Trypanosoma cruzi-elicited CD8⁺ T cell-mediated myocarditis: chemokine receptors and adhesion molecules as potential therapeutic targets to control chronic inflammation?

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In Chagas' disease, the establishment of inflammatory processes is crucial for Trypanosoma cruzi control in target tissues and the establishment of host/parasite equilibrium. However, in about 30% of the patients inflammation becomes progressive, resulting in chronic disease, mainly characterized by myocarditis. Although several hypothesis have been raised to explain the pathogenesis of chagasic myocardialopathy, including the persistence of the parasite and/or participation of autoimmune processes [1,2], the molecular mechanisms underlying the establishment of the inflammatory process leading to parasitism control but also contributing to the maintenance of T. cruzi-elicited chronic myocarditis remain unsolved.

Trying to shed light on these questions, we have for several years been working with murine models for Chagas' disease that reproduce the acute self-resolving meningoencephalitis [3], the encephalitis resulting of reactivation described in immunodeficient individuals [4] and several aspects of the acute and chronic myocarditis [5,6]. In the present article, our results are summarized and discussed under the light of the current literature. Furthermore, rational therapeutic intervention strategies based on integrin-mediated adhesion and chemokine receptor-driven recruitment of leukocytes are proposed to control T. cruzi-elicited unbalanced inflammation.

During an infectious process, the recruitment and migration of leukocytes towards and inside a target tissue is crucial for resolving the infectious and reestablishment of homeostasis. Both recruitment and migration are multi-step processes that depend on the nature and state of activation of the leukocytes, involve pro-inflammatory cytokines, adhesion molecules and extracellular matrix (ECM) components, being co-ordinated by a mosaic of chemoattractant molecules (chemokines) [7-13].

Several studies have demonstrated that CD8⁺ cells are the predominant cell population in the cardiac tissue of chronic chagasic patients [14-17]. Using C3H/HeJ mice infected with the Colombiana strain of T. cruzi we showed the prevalence of CD8⁺ T cells (CD4:CD8 ratio of 1:3) in the inflamed heart of chronically T. cruzi-infected experimental animals. Akin chagasic patients, amastigote forms and parasite antigens were scarce in the chronically inflamed myocardium and associated with the presence of rare IL-4 producing cells, while large amounts of the pro-inflammatory cytokines IFN-γ and TNF-α were detected [16,17,6].

Thus, our results led us to consider that the infection of C3H/HeJ mice with the Colombiana strain of T. cruzi reproduced several aspects of the chagasic chronic myocarditis and was an appropriate model to study the molecular mechanisms involved in the recruitment and accumulation of inflammatory cells into the cardiac tissue leading to the establishment of chronic CD8-mediated T. cruzi-elicited myocarditis. Initially, we showed that the predominance of CD8⁺ T cells in the cardiac tissue was not a result of the influx of the predominant cell population in the immune compartments, since during the chronic infection CD4⁺ cells represent the prevalent T cell population in peripheral blood and lymphoid tissues. Therefore, insight into the mechanisms participating and regulating the selective CD8⁺ T cell trafficking is crucial when trying to understand how T. cruzi-elicited myocarditis is established.

The requirement and functional participation of L-selectin (CD62L), leukocyte function-associated antigen 1 (LFA-1; CD11a,CD18) and very late activation antigen 4 (VLA-4; CD49d,CD29; α 4β1) in T cell activation and migration has been extensively demonstrated in other model [18,19]. For this reason, we investigated the expression of CD62L, LFA-1 and VLA-4 on cells isolated from heart tissue, and peripheral blood of T. cruzi-infected mice. A high percentage of circulating CD8⁺ T cells and most of the CD4⁺ and CD8⁺ T cells present in myocardium of T. cruzi-infected animals express an activation phenotype characterized as CD62LLow, LFA-
Interestingly, the distribution of CD4+ and CD8+ T-cells in the myocardium mirrors the frequency of cells expressing the CD62LLow/LFA-1High/VLA-4High activation phenotype among CD4+ and CD8+ peripheral blood T cells. In agreement with these results, a recent study demonstrated that patients with a mild form of Chagas' disease showed a lower percentage of CD3α4+ peripheral blood cells, in comparison with those with severe chronic cardiopathy. Increased expression of ECM components particularly fibronectin (FN), a VLA-4 ligand, have been previously detected in the myocardium of T. cruzi-infected mice during the acute and chronic phases. Studying Colombiana-infected C3H/HeJ mice, we confirmed this finding and showed that a fine interstitial FN mesh present in the inflamed heart involves VLA-4+ CD4+ and CD8+ T cells as well as the infected and non-infected myocytes. Interestingly, the FN filamentous network present in the inflamed myocardium may function as a pathway in T cell migration process via VLA-4 interactions but may also influence T cell activation. In fact, it was demonstrated that antibodies against VLA-4 increase the anti-CD3-induced T cell adherence to FN and sinergistically induce T cell proliferation. Then, it is reasoned to propose that interactions of VLA-4+ CD4+ and CD8+ T cells with the FN mesh present in the inflamed heart could contribute to the perpetuation of the inflammation in this tissue influencing antigen-specific T cell recognition, activation, proliferation, survival and effector activity. Consistently, besides the FN network surrounding VLA-4+ activated mononuclear cells, endothelial cells expressing vascular cell adhesion molecules 1 (VCAM-1), another VLA-4 ligand, were found in the inflamed myocardium, supporting the possibility that VLA-4-mediated interactions might be important for leukocyte entrance in the cardiac tissue, cell migration and positioning towards infected sites as well as persistence of leukocyte activation phenotype inside the inflamed heart.

More recently, we have demonstrated that when compared with non-infected or 7 days T. cruzi-infected animals, 14 days infected mice present a large proportion of circulating CD8+ T cells expressing the VLA-4+ activation phenotype. Further, the CD8-mediated myocarditis is established on day 28 post-infection. In this moment, most of the mononuclear cells forming local inflammatory infiltrates surrounding T. cruzi-infected cardiomyocytes are CD8+ lymphocytes, whereas CD4+ cells constituted diffuse infiltrates (APMP Marino and J Lannes-Vieira, unpublished results). Altogether, these results led us to speculate that the predominance of CD8+ T cells in the inflamed myocardium of T. cruzi-infected host might result of the predominance of CD8+ expressing the activated phenotype (CD62Low/LFA-1High/VLA-4High) in peripheral blood which is established during the early acute infection and persisted during the chronic disease. The use of immunomodulators such as IFN-β able to downregulate the expression of VLA-4 or monoclunal antibodies and antagonists specific for adhesion molecules are being proposed and successfully used to ameliorate inflammatory processes. Therefore, our results point to the possibility that adhesion molecules may constitute a logic target to control the inflammation elicited by T. cruzi infection. In this context, anti-VLA-4 monoclonal antibodies selectively inhibited the migration of activated T cells to the brain tissue of T. cruzi-infected mice. Presently we are testing the participation of VLA-4-mediated interactions in the establishment of T. cruzi-elicited acute and chronic myocarditis, thus paving the way for the development of a VLA-4-based therapy for T. cruzi-elicited myocarditis.

Our previous report showing that a CD8-mediated encephalitis is restricted to the acute phase in the C3H/He mice infected Colombiana strain of T. Cruzi, however, provides evidence that the persistence of chronic myocarditis with the predominance of CD8+ T cells does not only result from the activation phenotype expressed by peripheral blood CD8+ T cells, although it may represent a requirement for the entrance of these cells into the target tissue. Thus, other factors present in the cardiac environment could also be involved in the migration, retention and activation of inflammatory cells leading to perpetuation of the myocarditis in the T. cruzi-infected mice.

It is considered that the cytokines produced in the heart tissue during the initial immune response will influence the subsequent immune reaction. In this vein, our data indicate that TNF-α, IFN-γ and IFN-γ-induced chemokines RANTES (regulated upon activation, normal T cell expressed and secreted), MIG (monokine induced by IFN-γ) and CRG-2/IP-10 (cytokine response gene 2/interferon g-inducible protein 10), as well as JE/MCP-1 (monocyte chemoattractant protein-1) and MIP1α (macrophage inflammatory protein 1 a) were found to be the dominant cytokines expressed in situ during acute infection, persisting during the chronic phase of T. cruzi-elicited myocarditis and may contribute to the intense recruitment of activated T cells.

Previous report showed that during Listeria monocytogenes infection, CD8+ T cells are the main source of MIP-1α and that this chemokine has the ability to preferentially recruit the CD8+ T cell subpopulation. In addition, in vitro experiments showed that MIP-1α preferentially induces chemotaxis of CD8+ T lymphocytes, whereas MIP-1β is related to the preferential recruitment of CD4+ T cells. Altogether, these findings led us to propose that IFN-γ and IFN-γ-elicited RANTES, JE/MCP-1 and MIP-1α chemoattractants present in the inflamed heart of Colombiana-infected C3H/HeJ mice create a favourable environment for selective and
preferential migration of CD8+ T cells towards this tissue, leading to a CD8-mediated myocarditis [6].

The pro-inflammatory cytokines TNF-α and IFN-γ play a major role controlling tissue parasitism during T. cruzi infection [27,28]. These cytokines are present in the inflamed heart of chronic chagasic patients [14,17], and acute and chronic experimentally T. cruzi-infected mice [29,5,6], suggesting that besides contributing to parasitism control they could be also involved in the maintenance of chronic myocarditis [27,28]. Recently we showed that the heart and central nervous system are the main sites of reactivation of T. cruzi infection in mice lacking functional genes for IFN-γ and IL-12, respectively.

This reactivation was characterized by intense inflammation accompanied by remarkable enhancement in parasitism, particularly resembling the parasite-elicited mass encephalitis often found in chronic chagasic patients with AIDS[30]. Importantly, during reactivation of infection the inflammatory processes in the central nervous system from IL-12 knockout mice were mostly devoid of CD8+ T cells and mainly composed of polymorphic nuclear cells, macrophages and CD4+ T cells, whereas in the heart CD8+ T cells predominate over CD4+ T cells. Consistent with the hypothesis that CD8+ T lymphocytes are the major source of IFN-γ and IFN-γ-inducible chemokines that orchestrate the preferential CD8 migration during T. cruzi infection, we found IFN-γ-producing cells in the heart, but not in the central nervous system of these animals (V Michailowsky and J Lannes-Vieira, unpublished results).

Thus, considering the importance of CD8+ T cells in resistance to T. Cruzi [31,32], the increased susceptibility to parasite replication and mass encephalitis could be explained by the lack of CD8 T cell migration and local production of IFN-γ and IFN-γ-inducible chemokines in IL-12 deficient mice. Our results also showed that in contrast to IFN-γ knockout mice, splenocytes from IL-12 knockout mice infected T. cruzi produced low levels of IFN-γ upon stimulation with parasite antigens. Thus, our data suggest that the level of interferon-g deficiency is a major determinant of the site of reactivation of T. cruzi infection in immunocompromised host [4].

Chemokines are small (8-14 kDa) inducible cytokines that recognize a large group of 7 transmembrane-spanning G-protein-coupled serpentine receptors displayed on the leukocyte surface and are involved in normal trafficking of leukocytes to both lymphoid and nonlymphoid organs and recruitment of these cells to sites of injury and infection. Moreover, chemokines also appear to affect several other immunological phenomena, and play an important role in immune regulation, mediating leukocyte activation, costimulation and differentiation during innate and adaptive immune responses [11-13].

Recent studies have shown that human macrophages in vitro infected with T. Cruzi [33], and in vivo and in vitro T. cruzi-infected peritoneal macrophages and cardiomyocytes produce RANTES, MIP-1α and JE/MCP-1 and respond to these chemokines increasing T. cruzi uptake, enhancing nitric oxide production and controlling parasite replication [34,35]. Further, using knockout mice infected with T. cruzi we showed that both IFN-γ and TNF-α are essential for the production of RANTES and MIP-1α, respectively [36]. Also, IFN-γ enhanced the in vitro production of RANTES and IP-10 by macrophages infected with T. cruzi or treated with glycosylphosphatidylinositol-anchored mucin-like glycoproteins of trypomastigotes (tGPI-mucins) [36,5]. Thus, it is possible that IFN-γ-induced chemokines present in the cardiac tissue of T. cruzi-infected individuals could also participate in parasitism control.

Considering a general model for Chagas' pathogenesis, the results discussed above suggest that the predominance of CD8+ T cells in the myocardium of T. cruzi-infected individuals reflects the profile of adhesion molecules and chemokine receptors displayed by the circulating CD8+ T cells and point to the possibility that multiple IFN-γ-inducible molecules present in the inflamed tissue contribute to the genesis and maintenance of T. cruzi-induced myocarditis.

The corollary of this hypothesis is that activated CD8+ T cells present in the inflamed heart as well as a large proportion of circulating CD8 lymphocytes of T. cruzi-infected individuals differentially bear CC chemokines receptors such as CCR1, CCR3, CCR5 [37], in comparison with CD4+ T cells. Interestingly, recent reports show that the expression of CCR5 is increased on CD3+CD8+ peripheral blood T cells obtained from patients with mild cardiomyopathy compared with uninfected individuals or patients with severe disease [38]. In agreement with this finding, it has been shown that a CCR5 promoter point mutation was significantly increased in asymptomatic patients compared with patients with cardiomyopathy [39]. Moreover, it has been reported that most CD8+ T cells present in the heart of T. cruzi-infected mice express CCR5 in heart [40].

Altogether, these results and our data discussed above led us to consider that CC chemokines, specially RANTES and MIP-1α, and CC receptors, mainly CCR5, could be involved in differential cell migration and
pathogenesis of T. cruzi-elicited CD8-mediated myocarditis. To test this possibility we used Met-RANTES, a N-terminally modified human RANTES that inhibits agonist activity at CCR1 and CCR5 [41], to modulate the establishment of T. cruzi-elicited acute myocarditis. Our results showed that Met-RANTES treatment increased the survival of T. cruzi-infected animals when compared to saline treatment. Moreover, Met-RANTES treatment significantly decreased the inflammatory infiltrates due to CD4 and CD8 T cells, but decreases or does not interfere, depending on the therapeutic scheme used, with the heart parasitism [42].

These results indicate that the massive influx of CC chemokine receptors-bearing inflammatory cells into the heart tissue is not crucial for cell-mediated anti-T. cruzi immunity, however it seems to be critical for T. cruzi-elicited immunopathology. Thus, the beneficial role of Met-RANTES may relay on the reestablishment of balanced inflammation, however further investigation are required to unravel the beneficial effector mechanisms of Met-RANTES. Anyhow, the inhibition of CC chemokine receptors might become an attractive therapeutic strategy for further evaluation during T. cruzi infection.

Thus, further studies may provide additional insights into the mechanisms by which adhesion molecules, chemokines and chemokine receptors contribute to the establishment and regulation of the immune response in the inflamed heart during T. cruzi infection and may unravel novel therapeutic strategies aimed at limiting the inflammatory damage to cardiomyocytes, targeting integrin-mediated adhesion and chemokine-driven recruitment of leukocytes besides the specific anti-parasite drugs (Figure 1).

Figure 1: Leukocyte migration from the circulation into target tissues is a multi-step process. The first step involves transient selectin-mediated binding (rolling). Next, integrins on leukocytes are activated by chemokines (triggering), resulting in firm adhesion between leukocytes and activated adhesion molecules-bearing endothelial cells (EC). Finally, leukocytes extravasate through the vascular wall and into the tissue in response to chemokine-driven recruitment (transmigration) [19]. During T. cruzi infection, circulating CD8+ T cells express a differential profile of adhesion molecules (LFA-1Hi, VLA-4Hi). This could account for the preferential interaction with activated EC (ICAM-1, VCAM-1). Also, the inflamed tissue presents pro-inflammatory cytokines (IFN-γ, TNF-α) and chemokines (RANTES, MIP-1α, JE/MCP-1). These might selectively and preferentially recruit CD8+ T cells leading to the establishment of CD8-mediated myocarditis [5,6]. Novel therapeutic strategies aiming at limiting the inflammatory damage of cardiomyocytes, targeting integrin-mediated adhesion and chemokine-driven recruitment of leukocytes, besides the specific anti-T. cruzi drugs, should be evaluated.

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