

Chronic indeterminate phase of Chagas' disease: mitochondrial involvement in infection with two strains

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SUMMARY

Chagasic cardiopathy has become one of the most frequent causes of heart failure and sudden death, as well as one of the most common causes of cardio-embolic stroke in Latin America. The myocyte response to oxidative stress involves the progression of cellular changes, primarily targeting the mitochondria and modifying therefore the energy supply. In this paper we analysed the effect of the infection of mice with 2 different strains of *Trypanosoma cruzi* (Tulahuen and SGO Z12) in the chronic indeterminate stage (75 days post-infection), upon the structure and function of cardiac mitochondria. The structural results showed that 83% of the mitochondria from the Tulahuen-infected mice presented an increase in their matrix and 91% of the mitochondria from the SGO Z12-infected group showed a reduction in their diameter ($P < 0.05$). When the Krebs cycle and mitochondrial respiratory chain functionality was analysed through the measurement of the citrate synthase and complexes I to IV activity, it showed that their activity was altered in all cases in a similar manner in both infected groups. In this paper we have demonstrated that the chronic indeterminate phase is not 'silent' and that cardiac mitochondria are clearly involved in the genesis and progression to the chronic chagasic cardiopathy when different factors alter the host-parasite equilibrium.

Key words: Chronic indeterminate Chagas' disease, *Trypanosoma cruzi* strains, mitochondria, respiratory chain, myocardiopathy.

INTRODUCTION

One of the most important determinants of congestive heart failure and sudden death in Latin America is Chagas disease, provoked by infection with the intracellular protozoan parasite *Trypanosoma cruzi*. It affects approximately 20 million people (World Health Organization, 2007; Moncayo and Silveira, 2009) and represents a serious public health problem in Central and South America (Biolo *et al.* 2010). Chagasic cardiopathy appears to carry the worst prognosis (Pereira *et al.* 2009) and has become the most frequent cause of heart failure and sudden death, as well as the most common cause of cardio-embolic stroke in Latin America. Chagas disease also represents an increasing challenge for clinicians in the United States (Bern *et al.* 2007) and some European countries (Reesink, 2005) due to the continuous immigration of people from disease-endemic countries (Polo-Romero *et al.* 2011).

There are 3 stages in the evolution of Chagas disease: the acute phase with a local inflammatory lesion that appears at the site where the metacyclic trypomastigotes entered, and the parasite spreads throughout the host organism (Prata, 2001; Umezawa *et al.* 2000); the cardiac chronic phase in which the diversity and severity of the symptoms range from a mild electrocardiographic alteration to sudden death due to cardiac dysrhythmias, varying in different patients and in different regions (Storino and Milei, 1994). Between the acute and the cardiac chronic phases exists a period called the chronic indeterminate stage, which is generally symptomless and may last for 10 to 20 years (Macedo, 1999; Ribeiro and Rocha, 1998). More sensitive tests have demonstrated that patients in this stage may present significant abnormalities (Ribeiro and Rocha, 1998). According to cardiologists, this period could be the key to recognizing which patients will develop the cardiac form of infection (Elizari, 1999).

Trypanosoma cruzi invasion and replication in the cardiomyocytes provoke cellular damage and a cytotoxic reaction with inflammatory cytokines and nitric oxide production, both of them being a source of reactive oxygen species (ROS) in the acute

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(Cardoni *et al.* 1997) and cardiac chronic phases of the infection (Talvani *et al.* 2004). A similar inflammatory response has been described for the chronic indeterminate phase. These responses are triggered to control parasite reproduction, but it may also have toxic effects upon host cellular components (Ueda *et al.* 2002). The myocyte response to oxidative stress involves the progression of cellular changes, primarily targeting the mitochondria (Long *et al.* 2004) and modifying therefore the energy supply; this bioenergetic dysfunction could be involved in the genesis and progression of heart failure (Marin Garcia and Goldenthal, 2008; Tsutsui, 2006).

Previous work from our laboratory (Báez *et al.* 2008, 2011), and from other authors (Garg *et al.* 2003; Vyatkina *et al.* 2004), have demonstrated different structural and functional alterations in cardiac mitochondria isolated from mice infected with different *T. cruzi* strains in the acute and cardiac chronic stages.

Positive serological reactions for *T. cruzi*, normal thoracic radiography and normal electrocardiographic tracings are the elements used for the characterization of the chronic indeterminate form of Chagas disease. But the study of this long silent period of the disease, with more sensitive, invasive or non-invasive, diagnostic methods, demonstrates that most chagasic patients with normal electrocardiograms have some type of anatomical or functional alteration. Taking this into account, our current study investigates whether the structure and function of cardiac mitochondria are altered as a result of the infection with different *T. cruzi* strains, in order to determine their involvement in the pathophysiological mechanism of chronic chagasic myocardiopathy. These studies could help to elucidate which patient will develop a chagasic cardiopathy.

MATERIALS AND METHODS

Infection

Albino Swiss female and male mice weighing 30 ± 1 g ($n=90$) were used as follows: 30 mice were inoculated, by intraperitoneal injection, with 50 trypomastigote forms of *T. cruzi*, Tulahuen strain (Tulahuen), and 30 mice with 50 trypomastigote forms of the SGO Z12 isolate (SGO Z12), which was obtained from a patient from an endemic area. The number of parasites/ml of blood was determined in each group using a Neubauer haemocytometer. A non-infected group ($n=30$) was also studied.

Parasitaemias in all groups were determined in a Neubauer haemocytometer using blood samples obtained from the tail of the mice, twice a week, beginning 7 days after the infection until 75 days post-infection (chronic indeterminate stage).

The investigation was carried out according to the Guide for the Care and Use of Laboratory Animals

published by the US National Institute of Health, NIH publication (No. 85-23, revised 1996).

Serology

Blood samples were collected 50, 90, and 135 days post-infection, representing the acute, intermediate, and chronic phases of the experimental infection, respectively. These samples were then assayed for antibodies to 'cruzipain' antigen (see below), by ELISA (Gea *et al.* 1992).

Trypanosoma cruzi antigen

Epimastigote forms of *T. cruzi* SGO Z12 isolate were grown at 26 °C in liver digest neutralized medium (Oxoid America, Columbia, USA) supplemented with 0.5% tryptose, 10% FCS, 200 mg/ml hemin, 100 U/ml streptomycin. Parasites were harvested at the exponential growth phase, centrifuged at 5000 *g* for 10 min, and washed with PBS. One millimolar *N*- α -*p*-tosyl-L-lysine chloro methyl ketone (TLCK) was added as a cysteine protease inhibitor and the parasite pellet was lysed by 8 cycles of freezing and thawing. The parasite homogenate was centrifuged at 105 000 *g* and the supernatant was used for cruzipain purification. Cruzipain was purified by affinity chromatography as previously described (Labriola *et al.* 1993). The absence of enzyme activity was controlled in gels containing 0.1% gelatin (Sigma) incubated after the PAGE run at pH 5.7 and stained with Coomassie blue R250; the samples were neither reduced nor boiled.

Mitochondria isolation

Hearts were removed on day 75 post-infection (p.i.), which corresponds to the chronic indeterminate phase of the experimental infection (Lo Presti *et al.* 2008), obtaining both ventricles. They were washed and suspended in ice-cold isolation buffer (5 mM HEPES, pH 7.2, containing 210 mM mannitol, 70 mM sucrose, 1 mM EGTA, and 0.5% BSA (fatty acid-free), tissue/buffer ratio, 1:10 w/v) and immediately homogenized. Homogenates were centrifuged at 1500 *g* at 4 °C for 20 min and the supernatant transferred to a new tube. The pellet was re-suspended in isolation buffer, homogenized, and centrifuged again at 10 000 *g*, 4 °C for 5 min. The supernatant was discarded and the pellet was resuspended in buffer and centrifuged at 10 000 *g* at 4 °C for 10 min (twice = purification). The mitochondrial pellet was re-suspended in isolation buffer (tissue/buffer, 1:1 ratio, w/v), and the aliquots stored at -80 °C.

Respiratory complex and citrate synthase activity

The activities of the respiratory complexes (CI–CIV) and citrate synthase were monitored by spectrophotometric methods as previously described (Trounce *et al.* 1996; Jarreta *et al.* 2000; Vyatkina *et al.* 2004; Báez *et al.* 2008, 2011) with slight modifications. Protein concentrations were determined by Bradford assay (Bradford, 1976).

All assays were performed in 1 ml final volume with 30–40 mg (for complexes I and II), 20–30 mg (for complex III) and 15 mg (for complex IV) of mitochondrial protein, and the linear change in absorbance was measured for 3 min.

CI (NADH-ubiquinone oxidoreductase). The reaction mixture consisted of 10 mM Tris–HCl buffer, pH 8.0, 80 μ M 2,3-dimethoxy-5-methyl-6-decyl-1,4-benzoquinone (DB), 1 mg/ml BSA, 0.25 mM KCN. After incubating the mitochondria in the reaction mixture at 30 °C for 10 min., oxidation of NADH (200 μ M) was monitored at 340 nm (ϵ 8 mM⁻¹ cm⁻¹).

CII (succinate-ubiquinone oxidoreductase). Mitochondria were incubated in 1 M potassium phosphate buffer, pH 7.0, containing 0.1 ml succinate phosphate 0.1 M. After addition of assay mixture consisting of 50 μ M 2,6-dichlorophenolindophenol (DCPIP), 5 μ l EDTA 1 mM, 10 μ l of Triton X-100 1%; 50 μ l of DB. All the components were mixed. The reduction of DCPIP in association with CII-catalysed DB reduction was measured at 600 nm (ϵ 20.5 \times 10⁶ M⁻¹ cm⁻¹).

CIII (ubiquinol-cytochrome c oxidoreductase). Mitochondria were suspended in 50 mM Tris–HCl buffer, pH 7.4, containing 1 mM EDTA, 250 mM sucrose, 2 mM KCN and 50 μ M oxidized cytochrome c. After the addition of 80 μ M reduced DB (DBH₂), the reduction of cytochrome c was measured at 550 nm (ϵ 19.0 mM⁻¹ cm⁻¹).

CIV (cytochrome c oxidase). Mitochondria (2 μ g protein) were permeabilized in 10 mM Tris–HCl, pH 7.0, 25 mM sucrose, 120 mM KCl, and 0.025% n-dodecyl- β -D-maltoside, and 50 μ M reduced cytochrome c added. The oxidation of cytochrome c was measured at 550 nm (ϵ 19.0 mM⁻¹ cm⁻¹).

Citrate synthase. The mitochondrial pellet was added to 100 mM Tris–HCl buffer, pH 8, 0.3 mM acetyl CoA, 100 μ M 5,5-dithio-bis-2-nitrobenzoic acid (DTNB). The reaction was initiated by 0.5 mM oxaloacetate. Citrate synthase-catalysed reduction of acetyl CoA with oxaloacetate in conjunction with DTNB reduction was monitored at 412 nm (ϵ 13.6 mM⁻¹ cm⁻¹).

Histopathological studies

The hearts were removed from the infected mice on day 75 p.i., fixed in buffered (pH 7.0) 10% formaldehyde, and embedded in paraffin. Each heart was cut horizontally into 5 μ m sections from the apex to the auricles. The sections were stained with haematoxylin–eosin. In total 50 slices from each group were analysed with a 40 \times objective.

Parasite detection

For parasite detection in cardiac tissues, polymerase chain reaction (PCR) was performed for 40 cycles, using *T. cruzi*-specific oligonucleotides (Tcz1: 5'-CGAGCTCTTGCCCACACGGGTGCT-3' and Tcz2: 5'-CCTCCAAGCAGCGGATAGTTTCAGG-3') with 2.5 μ l of the total DNA as template. Denaturation, annealing, and elongation steps were performed for 30 sec at 95 °C, 30 sec at 60 °C, and 30 sec at 72 °C, respectively. A 10 μ l aliquot of each PCR reaction was resolved on a 1.2% agarose gel. The ethidium bromide-stained gels were visualized using long-wave UV light.

Electron microscopy studies

A 1 millimeter square section, from the tip of the left ventricle, was fixed immediately after extraction in a Karnovsky solution (4% formaldehyde and 1.5% glutaraldehyde in 0.1 M cacodylate buffer) for at least 2 h at room temperature. Then the tissues were washed 3 times in cacodylate buffer and post-fixed in 1% osmium tetroxide for 1–2 h. After dehydration in a graded acetone solution (50%, 70% and 90%), the inclusion of samples was performed in an Epoxy composite mixture of Araldite 506 (48.5%), dodecenyl succinic anhydride (48.5%), dibutyl phthalate (0.5%) and dimethyl aminobenzene accelerator (2.5%). Ultrathin cuts were stained with uranyl acetate and lead citrate and examined in a Zeiss electron microscope. In order to evaluate the changes on mitochondrial morphology observed in the different experimental groups (5 micrographs for each mouse), a 4-grade classification was used as follows. Grade 1: normal structure; Grade 2: normal size with dilated cristae; Grade 3: normal size and/or altered shape. Intact membrane with few cristae; Grade 4: mitochondrial swelling.

The 3-dimensional studies were carried out using the Femtoscan program. FemtoScan Online software analysed 5 micrographs for each mouse to build images from electron microscopy. The program runs on operating systems Windows XP, Vista, Windows 7, as well as their server analogues – Windows Server 2003, 2003 R2, 2008, 2008 R2.

Statistical analysis

Results are shown as mean \pm standard error. The data obtained were analysed by ANOVA (Fisher test),

Table 1. Mitochondrial enzymatic activity (η moles $\text{min}^{-1}/\text{mg}$ of prot) of hearts obtained from uninfected ($n=10$), Tulahuen infected ($n=10$) and SGO Z12 infected mice ($n=10$) at 75 days post-infection from different groups under study

(The results are expressed as mean \pm standard error unless otherwise indicated.)

Actividad Enzimática (η mol/min.mg de prot)	Uninfected	Tulahuen 75 dpi	SGO Z12 75 dpi	P-value
Citrate synthase	290 \pm 40 (B)	10 \pm 2.7 (A)	40 \pm 20 (A)	B vs A 0.0001
Complex I	50 \pm 20 (B)	3.9 \pm 0.43 (A)	10 \pm 1.1 (A)	B vs A 0.05
Complex II	1 $\times 10^{-6} \pm 0.00$ (A)	9.2 $\times 10^{-5} \pm 9.8 \times 10^{-6}$ (B)	1.4 $\times 10^{-4} \pm 1.09 \times 10^{-5}$ (C)	A vs B vs C 0.05
Complex III	180 \pm 20 (C)	10 \pm 0.44 (A)	90 \pm 10 (B)	C vs A vs B 0.001
Complex IV	290 \pm 30 (A)	280 \pm 30 (A)	650 \pm 30 (B)	A vs B 0.001
Comparison of all variables (Hotelling's test corrected by Bonferroni)	(C)	(A)	(B)	C vs A vs B 0.0001

MANOVA (Hotelling corrected by Bonferroni test) and Chi² test for categorical variables. Axiovision 3.0 software was used to quantify mitochondria. The significance level was set at $P < 0.05$ for all cases.

RESULTS

Detection of parasites in blood and cardiac tissue

Parasitaemias were analysed to assess the infection rate. Both infected groups showed the highest parasitaemia levels by day 14 p.i., with the levels of the SGO Z12 group being significantly lower ($P < 0.01$) than those presented by the Tulahuen group (87.58 ± 7.64 and 225.39 ± 19.11 parasites/ μL , respectively). Parasitaemias were negative from day 42 onwards in the Tulahuen-infected mice and from day 49 onwards in the SGO Z12-infected mice.

The PCR analyses carried out in cardiac tissues were positive in Tulahuen and SGO Z12 infected groups.

Serology

CI was positive at 50, 90 and 135 days post-infection.

Mitochondrial respiratory complexes and citrate synthase activity

In order to determine the mitochondrial function, we analysed the enzymatic activity of the mitochondrial respiratory chain complexes I to IV (cristae) and citrate synthase (matrix) in the myocardium from uninfected and Tulahuen- and SGO Z12-infected mice on day 75 p.i. As can be observed in Table 1, citrate synthase enzyme activity from Tulahuen and SGO Z12 mice was significantly decreased compared to the control group ($P < 0.0001$); there were no significant differences between parasite strains.

CI complex activity significantly decreased in both experimental groups, $P < 0.05$ when compared with

the control. CII complex enzyme activity increased in both infected groups 75 days p.i. ($P < 0.05$) but the increment was higher in the SGO Z12-infected mice than in the Tulahuen mice. As shown in Table 1, CIII complex enzyme activity significantly diminished with infection with either parasite strain, but the Tulahuen group presented a greater reduction in its activity ($P < 0.001$). The CIV complex showed an increase in its specific activity in chronic indeterminate infection in the SGO Z12-infected group ($P < 0.001$).

Cardiac histopathological studies and mitochondrial ultrastructural analysis during chronic indeterminate T. cruzi infection

In the chronic indeterminate stage of the infection, the Tulahuen strain-infected group showed lymphomonocitary infiltrates (Fig. 1 D). Hearts from mice infected with SGO Z12 isolate presented few inflammatory infiltrates (Fig. 1 G). Sections with no cardiac alterations were also observed in this latter group.

At this time, 83% of the mitochondria from the Tulahuen-infected mice presented at least 1 significant abnormality such as an increase in their matrix or disorganization in their cristae (Fig. 1 E, F) when compared with the non-infected group (Fig. 1 B, C). The average area of mitochondria from this group was greater than the control (Table 2). Some alterations of different importance were detected in the 91% of the mitochondria from the SGO Z12-infected group, showing a considerable reduction in their area ($P < 0.01$); a greater number of mitochondria was also found (see Fig. 1H and I and Table 2); cristae dilatation and increase in mitochondrial matrix, can be observed in the figures (see Fig 1 E, F, H and I).

Figure 2 shows a 3-dimensional view of cardiac mitochondria obtained from non-infected and

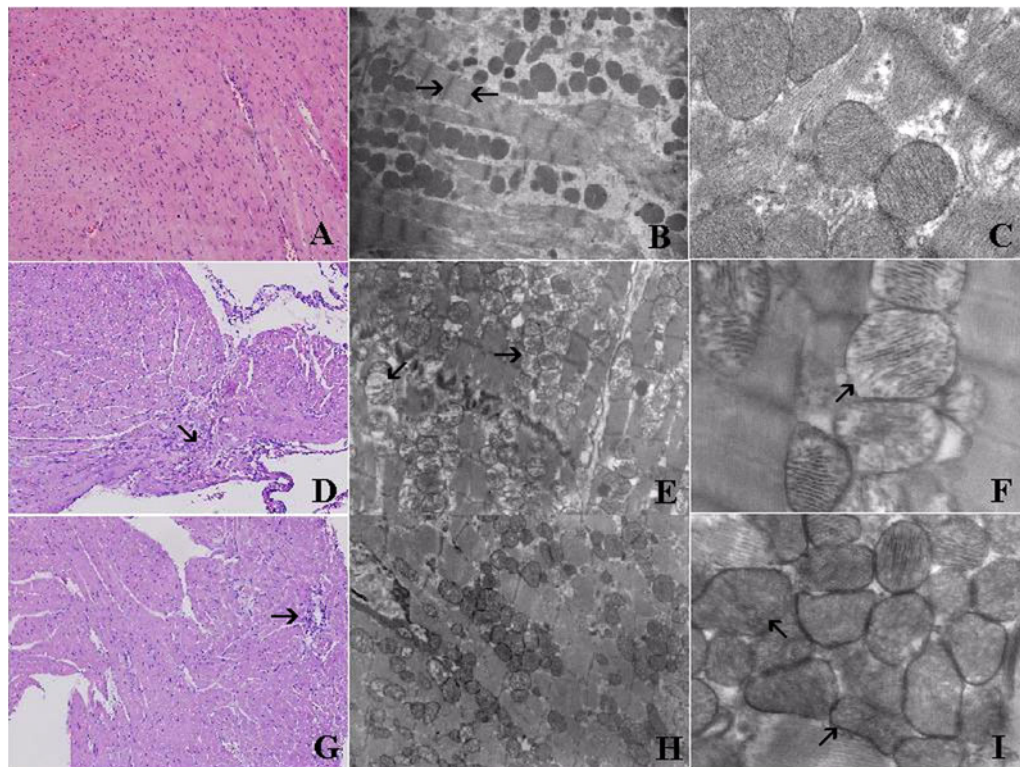


Fig. 1 (A) Cardiac histological sections from non-infected mice, 200 \times . (B) Cardiac ultrastructure from non-infected mice where sarcomere and mitochondria can be observed, 6000 \times . (C) Cardiac ultrastructure from non-infected group where mitochondria can clearly be observed, 27 800 \times , bar: 2.2 micrometer. (D) Cardiac histological sections from the Tulahuen-infected group (50 trypomastigotes of *Trypanosoma cruzi*/animal), 75 days post-infection showing cell infiltrate, 200 \times . (E) Ultrastructure of cardiac mitochondria from the same group showing separate cristae with increase of matrix volume, 6000 \times . (F) Ultrastructure of cardiac mitochondria from the Tulahuen-infected group, 75 days post-infection; increase of matrix can be observed (dilation), 27 800 \times . (G) Hearts from the isolate SGO Z12-infected group (50 trypomastigotes of *T. cruzi*/animal), 75 days post-infection showing mononuclear cell infiltrates, 200 \times . (H) Ultrastructure of cardiac tissue from the isolate SGO Z12-infected group, 75 days post-infection showing mitochondria 6000 \times . (I) Ultrastructure of cardiac tissue from the same group showing mitochondria with different morphology and increase of matrix volume, 27 800 \times .

SGO Z12- and Tulahuen-infected mice. In this figure an increase in the matrix and disorganization of the cristae can be observed in both experimental groups.

DISCUSSION

The globalization of Chagas disease, due to migration from endemic countries to North America, Europe, Asia and Oceania has created a new epidemiological, economic, social and political concern in non-endemic countries (Schummis, 2007; Coura and Borges-Pereira, 2010). The main issue in non-endemic countries is the risk of transfusion and congenital transmission, while infected triatomines are still the common way of transmission in endemic countries.

The determinants of Chagas disease come from several, complex and different factors: quantity of parasites that provoke the infection, *T. cruzi* lineages, specific histotropic receptors of the host and patient's initial immune response, among others (Coura, 1988;

Andrade *et al.* 2006; Bustamante *et al.* 2003; Zingales *et al.* 2009); despite this, all infected individuals present a symptom or symptomless acute phase that will evolve into the chronic indeterminate phase.

In this asymptomatic chronic indeterminate stage no anatomo-pathological manifestations are found, except for isolated inflammatory foci with mild reduction of cardiac neurons, which are not enough to provoke clinical manifestations. In the present paper, myocardium samples from Tulahuen- and SGO Z12-infected mice presented cardiac isolated inflammatory infiltrates and absence of amastigote nests. Even though these isolated inflammatory infiltrates described are the result of *T. cruzi* persistence in host tissues by successful evasive strategies used by the parasite: it releases proteases that activate TGF- β (Araujo Jorge *et al.* 2008; Waghabi *et al.* 2009; Maya *et al.* 2010). Additionally, phagocytosis and apoptotic bodies originated from T cells or neutrophils induce a prostaglandin-dependent production of TGF- β in macrophages. All this provokes a diminished nitric oxide production and the INF- γ

Table 2. Results of the ultrastructural analysis of cardiac mitochondria of uninfected ($n=5$ micrographs/mouse), Tulahuen infected ($n=5$ micrographs/mouse) and SGO Z12 infected ($n=5$ micrographs/mouse) mice at 75 days post infection

(The results are expressed as mean \pm standard error. Different letters mean statistically significant differences, $P<0.05$ within each row.)

Measurements	Uninfected ($n=5$)	Tulahuen 75 dpi ($n=5$)	Z12 75 dpi ($n=5$)
Total area occupied by mitochondria	36152.02 μm^2	40885.56 μm^2	49482.57 μm^2
Area average mitochondria	903.80 \pm 76.48 μm^2 (b)	1202.51 \pm 96.18 μm^2 (c)	494.83 \pm 32.44 μm^2 (a)
Grade Alteration			
	80% grade 1	17% grade 1	9% grade 1
	20% grade 2	32% grade 2	17% grade 2
	0% grade 3	37% grade 3	43% grade 3
	0% grade 4	14% grade 4	31% grade 4
Total number of mitochondria	40	34	100
Average number of mitochondria	8 \pm 2.30	4.20 \pm 0.58	12.00 \pm 1.70

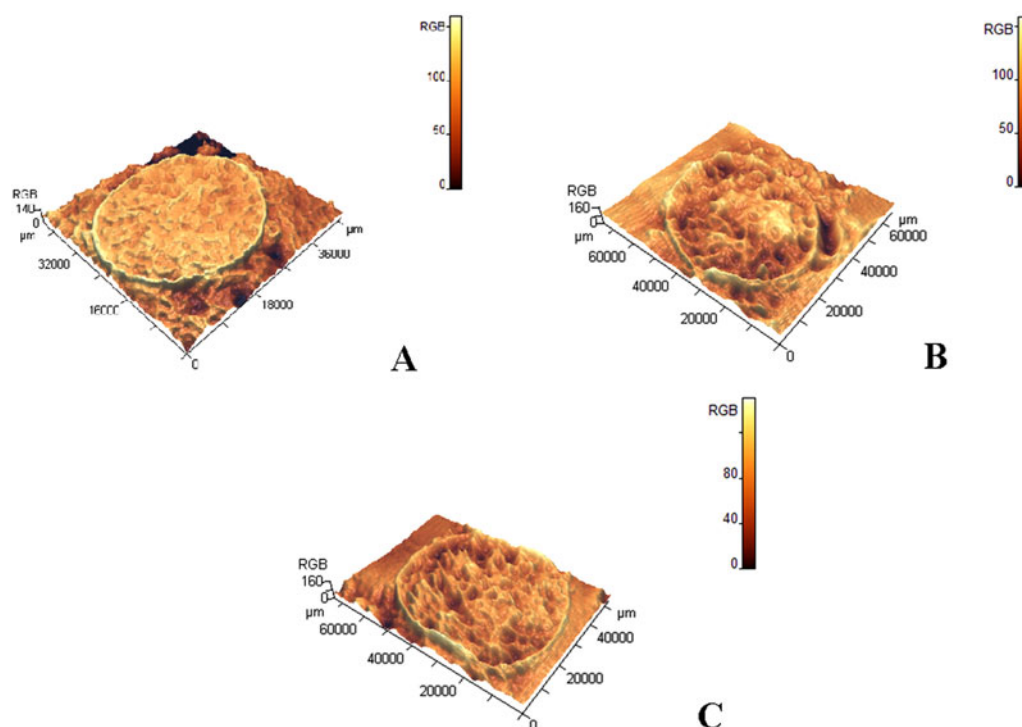


Fig. 2. Three-dimensional view of cardiac mitochondria obtained from non-infected (A), SGOZ12- (B) and Tulahuen- (C) infected mice. We observed an increase both the size of the mitochondria of the infected groups and the matrix. Measurements were made in pixels.

induced inflammatory response is attenuated; consequently, the parasite proliferates (Gutierrez *et al.* 2009). This may be the cause of the low-grade of inflammatory response described in the chronic indeterminate phase, that we also demonstrated in our results. But all of these changes that have no clinical evidence are generating cellular structural damage that can affect cardiac mitochondria either at or near the inflammatory formation (Tsutsui, 2006).

In this paper we analysed the effect of the infection with 2 different strains of *T. cruzi* (Tulahuen and SGO Z12) in this 'clinically silent' stage, upon the

structure and function of cardiac mitochondria. The structural results, using a 4-grade classification of mitochondrial damage, showed that at this time, 83% of the mitochondria from the Tulahuen-infected mice presented at least 1 significant abnormality such as an increase in their matrix or disorganization in their cristae (14% Grade 4) that are probably related to the enzymatic dysfunction.

Alterations of different importance were detected in the 91% of the mitochondria from the SGO Z12-infected group (31% Grade 4) ($P<0.05$). The abnormalities described were corroborated using the 3-dimensional studies (Fig. 2).

The number of mitochondria in cardiac tissue was not modified by the *T. cruzi* infection in the Tulahuen group, showing a significant increase in the average area of the organelle ($P < 0.01$). These results are different from those previously obtained in the cardiac chronic phase (Báez *et al.* 2011) when a clear reduction in the mitochondrial size was described. The SGO Z12-infected group showed a significantly diminished area occupied by mitochondria ($P < 0.01$). The greater number of mitochondria found in this group probably compensates the important number of them with several alterations.

Additionally, in this 'silent' stage of the *T. cruzi* infection we demonstrated functional changes in cardiac mitochondria of different involvement related to the strain that infected the host.

When we studied the Krebs cycle functionality through the measurement of the specific citrate synthase activity, we found it to be significantly diminished, in contrast to the increased activity found in the acute and cardiac chronic stages of the infection (Báez *et al.* 2008, 2011). When the mitochondrial respiratory chain was analysed in the chronic indeterminate stage of the infection through the measurement of the specific activity of complexes I to IV, we found that the activity of CI and CIII were altered in a similar manner in either infected group. CII presented a significant increase in both experimental groups, while CIV incremented its activity in the SGO Z12-infected mice ($P < 0.0001$) while the Tulahuen group presented a conserved activity not different from the control group; this last result may be related to the maintenance of the respiratory chain redox potential efficiency. Additionally, the dysfunction in the citrate synthase activity and respiratory complexes could be achieved via the matrix and cristae disorganization found in both infected groups.

It was recently demonstrated that oxidative stress-induced protein modifications occur in the myocardium of *T. cruzi*-infected experimental animals (Wen and Garg, 2004; Wen *et al.* 2004; Dhiman *et al.* 2008) and in peripheral blood of seropositive chagasic patients (Wen *et al.* 2006; De Oliveira *et al.* 2007). This peripheral oxidative stress was associated with a poor glutathione antioxidant defence and an increased myeloperoxidase activity and ROS production (Dhiman *et al.* 2012). Besides, *T. cruzi* infection provokes a significant fall in mitochondrial membrane potential; the ROS-induced oxidation of mitochondrial membranes may constitute a secondary signal affecting its potential, and consequently the respiratory chain efficiency (Gupta *et al.* 2009).

The inflammatory pathology is not very important during the chronic indeterminate stage by enhancing an antioxidant status, which is beneficial to preserve the cardiac function for a long time; but as it has been demonstrated that the mitochondrial ROS release due to electron transport chain dysfunction and enhanced release of electrons to molecular oxygen is

the primary source of oxidative stress in the heart (Wen and Garg, 2008). During this long stage there is a permanent aggression to the cardiac cells.

Previous studies from our laboratory have demonstrated that the chronic indeterminate phase is not so 'silent' as has been described: modifications of cardiac contractility, alterations of some components of the cardiac β adrenergic system (Enders *et al.* 1995; Lo Presti *et al.* 2008), and electrocardiographic alterations (Bustamante *et al.* 2003) were previously found. Now we have demonstrated that the responsible organelle for cardiac energy supply is also compromised both structurally and functionally.

The hypothesis that the indeterminate chronic phase is a stage of host-parasite equilibrium rather than a process of progressive damage (Prata, 2001) explains the long-term and clinical silence of the chronic indeterminate phase; present results, however, point to the substantial importance of the parasite strain and the different damage that each of them is capable of inducing throughout this silent stage.

The mechanisms involved in the pathogenesis of Chagas disease and its progression to chronic cardiomyopathy are still under intense discussion. We have demonstrated in the present paper that cardiac mitochondria are clearly involved in the genesis and progression to the chronic chagasic cardiopathy when different factors alter the host-parasite equilibrium.

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