

Chronic *Trypanosoma cruzi*-elicited cardiomyopathy: from the discovery to the proposal of rational therapeutic interventions targeting cell adhesion molecules and chemokine receptors - how to make a dream come true

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One hundred years ago, Carlos Chagas discovered a new disease, the American trypanosomiasis. Chagas and co-workers later characterised the disease's common manifestation, chronic cardiomyopathy, and suggested that parasitic persistence coupled with inflammation was the key underlying pathogenic mechanism. Better comprehension of the molecular mechanisms leading to clinical heart afflictions is a prerequisite to developing new therapies that ameliorate inflammation and improve heart function without hampering parasite control. Here, we review recent data showing that distinct cell adhesion molecules, chemokines and chemokine receptors participate in anti-parasite immunity and/or detrimental leukocyte trafficking to the heart. Moreover, we offer evidence that CC-chemokine receptors may be attractive therapeutic targets aiming to regain homeostatic balance in parasite/host interaction thereby improving prognosis, supporting that it is becoming a non-phantasious proposal.

Key words: Chagas disease - *Trypanosoma cruzi* - cardiomyopathy - chemokine - chemokine receptors - cell adhesion molecules

In 1909, Carlos Chagas realised a unique medical finding describing a new disease known as American trypanosomiasis, as well as its vector and causative agent, the haemoflagellate protozoan *Trypanosoma cruzi* (Chagas 1909). This discovery initiated a movement to characterise the disease's clinical manifestations and epidemiological significance (Laranja et al. 1956). One hundred years after its discovery, Chagas disease is still epidemiologically relevant, afflicting 12-14 million people in 18 endemic countries in Latin America (Coura 2007, Dias 2007). Several hundred thousand *T. cruzi*-infected individuals may also be living in North America, Europe and other countries as a result of immigration to these areas in search of a better life (Schmunis 2007), demonstrating that Chagas disease is a far-reaching public health issue.

Chagas' (1909, 1911) initial studies described the peculiar clinical symptoms associated with American trypanosomiasis, including cardiac excitability disturbance, extrasystoles, abnormal conduction and complete

heart blockage. In 1916, Chagas compiled the histopathological findings from acute and chronic cases of human as well as experimental *T. cruzi* infection. Based on these data, he proposed that the pathogenic mechanism underlying cardiac symptoms was intense parasitism coupled with inflammation during the acute phase and parasite persistence, "although in small numbers inside cardiomyocytes making them difficult to find", in parallel with inflammation during chronic infection (Fig. 1). Curiously, Chagas described the presence of amastigote forms of the parasite in the heart tissue of chronically infected individuals (Fig. 1). Assuming that inflammation was not proportional to the small numbers of parasite nests, he claimed that the inflammatory lesions resembled "an allergic process". However, Chagas noted that in "most of the chronic cases with predominant cardiac signs a large number of protozoa is observed in the myocardium and the inflammatory reactions of the organ is also more intense" (Chagas 1916).

During the following years, Chagas and Villela (1922) adopting a new non-invasive procedure for recording the heart's electrical activity (electrocardiography, or ECG) and recently described parameters to characterize heart abnormalities defined the precise cardiologic profiles of acutely and chronically *T. cruzi*-infected patients. However, three decades passed before for Chagas' original observations of cardiac disturbances during chronic *T. cruzi* infection were reproduced in experimentally infected dogs (Laranja et al. 1950, 1956) and confirmed in a large number of patients (Laranja et al. 1956). Chagas' detection of parasite nesting in cardiomyocyte or phagocytized by macrophages in all studied chronic chagasic

Financial support: CNPq (301504/2005-9, 302534/2008-3), FAPERJ (E-26/111.756/2008), DECIT/MS/CNPq/MCT, The Millennium Institute for Vaccine Development and Technology (420067/2005-1), INCT/CNPq (fellowships from CNPq: JLV, IRP, JCS)

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Received 20 April 2009

Accepted 1 June 2009

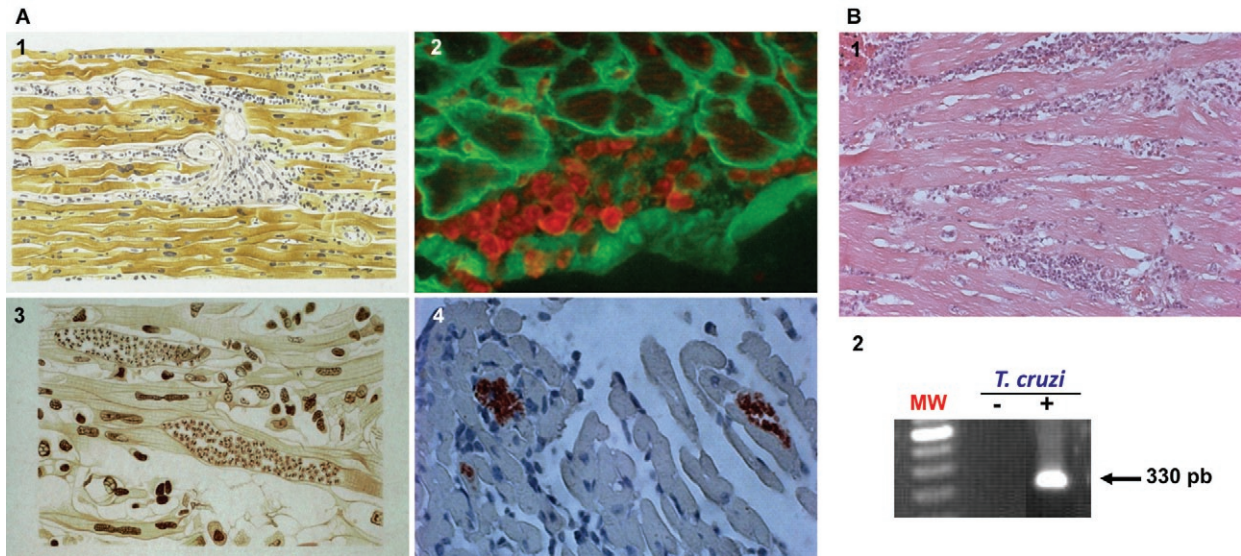


Fig. 1: histopathological alterations observed in the heart tissue of Colombian strain infected C3H/He mice reproduce the original findings described by Chagas (1916) in patients. A: in 1916, Chagas showed the presence of an intense inflammatory infiltrate in the heart of patients acutely infected by *T. cruzi* (1). The presence of inflammatory cells in the heart tissue of acutely Colombian-infected mice (2) is shown using confocal microscopy (1,000X) (Roffê et al. (2003) (red: inflammatory cell; green: cardiomyocytes). Similarly to Chagas's description showing the presence of amastigote forms of the parasite in the heart tissue of chronically infected individuals (3), we detected the presence of *T. cruzi* antigens (4) (400X), immunohistochemistry carried out (dos Santos et al. 2001); B: in the case of apparent absence of the parasite in inflammatory chronic heart lesions (1) (200X, H&E staining), parasite DNA was detected during the chronic phase of experimental infection (2). PCR conditions described in Carvalho et al. (2003). Authorized reproduction of drawings published in Chagas (1916).

cardiomyopathy (CCC) patients (Chagas 1916, Chagas & Vilella 1922), although “a time-consuming activity” was also confirmed by Laranja et al. (1956). Moreover, cardiac histopathological abnormalities were characterised as consisting of diffuse and focal heart inflammatory foci chiefly composed of lymphocytes, plasma cells and macrophages, as well as eosinophils and neutrophils. Focal and diffuse fibrosis of the myocardium was also evidenced in all cases, its intensity varying from case to case and from one area to another within the same case (Laranja et al. 1956). Together, these findings provide a firm basis for defining CCC as a distinct clinical and anatomical entity.

Intriguingly, *T. cruzi* parasites were not observed in the cardiac tissue lesions of most patients with chronic Chagas disease for decades. Scientists began to ascribe a major role to autoimmune reactions in CCC pathogenesis (Santos-Buch & Teixeira 1974, Ribeiro dos Santos & Hudson 1980, Kierszenbaum 1985), a theory that hampered efforts to develop chemotherapy and vaccine against *T. cruzi* (Kierszenbaum 2005). The importance of parasite persistence in CCC pathogenesis was finally reaffirmed nearly 80 years after Chagas' pioneering work (Chagas 1916) using immunohistochemistry and molecular biology to analyse the heart tissue of CCC patients (D'Avila Reis et al. 1993, Higuchi et al. 1993, Jones et al. 1993, Reis et al. 1997), as well as ingenious cell transfer studies in experimental models (Tarleton et al. 1997). Currently, it is widely believed that chagasic

cardiac injury results from an imbalanced effector immune response elicited by parasite persistence (Higuchi et al. 2003, Kierszenbaum 2005).

Adopting the experimental model consisting of female C3H/He mice infected with low-dose inocula of the Colombian *T. cruzi* strain, we demonstrated the presence of parasite nests and the persistence of parasite DNA in inflammatory cardiac lesions (Fig. 1), reproducing earlier findings in chronic chagasic patients (Higuchi et al. 1993, Jones et al. 1993). Our model also replicates Chagas' main findings in CCC patients (Chagas 1916, Chagas & Vilella 1922), including (i) parasitaemia control and acute phase survival; (ii) chronic increase in relative heart weight; (iii) acute and chronic myocarditis; (iv) cardiomyocyte lesions and (v) electrical abnormalities (Fig. 2).

In the past several years, many groups have sought to elucidate the physiopathogenesis of CCC and cardiomyopathy (Higuchi et al. 2003, Marin-Neto et al. 2008). Nevertheless, new therapeutic approaches, beyond already available anti-parasite drugs and interventions targeting symptoms, have not been developed (Marin-Neto et al. 2008, Soeiro & de Castro 2009). Since cardiac transplant is currently the only effective treatment for Chagas disease-induced heart failure, cell-based therapy aiming to recover heart function is a desirable alternative, although expensive and still restricted to clinical trials (Soares & Ribeiro dos Santos 2009). The major challenge in designing efficacious treatment for *T. cruzi*-elicited cardiomyopathy is to reduce inflammation,

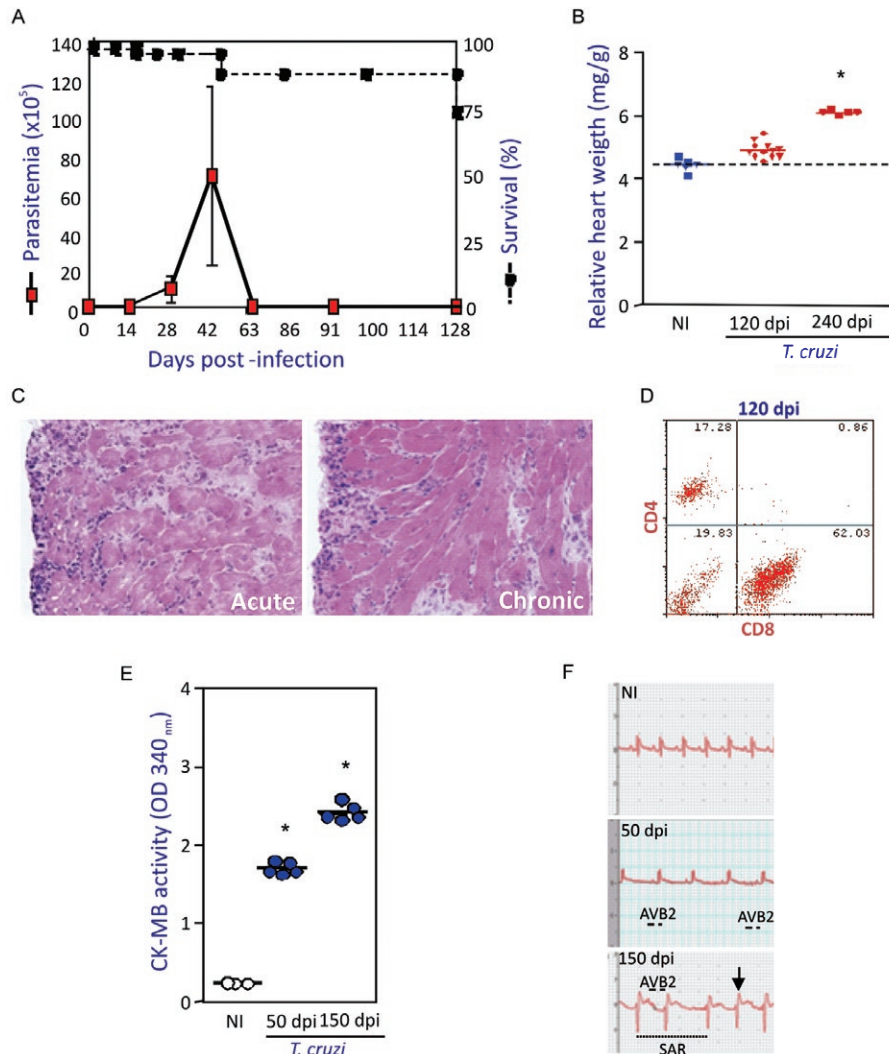


Fig. 2: experimental model reproduces different findings of the cardiac form of Chagas disease. Five to seven-week-old female C3H/He (H-2^b) mice were infected with 100 trypomastigote forms of the Colombian *Trypanosoma cruzi* strain (dos Santos et al. 2001). Panel A shows parasitemia (red squares) and survival curves (black squares). Increase in relative heart weight is observed at 120 days post-infection (dpi) and mainly, at 240 dpi (red triangles and squares, respectively) compared to non-infected C3H/He mice (blue squares/triangles). In panel C, histopathological analysis (400X, H&E) shows acute (50 dpi) and chronic (120 dpi) mononuclear cell inflammation in the cardiac tissue of infected mice. Flow cytometry analysis of the mononuclear cells isolated from cardiac tissue of chronically (120 dpi) *T. cruzi*-infected mice (dos Santos et al. 2001) shows the predominance of CD8⁺ T cells over CD4⁺ T cells (D). Cardiomyocyte damage was evaluated by CK-MB activity in sera of acute (50 dpi) and chronically (150 dpi) infected (blue circles) in comparison with non-infected (white circles) C3H/He mice (E) (Kroll-Palhares et al. 2008). ECG recordings from non-infected and *T. cruzi*-infected mice (F) [ECG recorded as shown in Silverio et al. (2009), unpublished] Mean \pm SD values for 5-10 mice are shown and data are representative of three independent experiments. *: $p < 0.05$; **: $p < 0.01$.

which causes severe tissue damage, without hampering immunological control of parasitism. A more thorough understanding of CCC pathogenesis and its underlying molecular mechanisms may thus help to delineate new therapeutic strategies.

Here, we critically review our and other recent data exploring the participation of cell adhesion molecules (CAM) and CC-chemokines and their receptors in the multi-step process of cell migration during *T. cruzi*-elicited myocarditis and CCC formation. We also offer evidence that these molecular pathways are rational

therapeutic targets for modulating detrimental cardiac inflammation during chagasic infection without hampering parasite control.

Key players in leukocyte migration to injured tissues

The three crucial steps of leukocyte migration to injured non-lymphoid tissues (rolling, adhesion and transmigration) were first described in the XIX century (Ley et al. 2007). However, only in the last decade of the XX century, after the discovery of molecules involved in the leukocyte migration cascade, has this multi-step

model emerged as a concept and begun to unravel the leukocyte recruitment process to injured tissue (Butcher 1991, Springer 1995). In the original model, Butcher and Springer proposed three steps underlying leukocyte migration: (i) selectin-mediated rolling; (ii) chemokine-driven activation and (iii) integrin-dependent arrest or firm adhesion. However, recent findings have revealed that the leukocyte migration cascade is more complex, involving tethering or capture, rolling, arrest or firm adhesion, recruitment, transendothelial migration (intraluminal crawling and paracellular and transcellular migration), extravasation through the basement membrane and positioning (Ley et al. 2007, Sigmundsdottir & Butcher 2008). Notably, all of these studies confirmed the participation of chemokines, chemokine receptors and CAM in leukocyte trafficking to injured tissues.

Chemokines are small (8-14 kDa) chemoattractant cytokines that are either constitutively expressed or induced in response to inflammation. These proteins eq o rtkug" hqwt" uwdhc o knkgu" *EZE" qt" . "EE" qt" . "E" qt" " and CX₃E" qt" + "v jcv" cev" qp" I / r tqv gkp/ eqwrngf" ugtrgpvkpg" receptors on target cells. Generally, CXC-chemokines act on neutrophils and on T and B cells, while CC-chemokines affect a wider spectrum of cell populations, including monocytes, basophils, eosinophils, T cells, dendritic cells and natural killer cells, but do not act on neutrophils. It is likely that all cell types, such as endothelial cells, cardiomyocytes, fibroblasts, T cells and macrophages, produce a variety of chemokines when adequately stimulated (Bromley et al. 2008, Viola & Luster 2008). Chemokines act in a context-dependent manner, varying in function according to whether they are in soluble or extracellular matrix-anchored form (Sigmundsdottir & Butcher 2008). Their biological functions are numerous, including facilitating (i) cell adhesion to the endothelium and extracellular matrix components; (ii) leukocyte chemotactic migration; (iii) angiogenesis control; (iv) extracellular matrix production control; (v) regulation of T-lymphocyte growth, activation and differentiation (including Th1/Th2 commitment); (vi) release of intracellular reserve of enzymes; (vii) oxygen radical formation; (viii) cytoskeleton alteration; (ix) generation of lipid mediators of intracellular activation signals; (x) haematopoietic precursor proliferation and (xi) control of pathogen invasion and propagation. Chemokines may thus play an important role in the regulation of host resistance to pathogens and in the immunopathogenesis of various conditions, including autoimmune and infectious diseases (Proudfoot 2002, Bromley et al. 2008, Viola & Luster 2008).

CAM, which are mainly members of the selectin, integrin and immunoglobulin superfamilies, participate in all cell migration steps involving cell-cell and cell-extracellular matrix interactions (Luo et al. 2007). These molecules are widespread, present on a variety of cell types and involved in diverse biological events, including (i) ontogenesis; (ii) immunological synapse formation; (iii) cell activation; (iv) immunological surveillance; (v) cell extravasation to injured tissue; (vi) cell positioning and (vii) cell-pathogen interactions (Luo et al. 2007, Abram & Lowell 2009). Importantly, chemokine-triggered mod-

ulation of integrin affinity is a crucial step in cell arrest or firm adhesion to the activated endothelium, making this process a potential target for impairing cell extravasation during chronic inflammatory diseases (Kummer & Ginsberg 2006, Ley et al. 2007, Luo et al. 2007, Viola & Luster 2008).

T. cruzi-elicited myocarditis

Although an inflammatory response seems to be required for parasite control in acute Chagas disease, about 30-40% of patients also develop chronic myocarditis with prominent fibrosis and organ dysfunction (Higuchi et al. 2003, Marin-Neto et al. 2008). During the chronic phase, inflammatory heart infiltrates, which are not directly related to the rare parasite presence, are mainly composed of mononuclear cells, lymphocytes and macrophages, although neutrophils and eosinophils are also occasionally detected (Chagas 1916, Laranja et al. 1956, Reis et al. 1993, Higuchi et al. 1997, dos Santos et al. 2001). Among the lymphocytes, CD8⁺ T cells outweigh CD4⁺ T cells in the focal and diffuse heart infiltrates of CCC patients (Reis et al. 1993, Higuchi et al. 1997) and chronically infected experimental models (dos Santos et al. 2001, Medeiros et al. 2009). Conversely, there is a direct relationship between parasite presence and CD8⁺ T-cell prevalence during cardiac inflammation (Higuchi et al. 1997). However, the molecular forces governing CD8⁺ T cell-enriched myocarditis, as well as the chemokines and CAM influencing cell migration and positioning in the cardiac tissue, are not yet well defined. The functional role of invasive inflammatory cells in parasite control and cardiomyopathy is also still unknown.

Participation of CAM in T. cruzi-induced myocarditis and cardiomyopathy

Initial studies of CAM participation in Chagas disease revealed high serum levels of soluble vascular CAM-1 (sVCAM-1 or CD106), intercellular adhesion molecule-1 (ICAM-1 or CD54) and CD44 (ligand of fibronectin and hyaluronic acid) during the acute phase of infection, whereas soluble P-selectin (sP-selectin or CD62P) was associated with chronic disease severity (Laucella et al. 1996a). Furthermore, sP-selectin and sVCAM-1 levels were found to be elevated in children with indeterminate Chagas disease, but decreased after benznidazole treatment, as compared with children receiving a placebo (Laucella et al. 1999). CAM release is triggered by the same stimuli that increase their cell surface expression, uwe j" cu" kpvgtgtqp/ " *KHP/ + " cpf" vw o qwt" pgetqku" hce- tor (TNF). In this context, CCC is associated with (i) increased expression of both ICAM-1 and VCAM-1 on the cardiac vascular endothelium; (ii) increased numbers of cardiac tissue-invading lymphocytes expressing leukocyte function-associated antigen-1 (LFA-1 or CD11a/CD18; ligand of ICAM-1), very late antigen-4 (VLA-4 qt" EF6; fIEF4; . " 6 3=" nkicpf" qh" XECO/3" cpf" EF66" and (iii) upregulation of class I major histocompatibility complex on cardiomyocytes (Benvenuti et al. 2000) in the presence of parasite antigens and CD8⁺ T cell-enriched inflammation (D'Avila Reis et al. 1993, Higuchi et al. 1993, Reis et al. 1997). Based on these in situ find-

ings, it is reasonable to propose that CAM contribute to chagasic myocarditis formation by mediating lymphocyte adhesion to the cytokine-activated endothelium of cardiac blood vessels. Moreover, one could speculate that LFA-1⁺ lymphocytes may adhere to ICAM-1-bearing cardiomyocytes, contributing to parasite control but also causing tissue lesions (Benvenuti et al. 2000). CAM may thus play an integral role in creating an appropriate scenario for chagasic heart failure, a role that is currently being elucidated experimentally.

Adopting C3H/He mice infected with the Colombian *T. cruzi* strain, which develop CD8⁺ T cell-enriched cardiomyopathy and increased heart relative weight, cardiomyocyte lesions and ECG abnormalities (Fig. 2), we demonstrated that both CD4⁺ and CD8⁺ T cells infiltrating the heart tissue exhibit the activation phenotype CD62L^{-/low}, LFA-1^{high} and VLA-4^{high}. In addition, a higher proportion of circulating CD8⁺ T cells than CD4⁺ T cells are LFA-1^{high} and VLA-4^{high}, which may allow interaction of these circulating cells with ICAM-1⁺ and VCAM-1⁺ activated endothelium (dos Santos et al. 2001). This potential interaction could facilitate the preferential migration to and predominance of these T cells in the myocardium of chronically *T. cruzi*-infected animals. Interestingly, in *T. cruzi*-infected C3H/He mice, the increased frequency of LFA-1⁺ and VLA-4⁺ circulating T cells and the upregulation of ICAM-1 and VCAM-1 on the cardiac vascular endothelium coincide with inflammatory cell invasion of this tissue (Marino et al. 2003). Wukpi "kpvtngwmp/34" *KN/34+/cpf"KHP/ /fghkekgpv" okeg." yg"jcxg"cnug"hgwpf"vjcv"KHP/ " oqfwncvgu"KECO/3" gzpression on endothelial cells and on the sarcolemma of cardiomyocytes (Michailowsky et al. 2004). However, c" uwwf{" k o r ng o g p w k p i " K H P / " t g e g r v q t / f g h k e k g p v " o k e g " claimed that cardiac ICAM-1 expression in *T. cruzi*-infected mice is upregulated, suggesting that other inflammatory factors or parasite products *per se* may also contribute to CAM upregulation during *T. cruzi* infection. In fact, in vitro *T. cruzi* infection of endothelial cells induces VCAM-1 and E-selectin and upregulates ICAM-1 via "pwengct" hcevqt/ D" *PH/ D"cevxcvqpp" *Jwcp"gv"cn0"3; ; ; +0"

Infection of ICAM-1^{-/-} mice indicated a role for ICAM-1 in *T. cruzi*-elicited myocarditis formation. The high susceptibility of ICAM-1-deficient animals to *T. cruzi* infection was associated with a decreased number of CD4⁺ and CD8⁺ T cells invading the cardiac tissue, implying that LFA-1/ICAM-1-mediated cell migration to the heart is essential for myocarditis formation and thus for protective anti-*T. cruzi* responses (Michailowsky et al. 2004). However, whether different CAM have distinct roles in protective immunity and in the induction of chronic inflammation in Chagas disease is still unclear. A better understanding of the role of CAM in *T. cruzi*-elicited myocarditis and CCC may pave the way to facilitate targeting of these molecules to treat Chagas disease. CAM antagonists have already been implemented in the treatment of chronic inflammatory diseases, including rheumatoid arthritis, multiple sclerosis and Crohn's disease (Kummer & Ginsberg 2006, Abram & Lowell 2009), indicating the feasibility of this therapeutic approach.

Chemokines and chemokine receptors in *T. cruzi*-elicited myocarditis and cardiomyopathy

During *T. cruzi* infection, cytokines, chemokines and other immune mediators are generated in the affected heart tissue, triggering intense leukocyte migration (Talvani et al. 2000, dos Santos et al. 2001, Marino et al. 2004). CC-chemokines appear to be particularly beneficial to the host during *T. cruzi* infection. The parasite or its molecules have been found to induce human and murine macrophages as well as murine cardiomyocytes to produce CC-chemokines in vitro (Lima et al. 1997, Villalta et al. 1998, Aliberti et al. 1999, Machado et al. 2000, Talvani et al. 2000). Ad-fkvpq"qh" o cetqr j c i g " k p h n c o o c v q t { " r t q v g k p / 3 " * O K R / 3 . " qt" EEN5+." OKR/3 " *qt" EEN6+ "qt" TCPVGU" *Tg i wncvqpp" on Activation, Normal T-cell Expressed and Secreted or CCL5) to such infected macrophages and cardiomyocytes resulted in increased parasite up-take and destruction by a nitric oxide-dependent mechanism (Aliberti et al. 1999, Machado et al. 2000). This finding suggests that parasites regulate chemokine production, which keeps the parasite under control and thus prevents host death.

Treatment of *T. cruzi*-infected cardiomyocytes with TNF also leads to induction or augmentation of CC-chemokine expression (Machado et al. 2000). Interestingly, *T. cruzi*-infected macrophages express more CC-chemokines than untreated cells, while CC-chemokine production, suggesting that pro-inflammatory and regulatory cytokines control parasitism and myocarditis via chemokine-dependent mechanisms (Aliberti et al. 1999). Additionally, the CC-chemokine monocyte chemoattractant protein-1 (MCP-1 or CCL2) directly triggers the chemotaxis and morphogenesis of the trypomastigote forms of *T. cruzi* (Yamauchi et al. 2006), implicating CC-chemokines as crucial actors in directing parasite to certain tissues and contributing to explain *T. cruzi* tissue tropism (Melo & Brener 1978).

Elevated mRNA levels of several chemokines [CCL2/OER/3."EEN51OKR3 ."EEN61OKR/3 ."EEN71TCPVGU." o q p q m k p g " k p f w e g f " d { " K H P / " * O K I " q t " E Z E N ; + . " k p v g t h g t - o n - i n d u c i b l e p r o t e i n - 1 0 (I P - 1 0 o r C X C L 1 0) a n d R A N T - E S] have been observed in the cardiac tissue, associated with the formation of inflammatory infiltrates of various murine models of acute and chronic *T. cruzi* infection (Talvani et al. 2000, Aliberti et al. 2001, dos Santos et al. 2001, Marino et al. 2004). Indeed, we recently demonstrated that myocarditis intensity during both the acute and the chronic phases of infection in C3H/He mice is directly correlated with the CC-chemokines CCL3/MIP-3 " c p f " E E N 7 1 T C P V G U " d w v " p q v " j g " e { " v q m k p g u " K H P / " c p f " TNF, concentrations in the cardiac milieu (Medeiros et al. 2009), reaffirming CC-chemokine participation in *T. cruzi*-elicited myocarditis formation and perpetuation.

Treatment of Colombian *T. cruzi*-infected BALB/c mice with antibodies recognizing CXCL10/IP-10 or CCL5/RANTES during the acute phase did not alter parasitaemia, heart parasitism or inflammation. However, treatment with anti-CXCL10/IP-10 led to increased

parasitaemia and parasitism but did not affect either myocarditis severity or the transcript levels of inflammatory cell markers (CD4, CD8, F4/80), cytokines (TNF- α , IL-10, or CCL5/RANTES) in the heart tissue, as compared with isotype-treated control mice (Hardison et al. 2006c). These data indicate that CXC-chemokines promote a protective immune response to parasite without contributing to cardiac inflammation.

The role played by CC-chemokines in myocarditis and cardiomyopathy development during *T. cruzi* infection has also been explored. Treatment of acutely infected mice with neutralising antibodies revealed that EEN51OKR/3 "ku" kpxqnxgf" kp" o cetqr jci g" tgetwkv o gpv" to infected tissue (Petray et al. 2002). Moreover, CCL3/OKR/3 /fghkekgpv" o keg" kphgevgf" ykvj" v jg" Dtc |kn" *T. cruzi* strain contrasting with wild-type mice presented normal ectfke" cevxkv{. " uwi i guvki" v jcv" EEN51OKR/3 " rnc{u" an important role in cardiac remodelling during infection (Durand et al. 2006). Our recent data indicate that EEN51OKR/3 " eqpvtqu" o cetqr jci g" cpf" EF:⁺ T-cell, but not CD4⁺ T-cell, migration to the heart tissue, particularly during the chronic phase. In addition, CCL3/OKR/3 " kphnwgpegu" EF:⁺CCR5⁺ T-cell compartmentalisation, which may explain the reduced migration of these cells into the cardiac tissue. Our data also indicate that EEN51OKR/3 " eqpvtqu" ectfke" rctcukvku o " cpf" cardiomyocyte lesion formation (Silverio et al. 2009). Ogcpy j kng. " EEN61OKR/3 " o c{ " rtgxgpv" gzequuxg" kpf" inflammation and *T. cruzi* pathogenesis, as indicated by a recent study showing that immunisation of animals with c" EEN61OKR/3 " FPC" xceekpg" ngf" vq" v jg" rtqfwevkq" qh" antibodies that exacerbated cardiac inflammation and fibrosis without affecting parasitism (Roffê et al. 2006).

Elevated plasma chemokine levels have been correlated with worse disease in patients with heart failure (Filippatos & Kardaras 2002). For example, high levels of EEN41OER/3. " dwv" pqv" EEN51OKR/3 " ctg" fktgevn{ correlated with heart dysfunction in CCC (Talvani et al. 2004a). Since CCC progression is probably multifactorial, involving an interplay between host genetics, the parasite, and environmental factors, characterisation of host genetic predisposition may help to identify risk factors for disease progression. For example, a study of a CCL2/MCP-1 single nucleotide polymorphism at the -2518A/G position revealed that subjects with the CCL2/MCP-1-2518AA genotype have a 4-fold greater risk of developing CCC than those without this genotype (Ramasawmy et al. 2006). Although the authors proposed that the CCL2/MCP-1 variant A, correlated with a low transcriptional level, is a genetic modifier of clinical outcome for *T. cruzi* infection, the CCL2/MCP-1 status of these patients and the participation of CCL2/MCP-1 in CCC progression remain to be clarified. CCL2/MCP-1 signals exclusively through CCR2, which also recognises CCL8/MCP-2, CCL7/MCP-3 and CCL13/MCP-4 (Viola & Luster 2008). CCR2-deficient mice infected with the Colombian *T. cruzi* strain exhibit a transient increase in heart parasitism but not parasitaemia during acute infection. Neither the intensity of their cardiac inflammation nor the

levels differs from those of wild-type mice (Hardison et al. 2006a). Conversely, CCL2^{-/-} mice infected with the Y strain of *T. cruzi* are more susceptible to infection, presenting higher parasitaemia and parasitism.

In CCL2/MCP-1-deficient mice, macrophages display reduced trypanocidal activity and CCL2/MCP-1 has been shown to be important for macrophage and CD8⁺ T-cell activation, cell migration to the cardiac tissue and inflammatory cell retention in focal infiltrates (CN Paiva et al., unpublished observations). Myocarditis intensity in acutely and chronically Colombian strain-infected C3H/He mice is thus directly correlated with the cardiac concentration of the CC-chemokine CCL2/MCP-1 (our unpublished observations). Together, these data indicate that the effects of CC-chemokines on the differential migration of distinct cell populations into the cardiac tissue and effector activity of the infiltrating inflammatory cells during *T. cruzi* infection should be further explored.

Chemokine receptor binding leads to integrin activation, encouraging firm leukocyte adhesion to the activated endothelium and extravasation into the inflamed tissues (Viola & Luster 2008). Thus, characterisation of chemokine receptor expression on circulating leukocytes may foster better comprehension of the roles of different cell populations in tissue injury. Peripheral blood mononuclear cells (PBMCs) of chagasic patients tend to express CCR5 and CXCR4 at a higher frequency than those of non-infected individuals (Talvani et al. 2004b). A high frequency of circulating CCR5⁺ lymphocytes, which were mainly CD8⁺, was also detected in acutely *T. cruzi*-infected mice (Marino et al. 2004). Interestingly, this increase in CCR5⁺ cells among the PBMC population occurs 14 days post-infection (Silva et al. 2007), coinciding with augmentation of LFA-1^{high}VAL4^{high} activated PBMC numbers and myocardium colonisation by inflammatory cells (Marino et al. 2003). Studies of CCR5 59029A/G gene polymorphism in Peruvian (Calzada et al. 2001) and Venezuelan patients (Fernandez-Mestre et al. 2004) revealed that the G allele, which reduces CCR5 expression, is found more frequently in asymptomatic patients than in cardiopatics. Together, these data suggest that CCR5 may be linked to CCC severity. In vitro stimulation of PBMCs from CCC patients with *T. cruzi* antigens also enhances the frequency of CD4⁺ and CD8⁺ V α 24⁺ T lymphocytes (EET7|KHP/ " EZET5|KHP/ " cpf" EZET5|VPH" * I q o gu" gv" cn|" 4227-4). Meanwhile, asymptomatic patients had an increased frequency of CD4⁺ T and CD8⁺ lymphocytes co-expressing CCR3/IL-10 and CCR3/IL-4. Therefore, CCC genesis may be correlated with the migration of cells bearing chemokine receptors and expressing specific effector functions from the peripheral blood to the cardiac tissue (Lannes-Vieira 2003). In the cardiac tissue of Colombian strain-infected C3H/He mice, there is a predominance of CD8⁺ CCR5⁺ T lymphocytes relative to CCR5⁺CD4⁺ T cells (Marino et al. 2004). To modulate heart inflammation, mice were treated with N-terminal-methionylated RANTES (Met-RANTES), a selective partial antagonist that acts on CCR1 and CCR5. This drug did not interfere with heart parasitism but significantly reduced the

numbers of CD4⁺, CD8⁺ and CCR5⁺ lymphocytes in the tissue. Furthermore, decreased fibronectin deposition paralleled an increase in the survival of treated animals. Pqvcn{."v jg"pw o dgt"qh"KHP/ + cells in the heart was similar in saline-injected and Met-RANTES-treated mice, while the IL-4⁺ cell count was drastically reduced in the treated mice (Marino et al. 2004). The cardiac tissue of the Met-RANTES-treated mice also exhibited lower TNF and higher IL-10 levels (Fig. 3), reflecting a more balanced cytokine milieu. These data strongly suggest that while CCR5⁺ cells are not crucial for parasite control, they may play an important role in the pathogenesis of chagasic cardiomyopathy. Importantly, Met-RANTES administration also demonstrated that *T. cruzi*-elicited acute meningoencephalitis is a CCR5/CCR1-independent process (Silva et al. 2007), supporting CCR5 targeting as a safe potential therapeutic strategy for modulating chagasic myocarditis.

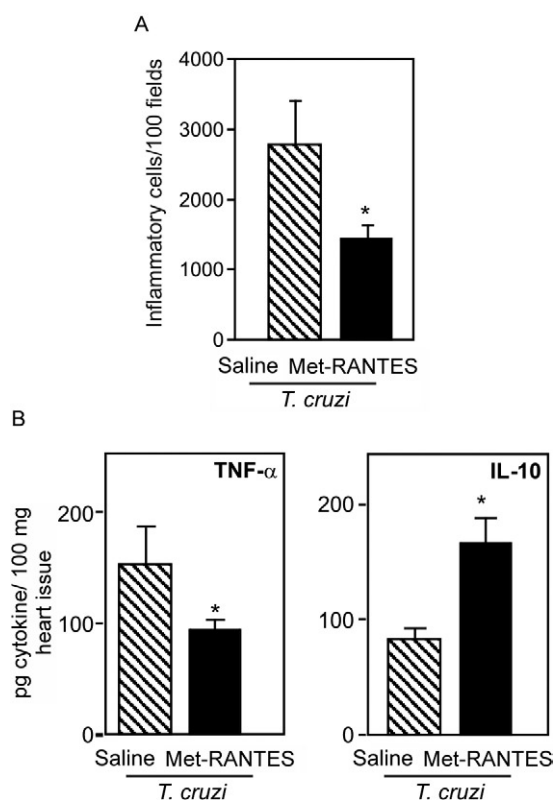


Fig. 3: total inflammatory mononuclear cells in the cardiac tissue of infected mice and concentration of the cytokines TNF and IL-10 in heart extracts of *T. cruzi*-infected C3H/He mice treated daily from 14-28 dpi with saline (striped bar) or methionylated-Regulation of Activation, Normal T-cell Expressed and Secreted (Met-RANTES) [dncem" dct-" 32" i" qh" Ogv/TC PVGUl o qwug" cu" fguetldgf" kp" Octkpq" et al. (2004)]. A: numbers of total inflammatory mononuclear cells (CD4⁺ and CD8⁺ T cells as well as macrophages) in the cardiac tissue of infected mice treated daily with saline or Met-RANTES (Medeiros et al. 2009); B: detection of TNF and IL-10 in heart extracts of infected mice treated daily with saline or Met-RANTES (Medeiros et al. 2009). Each bar represents the mean \pm SD of experimental group composed of eight mice. The data represent two independent experiments. *: $p < 0.05$.

The predominance of CD8⁺CCR5⁺ cells in the heart tissue of *T. cruzi*-infected animals was corroborated using C57BL/6 mice infected with the Y *T. cruzi* strain (Machado et al. 2005), indicating that CD8⁺CCR5⁺ cell prevalence results from *T. cruzi* infection and is not an artefact of a particular model. Increased CCR5 expression by CD8⁺ T cells was detected after in vitro stimulation with a *T. cruzi* antigen (Machado et al. 2005), demonstrating the central role of the parasite in CCR5 modulation. We have also recently demonstrated that TNF signalling via TNFR1 is crucial for CD8⁺CCR5⁺ differentiation, CD8⁺CD44⁺CD62L^{-low} compartmentalisation and CD8⁺ T cell-enriched myocarditis formation during *T. cruzi* infection (Kroll-Palhares et al. 2008).

CCR5-deficient mice present an intense decrease in macrophage and T-cell migration to the heart, coupled with high parasitaemia and cardiac parasitism, especially during early infection (Hardison et al. 2006b). We recently confirmed that CCR5 is not requisite to the acute-phase survival of Colombian strain-infected mice (NF Vinagre et al., unpublished observations), supporting our proposal that CCR5 may be a rational target for immunoregulation of *T. cruzi* infection.

The intensity of chronic *T. cruzi*-elicited myocarditis is related to the concentrations of the CC-chemokines EEN51OKR/3 " cpf" EEN71TCPVGU" kp" v jg" jgctv" vku-sue and to the frequency of peripheral CCR5⁺LFA-1⁺ T lymphocytes (Medeiros et al. 2009). Given the lack of therapeutic prospects for chronic *T. cruzi*-elicited cardiac inflammation and our promising results using Met-RANTES to treat acute infection (Marino et al. 2004), we administered this partial CCR1/CCR5 antagonist to chronically Colombian strain-infected mice. This treatment led to a 20-30% decrease in CD4⁺ cell numbers, reduced IL-10, IL-13 and TNF expression and impaired re-compartmentalisation of activated CD4⁺CCR5⁺ lymphocytes. Notably, a beneficial effect of CCR1/CCR5 blockage during chronic infection was revealed by protection against connexin 43 loss in the heart tissue and creatine kinase cardiac isoenzyme MB (CK-MB) level enhancement, both of which are typically markers of heart dysfunction (Medeiros et al. 2009).

Trafficking of immunocompetent CD8⁺ T cells to the cardiac tissue contributes to chronic *T. cruzi*-elicited myocarditis in chagasic patients (Reis et al. 1993, Higu-chi et al. 1997) and experimental models (dos Santos et al. 2001, Medeiros et al. 2009). There is also a correlation between CD8⁺ T cell numbers and parasite persistence in the heart tissue of chagasic patients (Reis et al. 1997). Emerging evidence suggests that CD8⁺ effector-cell subpopulations with varied cytotoxic and pro-inflammatory *KHP/ + abilities may determine the outcome of an inflammatory process (Sandberg et al. 2001). For instance, there is a sensible correlation between the numbers of KHP/ + and TNF⁺ cells and CD8⁺ T cells and successful parasite control in CCC patients (Reis et al. 1997). In contrast, the presence of heart-infiltrating, granzyme A-expressing CD8⁺ cytotoxic T lymphocytes is associated with severe CCC (Reis et al. 1993). In this vein, our recent data showed that the number of perforin⁺ and KHP/ + cells invading the cardiac tissue of *T. cruzi*-infected

ed mice was associated with CK-MB release and electrical abnormalities as infection progressed. Although the perforin-dependent pathway played a role in anti-*T. cruzi* immunity and acute-phase survival, it was not essential when mice were infected with a low inoculum of the Colombian strain. Moreover, perforin deficiency resulted in lower CK-MB isoenzyme activity in the serum and restricted connexin 43 loss, both markers of cardiac lesions and hampered the development of electrical abnormalities. *Vjg"pw o dgtu"qh" KHP/ /rtqfwepi" egmmu" ygtg" cnuq"* enhanced in perforin-deficient animals, suggesting that this cell population may play a protective role in *T. cruzi*-induced cardiomyopathy (Silverio et al. 2009).

The data discussed above support the idea that an appropriately intense and focussed immune response can control *T. cruzi* dissemination in the absence of disease, as observed in patients presenting an indeterminate form of Chagas disease. Hence, a thorough understanding of the biological forces governing leukocyte trafficking to *T. cruzi*-infected heart tissue, such as cytotoxic or inflammatory CD8⁺ T cells, may facilitate the design of treatments that regulate CCC without hampering parasite control. Currently, therapies using antibodies to and antagonists of CAM and chemokine receptors are being developed to selectively curb chronic inflammation in degenerative diseases (Kummer & Ginsberg 2006, Ley et al. 2007, Viola & Luster 2008, Horuk 2009). Therefore, the application of these strategies in parasite-triggered chronic inflammation, such as Chagas disease, to regain homeostatic balance in parasite-host interactions is an increasingly rational, attractive and realistic possibility.

ACKNOWLEDGMENTS

To Dr. Ricardo Lourenço (Editor of the Memórias do Instituto Oswaldo Cruz), for authorizing the reproduction of the original drawing made by Castro Silva (Chagas 1919), reproduced in Fig. 1.

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