role of

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the phlebotomine vector's saliva was observed in order for *Leishmania* to be successfully established in the host organism

(TITUS et al. 1988). Studies indicate that some proteins present

in the vector's saliva may induce a protective immunological

profile in the host when it is challenged. This immunomodulatory

process could be capable of hindering the initial infection

of Leishmania promastigotes in the target cell, creating the

expectation of formulating a vaccine against Leishmaniasis

(ENSERINK 2001; OLIVEIRA *et al.* 2015). In this sense, the objective of this manuscript is to discuss representative studies on the

biological properties of sandfly salivary molecules in Leishmania

GENERAL ASPECTS OF THE IMMUNOLOGICAL

RESPONSE UPON LEISHMANIA INFECTION

From the inoculation of the promastigote forms and saliva during the blood meal, a complex interaction between the parasite and

the host's immune response begins, where they are internalized

giving rise to phagosomes which fuses to the lysosomes forming a

cellular compartment rich in proteolytic enzymes with digestive

and microbicidal functions (RITTIG & BOGDAN 2000; SEMINI

infection, highlighting their biotechnological interest.

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et al. 2017). The vector's saliva induces cellular recruitment, influences the modulation of the T lymphocyte population, and inhibits the complement system and phagocytic cells, such as dendritic cells and neutrophils, as well as the appearance of late hypersensitivity reaction to saliva antigens and presence of IFN- γ at the site of the sting (ANDRADE *et al.* 2005; VINHAS *et al.* 2007).

Initially, the presence of inoculated sentinel cells elicits a nonspecific acute inflammatory response in which they participate in pathogen-associated molecular patterns (PAMPs) and Toll-Like Receptors (TLR) rapidly leading to the activation of transcription factors in the hyaloplasm (GAZZINELLI & DENKERS 2006). Once activated, the TLR signal through the formation of a TLR2-PI3K-phospholipase C complex (PLC) which induces activation of cytoplasmic phospholipase A_2 and Cyclooxygenase-2, which are essential for the formation of prostaglandins (PGE2) (BHATTACHRJEE *et al.* 2016).

In this context, the analyze of eicosanoids, in C57BL/6 mice measured ex vivo from leukocytes harvested by peritoneal cavity washing, induced by *Lutzomyia longipalpis* (Lutz & Neiva) saliva, main vector of American visceral leishmaniasis (AVL), suggests a regulation in the early stages of *Leishmania*. *infantum* infection leading to implications in the interaction between PGE2 and Leukotrienes B4 (LTB4) on immunopathogenesis (ARAÚJO *et al.* 2014). These mediators involved in different mechanisms, where PGE2 suppresses the host immune response through inhibition in the production of proinflammatory cytokines, for example: IFN- γ , TNF- α , IL-2 and IL-1 β (HARRIS *et al.* 2002). While LTB4 plays a role in increasing proinflammatory cytokines, as well as a decreasing the infection in several pathologies, since nitric oxide production may increase (PETERS-GOLDEN *et al.* 2005; ROGERIO & ANIBAL 2012).

In addition to these inflammatory inducers, the components of the complement system represent the first line of immune defense against Leishmania. The initial events of complement activation (fragments C3a, C4a and C5a) act as potent chemotactic stimuli for neutrophils, in addition to activating histamine-producing (vasodilator) mast cells, cytokines and other proinflammatory substances (Delves & Roitt 2000). Despite these obstacles, it is known that metacyclic promastigote forms express a large amount of kinases that phosphorylate several proteins of the host's complement system (C3, C5 and C9), which can inactivate it, thus increasing its survival inside macrophages by promoting low oxidative metabolism (HERMOSO et al. 1991). Recent studies have identified a protein present in sandfly saliva called SALO which specifically inhibited the classical pathway of the human complement system by a mechanism that appears to act on C1q through the displacement of serine proteases from the C1 complex without inhibiting the binding of C1q to IgG (FERREIRA et al. 2016). This synergistic action of proteins from the promastigote forms and saliva components may protect the parasite from mediated lysis during the initial stage of infection. Another study found that increased serine peptidase (ISP2) inhibitor in Le. donovani infected macrophages, etiologic agent of visceral leishmaniasis of the Old World, plays a critical role in the reduction of MASP2 via lectin, decreasing the formation of C3 and C5c and the membrane attack complex (VERMA et al. 2018).

In this environment, the development of a cellular response is of great importance in the control of Leishmaniasis, macrophages, Langerhans cells, Th1-type CD4+ cells and the cytokines they release like IFN- γ , IL-12 and TNF- α , are crucial in long-term resolution and protection against disease (Gomes *et al.* 2012). Once the Th2 lymphocyte response occurs, which is characterized by the intense release of IL-4, IL-5 and IL-10 cytokines, the host's initial inflammatory response is modified to a phenotype that is permissive to the infection. Connected to this phenomenon, higher levels of TGF- β and prostaglandin E_2 cytokines are able to inhibit IFN- γ production and deactivate macrophages (MONTEIRO *et al.* 2007).

Even with a Th1 effector host immune response, *Leishmania* is able to silence mechanisms of NO (nitric oxide) and reactive oxygen species, such as hydrogen peroxide, synthesis, which are potent microbicides produced by macrophages (GREEN & NACY 1993). For example, inhibition by membrane lipophosphoglycans (LPG) of promastigote forms acting on inducers of NO synthesis, such as PKC (protein kinase C), leads to a decrease in these potent effector molecules against the parasites (DESCOTEAUX & TURCO 1993). This delay in NO production can also occur through mechanisms provoked by synthesized TGF- β mainly CD4+ CD25+ T cells in individuals infected with *Leishmania* sp. leading to deleterious effects by increasing vulnerability (KARIMINIA *et al.* 2005).

Other mechanisms related to the evasion of *Leishmania* have been described; among these, the immunomodulatory effects due to salivary gland lysate (SGL) of *L. longipalpis* where an increase in IL-10 was observed stands out; similar to the SGL of *Phlebotomus papatasi* Scopoli, vector of *Leishmania major* L. the etiologic agent of zoonotic cutaneous leishmaniasis (ZCL), where IL-4 levels increased and IFN- γ , IL-12 and nitric oxide synthetase (iNOS) decreased with consequent worsening of saliva-induced skin lesions (NORSWORTHY *et al.* 2004).

THE ROLE OF SOME SALIVARY COMPONENTS IN BLOOD-FEEDING AND HOST IMMUNITY

Among the salivary properties of phlebotomines, the following stand out: the anticoagulant, vasodilator and antiplatelet factor, favoring the flow of blood and interstitial lymph for their feeding (ABDELADHIM *et al.* 2014). In addition to factors inherent to the host and the parasite, it has been demonstrated that some components of the insect's saliva also interfere in the host's immune response, favoring the persistence of the parasite in the macrophage, acting as one of the determinants of infection evolution (OLIVEIRA *et al.* 2013).

The salive of *L. longipalpis*'s saliva contains a potent vasodilator, maxadilan. This molecule is responsible for erythema caused by the bite during hematophagy inducing leukocyte chemotaxis and plasma extravasation in post-capillary venules through a mechanism associated with the type I polypeptide receptor of the pituitary adenylate cyclase activating peptide (PAC1) and C-X-C chemokine receptor type 1 (CXCR1) (LERNER *et al.* 1991; SVENSJO *et al.* 2011). In addition, the maxadilan, it has been shown to possess immunomodulatory characteristics with the elevation of intracellular macrophage cAMP, in order to reduce TNF- α production and increase IL-6 production (SoARES *et al.* 1998).

Analyzing the salivary gland of the Ph. arabicus transcriptome, which was recently shown to transmit Leishmania tropica, the causative agent of cutaneous leishmaniasis, four hyaluronidase coding sequences were identified (HOSTOMSKÁ et al. 2009). This enzyme hydrolyzes hyaluronic acid and chondroitin sulfate, important polymers in the constitution of animal connective tissue whose function is to open the way for other glandular components that diffuse through the adjacent cutaneous tissue. This process seems to aid in the diffusion of the anti-hemostatic agents in the tissue and to guarantee the establishment of microhemorrhaging caused by the bite, causing these telmophagous insects to better explore their blood supply (RIBEIRO et al. 2000). Similarly upon analysis, adenosine (ADO) and AMP present in Ph. papatasi saliva seem to utilize modulatory mechanisms through the A2a adenosine receptor (A2aR) to initiate the production of PGE2 and IL-10. Similarly, they may act in a paracrine manner since they induce regulatory T cells from effector T cells populations, implying a role in immunosuppression and establishment of the infection (CARREGARO et al. 2015).

SANDFLY SALIVA EFFECTS IN AN EXPERIMENTAL INFECTION BY LEISHMANIA

The immune system responses to the salivary effects of sandflies on hosts are complex, specific, and depend on both the phlebotomine species and the *Leishmania* species. The induction of *L. longipalpis* saliva, for example, in mice inoculated with parasites, had less marked parasitism and less infiltration of macrophages at the inoculation site, accompanied by lower production of IL-4 and IL-10 and higher levels of IL-12 (LAURENTI *et al.* 2009).

Antagonistic results for Th2 and also Th1 immune responses were found by Menezes *et al.* (2008), who demonstrated a reduction in IL-10 production in monocytes that was not stimulated by saliva, but rather by LPS; and that infection with *Le. braziliensis*, stimulated by LPS, led to a significant increase in the production of TNF- α , IL-6 and IL-8. The effects exerted by the saliva of *L. intermedia* in human monocytes were different from those with *L. longipalpis* saliva (Costa *et al.* 2004) and *Ph. papatasi* saliva (ROGERS & TITUS 2003); these differences may play an essential role in leishmaniasis.

The studies by CAVALCANTE *et al.* (2003) and COSTA *et al.* (2004) demonstrated the interference of *L. longipalpis* salivary gland extracts in the complement cascade. The saliva acted on the classical route and the alternative route. They also found that the inactivation is not related to the inhibition of serine proteases from the complement system. The saliva of *L. longipalpis* also affected the expression of costimulatory molecules on the surface of monocytes, macrophages and human dendritic cells. Exposure of monocytes and macrophages to saliva extracts resulted in decreased CD80 expression and increased HLA-DR, whereas CD86 was increased only in monocytes and remained unchanged in macrophages.

After the evaluation of maxadilan on the secretion of cytokines and NO on the survival of *Le. major* through stimulation in macrophages in mice with LPS, not only the increase in the response of type 2 cytokine (IL-10), but also a decrease in type 1 cytokines (IL-12p70 and TNF- α) and NO was observed (BRODIE *et al.* 2007). A subsequent study analyzed that dendritic cells of mice, when exposed to maxadilan, presented altered levels of the co-stimulatory molecules CD80 and CD86. In addition to increased secretion of type 2 cytokine (IL-10) and decreased secretion of type 1 cytokines (TNF- α , IL-12p40, and IFN- γ), this exposure to maxadilan consistently reduced the extent of dendritic cell-induced CD4+ T cell proliferation and boosted the targeting for a Th2 response (WHEAT *et al.* 2008).

It has been shown that the salivary gland lysate of *L. longipalpis* may be a facilitator in *Leishmania* infection, while modulating leukocyte recruitment and Th2 cytokine production in the inflammatory focus (MONTEIRO *et al.* 2007). However, in the species *L. intermedia* pre-exposure to saliva did not result in effective protection against *Le. braziliensis* infection. Mice immunized with saliva extract when inoculated with parasites and saliva showed continuous multiplication of parasites and a low proportion between IFN- γ and IL-4 levels (MOURA *et al.* 2007).

In order to reproduce the seasonality of host-vector interaction in nature, ROHOUŠOVÁ *et al.* (2011) exposed mice to *Ph. duboscqi* bites for long and short term periods and inoculated *Le. major* soon thereafter or after a period with no sandfly bites. Only animals exposed to short-term bites and inoculated with parasites and saliva extract soon after sensitization developed protection against infection, as demonstrated previously by KAMHAWI *et al.* (2000). They then suggested that the results obtained may explain the persistence of Leishmaniasis in endemic areas. Based on the frequent exposure of individuals from endemic areas, the study done by MUKBEL *et al.* (2016) investigated the human immune response in endemic areas of *Ph. papatasi* in volunteers exposed to frequent bites. It was found that the most reactive salivary proteins for the production of antibodies are SP28, SP32 and SP36. To date, it is known that the main reactive humoral salivary proteins are: SP28, SP32, SP36, SP42 and SP32, among them the immunodominant SP28, SP32 and SP36 (ABDELADHIM *et al.* 2011).

In mice deficient in B lymphocytes, immunized with the SP15 plasmid vaccine, *Le. major* infection was successfully controlled (VALENZUELA *et al.* 2001). Based on this, it was suggested that the salivary protein PpSP15 from *Ph. papatasi* as well as *Ph. dubosqui* would be the best mediator of immune cell activation and, thus, a promising candidate for the development of a future vaccine (ELNAIEM *et al.* 2005; LESTINOVA *et al.* 2015).

Analyses in dogs also detected a strong immune response to two vector salivary proteins. In a delayed-type hypersensitivity (DTH) test performed with 35 salivary proteins from *L. longipalpis*, it was found that the proteins LJL143 and LJM17 elicited the most intense responses, comparable to that of the whole saliva extract. The adaptive immune response was evidenced by the dominance of IgG2 antibodies and substantial production of IFN- γ by CD3⁺CD4⁺ T cells stimulated with recombinant proteins or saliva extract. *In vitro*, macrophages infected with *Le. infantum* efficiently killed parasites after the addition of autologous T cells from dogs immunized with LJL143 and LJM17 and stimulated with saliva, showing a marked reduction in infection in macrophages (COLLIN *et al.* 2009).

In hamsters, immunization promoted by vaccines of plasmids coding for LJM19 (salivary antigen) and KMP11 (*Leishmania* antigen) were evaluated. Animals immunized and infected with *Le. infantum* maintained a normal hematological profile even five months after infection, whereas unimmunized hamsters showed significant anemia (SILVA *et al.* 2011).

Specific markers of mosquito exposure were found to be exclusive to L. longipalpis saliva. Among nine proteins tested, in the study by TEIXEIRA et al. (2010), LJM17 was the only one recognized by the serum of three different hosts (humans, dogs and foxes); LJM11 was recognized by human and dog serum. They determined, through reactive intensity, that LJM17 and LJM11 are specifically recognized by the serum of humans inhabiting endemic areas for VL transmitted by L. longipalpis. Xu et al. (2011) report that in mice, LJM11 confers protective immunity against Le. major infection and add that splenocytes from mice sensitized with L. longipalpis bites produced IFN-y under LJM11 stimulus, demonstrating the induction of a Th1based immunospecific systemic response. Recents work's confirm the immunomodulatory role of Phlebotomium saliva, such as, improving the function of anti-leishmania macrophages with the production of nitric oxide (NO), tested in rabbits (PUSHPANJALI et al. 2016), Immunity to the sand fly salivary protein SALO protected hamsters against Leishmania infantum and Le braziliensis infection and, more recently, a vaccine combination of a genetically modified Leishmania with SALO conferred strong protection against Le. donovani infection (Asojo et al. 2017).

Studies on the salivary effects of sandflies reflect the importance of research to understand the challenge posed by Leishmaniasis. The advances have not yet been able to meet the real needs of the immense population exposed to the risk of the disease. Despite the withdrawn diversification over the last years, most published articles restrict their investigation to the relationship of the saliva of two species, *L. longipalpis* and *Ph. papatasi*, on infection by one parasite, *Le. major*. This concentration in the scientific production contrasts with the amplitude of relations undertaken by the various vectors in the infection of the most diverse types of *Leishmania*. We suggest that the wealth of information already described by other specific studies on these species or that the fact that these relationships already have a certain degree of understanding further drive the search for data in this way.

These reports consolidate knowledge about the salivary properties of sandflies in the intensification of *Leishmania* infection. From this, it is more evident that the same biochemical and pharmacological diversity of the salivary apparatus capable of favoring the parasite may be useful for the development of protective mechanisms for the hosts. The intense immunomodulatory activity of some compounds has been identified with the intention of reversing this role and stimulating an effective defense against the disease.

FINAL CONSIDERATIONS

The present review demonstrated that, according with the literature, the vector-host relationship is not limited to the simple inoculation of the parasite in the organism and that the influence determined by the insect on the induction of responses by the host is important in the dynamics of the disease and has been intensively investigated. Adding knowledge to the mechanism of triggering the initial immune response of the host organism is of fundamental importance, considering that this phase is decisive for the progression of the infection. However, managing variables from at least three different interrelated organisms evidencing processes with varying degrees of dependence is a major challenge. Advances in the development of vaccines for humans, and especially for dogs, are extremely important. In view of the fact that in many countries euthanasia of infected animals is still advocated. Therefore, an effective and low-cost vaccine as a control measure for visceral leishmaniasis would bring benefits. especially for dogs from endemic areas in low-income regions.

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