

Usefulness of PCR-based assays to assess drug efficacy in Chagas disease chemotherapy: value and limitations

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*One major goal of research on Chagas disease is the development of effective chemotherapy to eliminate the infection from individuals who have not yet developed cardiac and/or digestive disease manifestations. Cure evaluation is the more complex aspect of its treatment, often leading to diverse and controversial results. The absence of reliable methods or a diagnostic gold standard to assess etiologic treatment efficacy still constitutes a major challenge. In an effort to develop more sensitive tools, polymerase chain reaction (PCR)-based assays were introduced to detect low amounts of *Trypanosoma cruzi* DNA in blood samples from chagasic patients, thus improving the diagnosis and follow-up evaluation after chemotherapy. In this article, I review the main problems concerning drug efficacy and criteria used for cure estimation in treated chagasic patients, and the work conducted by different groups on developing PCR methodologies to monitor treatment outcome of congenital infections as well as recent and late chronic *T. cruzi* infections.*

Key words: Chagas disease - chemotherapy - *Trypanosoma cruzi* - molecular diagnosis - cure assessment

Chagas disease is an important cause of end-stage cardiomyopathy in Latin America (Rassi et al. 2000), with around 100 million people exposed to the disease (Coura 2007). In the early 1990s, an estimated 16-18 million individuals were reportedly infected with the kinetoplastid protozoan *Trypanosoma cruzi*, although this figure has recently been revised to around 11 million, revealing an important drop in prevalence (Guzmán-Bracho 2001). Recent surveys indicate that about 200,000 new cases occur yearly in areas where the disease is endemic, representing the third most common parasitic infection worldwide after malaria and schistosomiasis (WHO 2005). The urbanization process in Latin America and immigration trends, however, have led to the disease being diagnosed in non-endemic areas where, even in the absence of the vector, the infection can still be transmitted congenitally, by blood transfusion and by organ transplantation.

In humans, after a short incubation period, an acute phase occurs that, in the absence of specific treatment, persists for about 15-30 days featuring intense parasitaemia, fever and other symptoms. A strong immune response triggers both B and T lymphocytes. In this phase, morbidity and clinical symptoms are directly associated with parasitaemia level. Alternatively, the acute infection may also occur as a non-apparent form with few symptoms. After the acute phase of the disease, the

patients enter into an asymptomatic chronic stage (sub-acute or indeterminate phase) with the appearance of *T. cruzi*-specific antibodies in the blood stream, which lasts throughout life in about two-thirds of the patients. The remaining one-third of chronically infected individuals develop cardiac or digestive complications 10-30 years after the initial infection, at a stage of the disease in which blood and tissue parasites are scarce. About 20-30% of chronically infected individuals will develop a cardiac clinical form expressed by a large range of manifestations, which can lead to heart failure or sudden death in 70% and 30%, respectively, of those patients presenting chagasic cardiomyopathy (Manzullo & Chuit 1999). Approximately 8-10% of the patients develop digestive manifestations characterized by pathological dilatations of variable severity of the oesophagus and colon.

The pathogenesis of chronic Chagas disease is not completely understood. Parasite persistence with inflammatory reactions (Higuchi et al. 1993, Jones et al. 1993, Brandariz et al. 1995, Bellotti et al. 1996) and other alterations of the host's immune system have been implicated in progressive heart damage caused by infection (Kierszenbaum 1999, Machado et al. 2000). For many years, the traditional parasitological methods, such as haemoculture and xenodiagnosis (XD), could not demonstrate the presence of parasites in damaged tissues. Several studies have implicated autoimmune phenomena as the principal mechanism leading to late cardiac injury (Cunha-Neto et al. 1996, Rassi et al. 2001, Pontes de Carvalho et al. 2002, Iwai et al. 2005, Marin-Neto et al. 2007). This hypothesis is based on the apparent absence of parasites in cardiac inflammatory lesions and the presence of anti-self immune responses in chronic Chagas cardiomyopathy patients, caused by either auto-antibodies or autoreactive T cells, derived by molecular

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mimicry between parasite and host antigens (Kalil & Cunha-Neto 1996, Girones et al. 2005). This supposition suggested that etiologic treatment would be of little benefit (Cunha-Neto et al. 2006).

Despite the autoimmune response, the role played by the parasite is critical (Jones et al. 1993, Vago et al. 1996) and thus, anti-parasitic therapy may prevent disease progression in chronic Chagas disease (Urbina & Docampo 2003). The demonstration of *T. cruzi* components in myocardium by more sensitive methods, such as immunohistochemistry and polymerase chain reaction (PCR), suggested that vestiges of parasites are necessary to trigger the inflammatory process (Higuchi et al. 1993, Jones et al. 1993, Bellotti et al. 1996, Lane et al. 1997, Olivares-Villagómez et al. 1998). Jones et al. (1993) also demonstrated that cardiac tissue adjacent to inflammatory infiltrates yielded PCR-amplified *T. cruzi* DNA in patients who died of severe chronic chagasic cardiomyopathy. Similarly, Vago et al. (1996) revealed the presence of parasite DNA in oesophageal tissues of patients with chronic digestive Chagas disease. These findings suggest that heart parasitism is the primary stimulus for the perpetuation of myocardial inflammation and is responsible for increased serological response to the parasite (Levin 1996). It is likely that the autoimmune component of Chagas disease is a parasite-induced disorder; therefore, etiologic treatment should focus on clearance of parasites and not on suppressing a presumed pathogenic anti-self response. The few studies that attempted to demonstrate the effectiveness of etiologic treatment were only performed during the last decade of the 20th century, probably based on molecular biology evidence that, at that time, proved that the parasite remained in targeted tissues meaning that they could no longer attribute the etiopathogeny exclusively to immunological mechanisms (Villa et al. 2007). Observational studies found that benznidazole treatment for patients with chronic Chagas disease delayed or prevented clinical progression of heart damage, but was not believed to be able to eliminate the parasite during the chronic phase of infection (Viotti et al. 1994, Fabbro De Suasnábar et al. 2000, Cançado 2002). Conversely, immunosuppressive treatments aggravated inflammatory responses in both experimental models and humans (Andrade et al. 1987, Silva & Rossi 1990, Bocchi & Fiorelli 2001).

Although the recent advances in vector control in the Southern Cone countries by an initiative of the Pan American Health Organization (PAHO) and the World Health Organization (WHO) have decreased the incidence of new infections (Schofield & Dias 1999), two problems are still considerable: the treatment of chronic cases of the disease and the high level of acute cases in some Latin American countries, such as Bolivia and Mexico, where the incidence of infection in some regions reaches levels above 80% of the population (Medrano et al. 1996). In endemic areas, the transfusional transmission of Chagas disease, due mainly to urbanization and migration processes, represents a great threat (Wendel & Dias 1992). With this scenario, the major goal of research on Chagas disease is the development of specific chemotherapy that can eliminate the infection from indi-

viduals who are acutely or chronically infected, but who have not yet developed cardiac and/or digestive forms of the disease. Research and development of new diagnostic tests and drugs for the etiologic treatment of Chagas disease have been almost non-existent when compared with other non-neglected infectious diseases, such as acquired immunodeficiency syndrome (AIDS). Benznidazole and nifurtimox, the only two drugs available for treatment, have been in the market for more than 30 years and their pharmacokinetic characteristics, formulations, efficacy and secondary effects are far from ideal (Villa et al. 2007). The evaluation of cure of Chagas disease is certainly the more complex aspect of its treatment, leading several times to diverse and controversial results in relation to both parasitological and clinical cures.

Limitations for the use of specific therapy in Chagas disease

Currently, important questions remain of how best to design therapeutics or prophylactic drugs to treat or prevent Chagas disease as a parasitic infection (Tarleton 2001, Urbina 2001). The specific chemotherapy for human Chagas disease has several limitations, such as the requirement for long-term administration of highly toxic nitroimidazole derivatives that hamper treatment continuity, as well as their limited availability. In addition, natural drug resistance is quite common in the course of human infection, even among parasite populations without previous exposure to these drugs. In this sense, therapeutic results have differed between countries probably due to differences in *T. cruzi* strains from distinct geographical areas. The general assumption, however, is that the earlier the diagnosis is made and the specific treatment initiated, the greater the chance of parasitological cure (WHO 1991, Villa et al. 2007). Nevertheless, one of the greatest concerns in Chagas disease is the absence of reliable methods or a diagnostic gold standard for the evaluation of chemotherapy efficacy in treated patients. Clearance of parasitaemia and disappearance of antibodies (seronegative conversion) are taken together as cure criteria by some authors (Cançado 1963, 1997), whereas others, like Rassi and Luquetti (1992), Andrade et al. (1996) and Sosa Estani et al. (1998), admit a long period of reaction negativation and even low serological titres as criteria of cure. Effectiveness in eradicating the infection depends on the length of time between infection and treatment initiation.

According to the Brazilian Ministry of Health (MS 2005), treatment of *T. cruzi* infection is recommended during the acute phase, congenital and accidental infections, early chronic phase (children under 15 years old and elderly patients with evidence of recent chronic infection), for patients with AIDS and for those undergoing organ transplantation or receiving immunosuppressive drugs, which present a risk for reactivation of latent infection with *T. cruzi* (Fragata Filho et al. 1997).

Chemotherapy during the acute stage of infection leads to regression of clinical symptoms and parasitological cure (Cançado 1985), but its effectiveness during the indeterminate and chronic stages remains unclear (Marin-Neto et al. 2007). Studies on the clinical

evolution of Chagas disease after specific treatment are controversial and results are not convincing due to differences in casuistry, evaluation methods, time of follow-up and data interpretation (Coura & de Castro 2002). The main limitations in evaluating treatment for chronic Chagas disease arise from the need for long-term follow-up, which usually lasts several decades, and the lack of reliable tests to ensure parasite elimination (Cançado 1985, 1999). It has been proposed that latent forms of *T. cruzi* parasites present in tissues during the chronic phase of infection are not eliminated by chemotherapy (Brener et al. 2000).

Regardless of intensive research into the development of new drugs that are safer and more efficient for Chagas disease chemotherapy (Urbina 2001), the only drug now available in Latin America is benznidazole [BZ; Rochagan, Rodanil (Roche)], while nifurtimox (Lampit; Bayer) is the drug approved for chemotherapy in the United States. Their administration in the acute and recently acquired chronic phases of infection demonstrated significant trypanocidal activity (Rassi & Luquetti 1992, Andrade et al. 1996, 2004). Benznidazole (*N*-benzil-2-nitro-1-imidazole-acetamide) has direct action against both the circulating (trypomastigote) and tissular (amastigote) forms of *T. cruzi*. Its efficacy varies according to the phase of the disease, dose and duration of treatment, age, period of follow-up after therapy and the tests used to assess parasite clearance.

Nifurtimox and benznidazole have significant activities in the acute phase, with up to 80% of parasitological cure rates in treated patients (Andrade et al. 1992). However, their efficacies vary according to the geographical area, probably due to differences in drug susceptibility among different *T. cruzi* strains, a fact correlated with the biological characteristics of the parasite (Andrade et al. 1992, Cançado 1999). In the past decade, benznidazole has been reported to have significant curative activity in recent chronic disease (up to a few years post-infection), with up to 60% parasitological cure rates observed in infected children of Argentina and Brazil treated with this compound (Sosa-Estani & Segura 1999, Andrade et al. 2004). Similar results were obtained in Chile with nifurtimox (Solari et al. 2001). After the introduction of nifurtimox and benznidazole, few compounds were assayed in chagasic patients. The results obtained with allopurinol (ALLO) [4-hydroxypyrazolo (3,4-*d*) pyrimidine HPP] in experimental animals and the knowledge about its mode of action led to clinical assays for the treatment of Chagas disease (Lauria-Pires et al. 1988, Galleano et al. 1990).

The role of anti-parasitic treatment in the late chronic phase of Chagas disease remains unclear. A limited efficacy of available drugs has been observed in this phase of the disease, in which parasitological cure is achieved in 0-20% of treated patients (OPAS/OMS 1998, Lauria-Pires et al. 2000, Britto et al. 2001, Guedes et al. 2006). On the other hand, some studies performed with benznidazole have been remarkably consistent in showing moderate to significant efficacy in long-term chronic infections (de Castro et al. 2006, Sosa-Estani & Segura 2006, Viotti et al. 2006). For instance, Viotti et al. (2006)

reported the success of drug treatment in arresting disease progression in subjects who have been infected for more than 20 years.

The capacity of any given drug to eradicate the infection is dependent on intrinsic host features, such as immune status, the susceptibility of a particular *T. cruzi* genotype to the drug, as well as the drug pharmacokinetic properties and toxicity in the host and the duration of treatment. Other chemotherapeutic agents, such as ALLO and itraconazole (ITRA), are being evaluated for the treatment of chronic Chagas disease, but their advantages over benznidazole have not been determined.

Heterogeneity of *T. cruzi* populations

T. cruzi is composed of a heterogeneous population of clones with broad biological and genetic variability circulating in domestic and sylvatic cycles, which includes humans, insect vectors and animals. Moreover, this parasite undergoes long-term clonal evolution that predicts correlation among phylogenetic divergence. The parasite's biological properties play an important role in the disease pathogenesis (morbidity) and may affect drug efficacy (Andrade et al. 1975, Tibayrenc et al. 1986, Andrade & Magalhães 1996, Revollo et al. 1998, Toledo et al. 2003, 2004). The occurrence of naturally resistant *T. cruzi* strains demonstrated in experimental models (Filardi & Brener 1987, Murta et al. 2008) is supposed to be one of the most important factors explaining the low rates of cure in some treated chagasic patients from endemic zones, as seen in Central Brazil in patients infected with strains of the biodeme type III, Z1 (*T. cruzi* I) as compared with those infected with biodeme II, Z2 (*T. cruzi* II) (Andrade et al. 1992). In fact, the occurrence of predominant strains exhibiting different susceptibility levels (in specific geographic areas) has been associated with a high variability of clinical symptoms concerning treated patients. Studies have demonstrated that the so-called type III biodeme, of which the prototype is the Colombian strain, is highly resistant (Andrade et al. 1985, Andrade & Magalhães 1996, Toledo et al. 2003, 2004).

In experimental mixed infections with parasites from different *T. cruzi* genotypes, interactions between subpopulations may occur, resulting in important changes in the parasites' biological characteristics and the evolution of infection (Deane et al. 1984, da Silveira Pinto et al. 1998, 2000, Martins et al. 2006, 2007). Mixed infections show responses to benznidazole treatment distinct from the expected response based on single-infection analyses (Martins et al. 2007), suggesting that the expected correlation between susceptibility to treatment is difficult to establish for dual infections when genetic combination occurs.

The existence of a high incidence of mixed infections in humans (Solari et al. 2001) and vectors (Bosseno et al. 2000) has been verified and raises questions about its important consequences regarding morbidity, the dynamics of parasite transmission and the response to chemotherapy. The contradictory results obtained in human treatment by different authors are probably influenced by the occurrence of mixed infections, since individuals in areas where the disease is endemic may be re-infected

several times (Brenière et al. 1998). These results emphasize the importance of phylogenetic diversity of *T. cruzi* genotypes in approaches towards diagnosis and treatment of Chagas disease.

Usefulness of PCR for establishing drug efficacy

Most human infections with *T. cruzi* are only detected in the chronic phase, characterized by subpatent parasitaemia and scarce tissue parasitism. Demonstration of treatment efficacy is hampered by the lack of reliable criteria of cure, as well as the rather poor sensitivity of conventional parasitological practices. Microscopic methods lack sensitivity, and XD and haemoculture may require 30 days or more, and also lead to false-negative results. Both methods have a tendency to increase their positivity with the number of tests performed, amount of blood employed, cultivation medium, interval of time between blood collection and cultivation and other factors emphasized by Chiari et al. (1989). Moreover, these tests are time-consuming and laborious, requiring special laboratory biosecurity conditions (Brenner 1962, Gomes et al. 1999). In addition, post-therapeutic monitoring in the chronic phase of Chagas disease still constitutes a major challenge due to the long-term persistence of specific antibodies that are detected by conventional serology for several years, despite repeated negative direct parasite detection tests (Andrade et al. 1988, 1991); non-conventional serology is not frequently used (Kretzli & Brenner 1982). The presence of lytic antibodies has therefore been proposed as an indicator of active ongoing infection and treatment failure (Kretzli et al. 1982, Galvão et al. 1993). This test requires the use of live, infectious trypomastigotes and is not practical for routine use in the evaluation of chemotherapeutic agents or in clinical management of chagasic patients (Kretzli et al. 1984).

In an effort to develop more sensitive assays, PCR technology has been introduced to specifically detect *T. cruzi* DNA in blood samples from chagasic patients, opening new possibilities in the diagnosis and follow-up assessment of chemotherapy (Moser et al. 1989, Sturm et al. 1989, Avila et al. 1991, 1993, Britto et al. 1993, 1995, 2001, Wincker et al. 1994a, b, Junqueira et al. 1996, Gomes et al. 1998, 1999, Castro et al. 2002, Galvão et al. 2003). Reconstitution experiments showed that PCR procedures are able to detect the equivalent of a single parasite cell in 10-20 mL of whole blood (Moser et al. 1989, Avila et al. 1991, Britto et al. 1993). However, there are large genetic differences between distinct *T. cruzi* strains related to their biological variability, such as virulence or variable susceptibility to the immune response (control of parasitaemia) developed after the acute phase of human infection. Consequently, genetic strain differences may influence parasitaemia in man and partially explain the discrepancies of PCR sensitivity between studies carried out in different endemic areas. For instance, in Bolivia and most part of Brazil, in areas where *T. cruzi* II stocks circulate, PCR shows high sensitivity, contrasting with areas from the Brazilian Amazon and Mexico, where only *T. cruzi* I stocks occur (Avila et al. 1993, Wincker et al. 1994a, b, 1997, Zingales et al. 1998, Fernandes et al. 2001, Coura et al. 2002).

Several therapeutic studies confirm the usefulness of usual PCR strategies to evaluate treatment outcome in either acute or chronic cases of Chagas disease (Russo-mando et al. 1998, Lauria-Pires et al. 2000, Britto et al. 2001, Solari et al. 2001, Galvão et al. 2003, Schijman et al. 2003, Zulantay et al. 2004, Sánchez et al. 2005). PCR is a helpful tool for the early detection of treatment failure when comparing drug efficacy, tolerance, therapeutic schemes, periods of follow-up and cure criteria. Considering that the suppressive activity on parasitaemia is almost immediate after treatment initiation in the case of *T. cruzi* populations susceptible to the drug, the assessment of treatment outcome is mainly parasitological; anti-*T. cruzi* IgGs remain positive practically lifelong. The higher PCR sensitivity for *T. cruzi* detection confirms its potential use for evaluating chemotherapeutic efficacy compared with traditional parasitological methods, such as XD and haemoculture.

Other distinct factors may contribute to the overall performance of PCR assays: the epidemiological characteristics of the study populations, the blood volume collected, the method used to isolate DNA, parasite target-sequences [e.g., nuclear satellite DNA or kinetoplast DNA (kDNA)] and primers selected for PCR, reagents used and thermo-cycling conditions. The blood sample volume is an important factor to be considered in chronic patients with low levels of parasitaemia following treatment, and the differences in PCR sensitivity can be explained by the intermittent presence and quantity of circulating parasites at the time of blood collection (Castro et al. 2002).

Whether a positive PCR reflects detection of intact parasites or circulating DNA derived from lysed organisms is not clear. Tarleton and Zhang (1999) reported that after intramuscular injection of a large quantity of *T. cruzi* kDNA in mice, PCR performed on blood samples was positive up to 48 h after inoculation, suggesting that parasite DNA detected by PCR derives from intact, extracellular or recently lysed parasites.

If so, PCR is a rapid and safe indicator of the parasite's susceptibility to drugs, allowing early therapy modification in cases of resistance or reactivation of chagasic infection (Schijman et al. 2000, Lages-Silva et al. 2002). According to Galvão et al. (2003), there is no guarantee that a single "flash" of a negative PCR means parasitological cure, especially when the well-known waves of parasitaemia during the long course of Chagas disease are taken into account. A negative post-treatment PCR result may be indicative of the absence of parasite DNA at that moment. Negative results for both, serology and PCR are probably indicative of cure. The value of parasitological tests lies mainly in the positive results they yield; thus, a positive PCR in blood may reflect treatment failure. Hence, PCR can be used as an early marker of specific chemotherapy resistance years before serological reversion [long-lasting sustainability of negative seroconversion, as pointed out by Rassi and Luquetti (2003)]. If the persistence of positive PCR results is considered a therapeutic failure, assessment of parasite load by quantitative real-time PCR could still be correlated with the impact of trypanocidal treatment on the disease evolution.

Treatment of congenital infections - T. cruzi congenital transmission may occur in some or all pregnancies in the infected mother, whether in the acute or chronic stages, normally with oligoparasitaemia and usually as an asymptomatic disease (Bittencourt 1992). Variable rates of congenital transmission have been reported in different geographical areas where distinct parasite strains predominate, suggesting that parasite genotypes might play a role in the risk of congenital transmission. Moreover, in cases of transmission, it is unknown if the whole maternal *T. cruzi* population or certain clones are preferentially transmitted by the transplacental route (Burgos et al. 2007). The prevalence of *T. cruzi* infection among pregnant women ranges from 2-51% in urban areas and from 23-81% in rural regions of Latin America (Freilij & Altcheh 1995).

Standard serodiagnosis of *T. cruzi* infection in infants born to seroreactive women has low positive predictive value, since the presence of anti-*T. cruzi* IgG antibodies in the newborn may be due to passive transfer of maternal IgG antibodies, which in the non-infected infant would normally disappear around the sixth month of age (Moya et al. 1989, Freilij & Altcheh 1995, Luquetti 1997). Moreover, a small proportion of infected newborns is seronegative. The detection of IgM antibodies against *T. cruzi* is not satisfactory (Freilij & Altcheh 1995, Blanco et al. 2000). Therefore, the diagnosis of congenital infections usually relies on microscopic observation of bloodstream trypomastigotes, which is more effective by microhaematocrit concentration technique in infants less than six months of age (Freilij et al. 1983, Freilij & Altcheh 1995). Anti-parasitic treatment has greater success in newborns close to birth and is strongly indicated (Freilij & Altcheh 1995, Luquetti 1997). In search of more sensitive laboratory tests to detect infection and evaluate treatment outcome in congenital infections, PCR appears to be a promising laboratory tool (Russumando et al. 1998, Schijman et al. 2003, Virreira et al. 2005, Duffy et al. 2009).

Russumando et al. (1998) examined 58 infants born to seropositive mothers for congenital transmission of Chagas disease in Paraguay using direct microscopy, immunofluorescence for anti-*T. cruzi* IgM antibodies and PCR targeted to nuclear genomic markers (Moser et al. 1989). It was possible to detect *T. cruzi* DNA from 100 uL of plasma in six newborns (10%) on the day of birth who were IgM negative. Treatment with benznidazole (7 mg/kg of body weight) was given in two daily doses for 60 days and was initiated on the day *T. cruzi* infection was confirmed by direct microscopic observation or haemoculture and positive PCR. The infected infants were followed-up for four years by serology and PCR. Serology remained positive 3-8 months post-treatment and the evaluation of treatment effectiveness by PCR presented higher sensitivity, as soon as 15 days after treatment. These data suggested that the later the treatment is initiated, the later the negative seroconversion takes place. The authors concluded that conventional parasitological and serological techniques were not efficient enough to provide early diagnosis of con-

genital Chagas disease, especially in rural areas where the disease is endemic. PCR sensitivity was usually higher than haemoculture and practically confirmed 100% treatment efficacy in association with conventional serology negativation.

Schijman et al. (2003) performed the first prospective PCR study in a cohort of paediatric patients with congenital Chagas disease from a non-endemic area in Argentina, without risk of vectorial re-infections. The study focused on the evaluation of anti-parasitic therapy in patients with congenital infection, diagnosed and monitored by conventional and PCR-based assays targeting the parasite mitochondrial genome or kDNA. A competitive PCR was carried out in order to characterize the bloodstream parasite load in patients at the time of diagnosis, as well as during post-treatment follow-up. To evaluate the outcome of 40 patients undergoing anti-parasitic treatment after 2-3 years post-chemotherapy, the children were classified into three age groups: GI (< 3 months of age), GII (7 months-2 years old) and GIII (> 3 years old). Patients from both GII and GIII represented the undetermined phase of Chagas disease. Cure was demonstrated by both negativation of PCR and serology in 100% of infants from GI, in 66.7% of children from GII and in only 12.5% of GIII cases. In those infants who started therapy in their first months of life (< 3 months), cure was achieved early after treatment, in accordance with previous studies (Blanco et al. 1999, 2000). In these cases, PCR became negative earlier or at the same stage as serology did. The authors did not observe differences between nifurtimox or benznidazole for toxicity, clinical manifestations, serological and PCR outcome. They concluded that PCR is a helpful tool for early detection of treatment failure in patients who initiate therapy at the undetermined phase of the disease. The dissociation between post-treatment persistent negative PCR results with positive conventional serology after three years of follow-up observed in 20 out of 30 undetermined Chagas disease patients (GII + GIII) was in agreement with previous findings in treated chronic patients presenting reactivity by conventional serology with clearance of lytic antibodies (Galvão et al. 1993). The investigation by Schijman et al. (2003) reinforces the need to screen all pregnant women living in or emigrating from endemic areas in order to provide their newborns with an early accurate diagnosis for a more successful treatment outcome.

Treatment in early life - It is well documented that infants infected during the early years of life represent those patients amenable to specific treatment with the greatest possibility of cure (Rassi & Luquetti 1992, Villa et al. 2007). Children better tolerate a long-term treatment and present high indices of cure, as demonstrated by two randomized placebo-controlled trials with benznidazole in Brazil (Andrade et al. 1996) and in Argentina (Sosa Estani et al. 1998), where negative seroconversion was estimated in around 60% of the benznidazole-treated children. Thus, in acute and recent infection cases, an effective treatment might prevent the progression of infection to disease and its complications.

Solari et al. (2001) evaluated 66 Chilean chagasic children from 0-10 years old treated with nifurtimox and further monitored every three months in the first year and every six months during the second and third year post-therapy. Chemotherapy efficacy was assessed by using PCR for *T. cruzi* kDNA, XD and serology (ELISA-IgG and indirect haemagglutination). The results showed that serology still remained positive and serum titres unchanged at 36 months following treatment, except in the two youngest cases (1 and 14 months old) out of seven cases ranging from 1-20 months old. Therapeutic effectiveness, as determined by serological cure, seemed to depend on the delay between infection and the start of treatment. Before therapy, all children were positive by PCR. The rate of parasitological cure in 0-10 year-old children was variable and dependent on the region in which those children lived. Several months were required to obtain a prolonged negative result by PCR, even though positive XD rapidly dropped 3-6 months after treatment from 79.1-1.6%, probably due to the lower sensitivity of the method. In this study, the slow rate of PCR conversion could be explained by the late administration of the drug after infection as compared to the early efficacy of treatment in congenital cases, from a few days to three months (Russomando et al. 1998). According to Solari et al. (2001), PCR was the most effective test to monitor children three years post-chemotherapy, when all treated children converted from positive to negative, and no significance was found between patient age and PCR conversion. Conventional serology, however, remained positive three years post-chemotherapy in most cases.

The investigation conducted by Galvão et al. (2003) used a PCR based on kDNA to assess the rate of chemotherapy failure. They studied a well-characterized Brazilian cohort of *T. cruzi* seropositive schoolchildren previously enrolled in a field trial of benznidazole efficacy who lived in a highly endemic rural area in Central Brazil. Paired blood samples (5 mL) from 111 children were taken at baseline and 36 months after treatment with either benznidazole (n = 58) or placebo (n = 53). At the end of follow-up, PCR was positive in 39.6% of the treated group versus 64.2% of the placebo group, indicating that untreated patients had a 1.6-fold higher chance of remaining PCR-positive than those who received specific chemotherapy (p = 0.01). In the study, the possibility of re-infection was discarded. Among those who received benznidazole, 22.6% did not present a decrease in the indirect immunofluorescence (IIF) titres and PCR continued to be positive for 39.6% (p > 0.05), pointing out the proportion of benznidazole failure after a 3-year follow-up. An adequate correlation could be found between a high proportion of negative PCR results and a decrease in antibody titres in the treated group. PCR positivity occurred in patients without reductions in antibody titres. In the placebo group, a higher proportion of PCR positivity (64.2%) was detected in agreement with the absence of a decrease in IIF titres (77.4%). In this work, PCR was confirmed as a useful tool for revealing therapeutic failure of *T. cruzi* infection on a short-term basis, although the authors highlight the possibility that among treated children with negative PCR,

some of them were expected to shift to positive PCR during long-term follow-up, especially when the well-known waves of parasitaemia during the long course of Chagas disease are taken into account.

Treatment of chronic infections - Effective chemotherapy of *T. cruzi* infection is needed due to the enormous burden of chronic infections imposed on human populations living in endemic areas in which active insect-vector transmission occurs. While the elimination of Chagas disease has been considered a reasonable public health goal, controversies still remain about the efficiency of trypanocidal chemotherapy, especially in chronic asymptomatic individuals. It is recommended that late chronic infections without clinical manifestation or with mild cardiac or digestive symptoms should be treated during 60-90 days, in accordance with the tolerance to the drugs, aiming to prevent or reduce the evolution of Chagas disease to more severe forms (Luquetti 1997, Coura & de Castro 2002). In spite of the evidence that nitroimidazole derivatives improve parasite-related outcomes in both children and adults with chronic asymptomatic *T. cruzi* infections, it has not yet been definitely proven whether the reduction in parasite load translates into improved clinical outcomes (Marin-Neto et al. 2008). Up to 80% of patients treated during the late chronic infection do not achieve parasitological cure as assessed by the persistence of positive serology, now confirmed using PCR-based methods.

The great advances in vector control in the South Cone countries and demonstration of the parasite in chronic patients by more sensitive diagnostic tests indicated the urgency to discuss the etiologic treatment during this phase, and thus reinforcing the need to find drugs with more efficacy and less toxicity. Agents such as ALLO and ITRA have been evaluated for treatment of chronic infections, but their advantages over benznidazole have not yet been determined. One of these studies showed that ITRA, used as an anti-fungal drug for 20 years, produced parasitological cure in 53% of chronic chagasic patients (6 mg/kg/day for 120 days), whereas ALLO (8.5 mg/kg/day for 60 days) was able to eliminate *T. cruzi* infection in 44% of the treated patients (Apt et al. 1998). The criteria for parasitological cure were the maintenance of negative XD and/or complement-mediated lysis for at least four years of follow-up. Double-blind randomized longitudinal studies must still be performed to re-evaluate the efficacy of these drugs for treatment of Chagas disease.

Cure assessment in chronic infections is controversial and difficult to demonstrate, mainly due to the lack of sensitive or specific tests to document parasitological cure (Kretzli et al. 1984, Galvão et al. 1993). However, in the 1990s, assays based on PCR were used to detect *T. cruzi* DNA in the blood of chronic chagasic patients with an adequate sensitivity (a single parasite cell in 10-20 mL of whole blood) and proven to be a promising tool for evaluating parasitological failure after specific etiologic treatment in chronic infections (Avila et al. 1991, Britto et al. 1993, 1995, 2001, Wincker et al. 1994a, b, Junqueira et al. 1996, Gomes et al. 1999).

Britto et al. (1995) proposed for the first time the use of PCR tests to monitor chagasic patients submitted to specific treatment as a complement to serological investigation of cure. They assessed the performance of a PCR assay directed to *T. cruzi* kDNA for the evaluation of chemotherapy outcome in 32 benzimidazole-treated seropositive chronic patients attending the Hospital of Infectious Diseases from Fiocruz, in Rio de Janeiro. Positive amplification results were observed in only nine out of 32 treated patients who remained reactive with conventional serology, even after a mean period of five years post-treatment. XD was not able to detect parasite persistence in five out of the nine PCR-positive treated patients. Consequently, PCR appears remarkably useful for early detection of cases refractory to therapy and should be employed in association with serology for the assessment of cure in chronic Chagas disease infection.

Lauria-Pires et al. (2000) performed a 10-year follow-up study to determine the effectiveness of nitro-derivative therapy in a cohort consisting of 45 treated chronic patients with severe electrocardiographic alterations (62% treated with nifurtimox and 38% with benzimidazole), 46 chronic chagasic individuals who did not receive therapy and 41 uninfected controls. Anti-*T. cruzi* antibodies were consistently lower one year after treatment than 10 years thereafter ($p < 0.001$). Blood from all treated and 93.7% of untreated chagasic patients yielded PCR-amplified product from the parasite nuclear DNA, indicating active infection. A competitive PCR demonstrated no significant difference in parasite load between treated (13.8 ± 14.9 *T. cruzi*/mL of blood) and untreated (20.1 ± 22.6) patients. The evidence of ongoing infections was in accordance with the immunological results, which were indistinguishable in the treated and untreated groups of patients. In this study, treatment with nitrofurans and nitroimidazole compounds did not lead to parasitological cure and did not alter the progression of heart disease in chronic chagasic patients. Progressive electrocardiographic alterations were recorded in treated as well as untreated patients, the proportions not being statistically different. The authors suggest that nitro-derivative therapy may discharge parasitaemia as a consequence of drug-induced immunosuppression and parasite resistance, probably through the selection of highly virulent parasite clones.

In the study by Britto et al. (2001), PCR and XD performed 20 years after trypanocidal chemotherapy were compared aiming to investigate parasite clearance in 85 chronic asymptomatic individuals who were previously diagnosed as unquestionably chagasic by XD. These individuals inhabited two distinct geographic areas from Brazil in which vector transmission was interrupted for more than 20 years, and were submitted to specific treatment with either benzimidazole or nifurtimox (37 in the acute phase and 48 in the chronic phase). The untreated group consisted of 15 xenopositive chronic asymptomatic individuals that received placebo. Treatment in the acute phase yielded PCR-negative results in 73% of the cases (endpoint analysis), while XD negativation was observed in 86%. Regarding the individuals treated during

the chronic phase, PCR was negative in 65% and 83% had XD-negative results 20 years after treatment. For the untreated group, XD was not able to find the parasite in 73%, in which 36% were positive by the molecular approach. It is important to highlight that a negative PCR result might be indicative of the absence of parasite DNA at that moment. Herein, as only one point in time was analyzed, a negative result could not be assumed as definitive cure. The agreement of negative results between serology and PCR is probably indicative of cure. In this investigation, only 10 individuals that were considered seronegative several years after the start of chemotherapy unequivocally presented negative results by PCR, demonstrating the elimination of circulating parasites in these cases. Seventeen individuals had their antibody titres decreased to such a level, and the final results were considered doubtful; 16 of them presented negative PCR. The results supported the advantage of PCR over conventional parasitological techniques to demonstrate persistent infections in chronic chagasic patients undergoing chemotherapy.

Zulantay et al. (2004) investigated the use of PCR and hybridization assays to detect *T. cruzi* in 52 chronic chagasic patients from a highly endemic region of Chile six years following therapy with either ITRA or ALLO. All patients yielded negative XD six years post-treatment with either drug; XD was performed at the same time as blood sample collection for PCR. The PCR alone enabled the identification of *T. cruzi* kDNA in 40% of patients treated with ITRA and in 60% of those ALLO-treated. When associated with an isotopic hybridization step (kDNA as probe), PCR positivity was higher in the group treated with ITRA (60%) and a slight reduction of PCR sensitivity was observed in patients that received ALLO, due to only one case where the hybridization result was not concordant and, consequently, the fraction of PCR positivity dropped to 53%. The authors reinforced the use of PCR coupled to hybridization to demonstrate parasite persistence in chemotherapeutic evaluation.

In the work by Sánchez et al. (2005), a combination of PCR and flow cytometry analysis of anti-live trypanomastigote antibodies (FC-ALTA) was performed to monitor parasite clearance in 54 chronic patients treated with ALLO ($n = 31$) or ITRA ($n = 23$) 10 years earlier. All patients maintained positive conventional serology up to 10 years follow-up. XD gave positive results in 35.5% and 60.9% of the ALLO and ITRA-treated groups, respectively, while both PCR and FC-ALTA were positive in 74.2% ALLO and 87.0% ITRA. Discordant results between PCR and FC-ALTA were as follows: (i) 13% of total patients were negative by PCR and positive by FC-ALTA and (ii) 5% gave positive PCR results with negative FC-ALTA. Only one case where both tests were negative was considered cured. The results did not show differences in efficacy between the drugs and reinforced the relevance of using sensitive tools such as PCR and FC-ALTA to follow-up patients with chronic Chagas disease.

Coronado et al. (2006) studied the distribution of *T. cruzi* clones after treatment failure with ITRA or ALLO

12 years after therapy completion in chronically infected individuals in an endemic area free of vectorial transmission in Chile. Blood samples from treated and untreated individuals were used to detect the parasite by PCR following confirmation with hybridization using total kDNA as the universal probe; FC-ALTA was also employed to detect active infection. In parallel, XD was performed with *Triatoma infestans* fed from the same group of patients. Southern-blot analysis of PCR products using two samples from each patient (blood and XD) was performed with a panel of four genotype-specific probes corresponding to *T. cruzi* clones 19 (TcI), 32 (TcIIb), 39 (TcIIId) and 43 (TcIIe). The results revealed complex hybridization patterns for both blood and insect samples, indicative of infection with more than one clone in some patients. No evidence of cure was assessed by FC-ALTA or PCR with either blood or XD samples. The percentages of PCR-positive samples in blood (89.5%) and XD samples (91.2%) increased to 100% and 96.5%, respectively, after hybridization. Although the quantification of parasitaemia levels was not performed, the authors predicted that the lower or higher percentage of a particular *T. cruzi* clone may be indicative of parasite susceptibility or resistance to chemotherapy. Clone TcI was present in 69.2% of the XD samples from the ALLO-treated group compared to the control group (non-treated, 29.6%; $p = 0.0178$), suggesting resistance of this clone to ALLO. When the ITRA-treated group was compared with the control group, significant differences were found in both blood and XD samples. In blood, the clone TcIIb was detected in 35.5% of treated individuals and in 66.7% of the control group ($p = 0.0207$), which suggested susceptibility of this clone to ITRA. Regarding the XD samples, clone TcI was observed in 82.3% of the ITRA-treated individuals but only in 29.6% of the control group ($p = 0.0006$), thus also indicating resistance of this clone to ITRA. No significant differences were found for the *T. cruzi* clones TcIIId and TcIIe in the treated (ALLO and ITRA) and non-treated groups.

The results from Coronado et al. (2006) obtained for the first time in humans were able to confirm previous data using murine models concerning the association between *T. cruzi* phylogenetic diversity and chemotherapeutic response (Andrade et al. 1985, Toledo et al. 2003). In general, these studies suggest that susceptibility or resistance to a drug depends on the *T. cruzi* genotype, thus indicating that the appropriate drug or combination of drugs should depend upon the infective parasite clone or mixture of clones present in a particular host. In situations relating to parasitological cure failure, the best adapted *T. cruzi* clone to a host would be associated with the resistance to chemotherapy (TcI increased in patients treated with ALLO and ITRA). In contrast, the lower adaptation of *T. cruzi* clones to a host would be correlated with drug susceptibility (TcIIb decreased only in the ITRA-treated patients).

The efficacy of benznidazole therapy in preventing clinical complications in patients with pre-existing cardiac disease is now being evaluated in the scope of the BENznidazole Evaluation for Interrupting Trypano-

somiasis (BENEFIT) study, conducted by the Canadian Institutes of Health Research and WHO/TDR in collaboration with investigators in Latin America and expected to be completed in 2011 (5 years follow-up). BENEFIT is a multicentre, randomized, double-blind, placebo-controlled trial of 3,000 patients with Chagas cardiomyopathy in Latin America, thus representing the largest trial yet conducted on Chagas disease (Marin-Neto et al. 2008). The study was designed to clarify the role of trypanocidal therapy for prevention of cardiac disease progression and death and also comprises a sub-study to evaluate the effects of benznidazole on parasite clearance using PCR-based methods (qualitative and quantitative assays). The BENEFIT program will provide a unique opportunity for better understanding of the clinical progression of chronic Chagas cardiomyopathy and it is expected to provide conclusive information on the role of etiologic treatment in this phase of the disease (Marin-Neto et al. 2008).

Final comments

Great efforts are being made in countries of South America to control Chagas disease transmission by either insect vectors or blood transfusion, although more effective chemotherapy is needed for the million people who are already infected. There is a need for new drugs with fewer secondary effects and higher trypanocidal activity. The advancement in studies with posaconazole, already commercialized for systemic mycotic infections, and ravuconazole, in the animal experimental phase, is urgent (Villa et al. 2007). However, one of the key issues concerning Chagas disease is still that of diagnosis. Without effective diagnostics, infected individuals cannot be identified and hence treated, and the success of treatment cannot be efficiently assessed. The absence of a true gold standard (i.e., a method to consistently detect the presence of parasites in those *T. cruzi*-infected individuals) makes it difficult to evaluate the sensitivity of serological tests (Tarleton et al. 2007). The development of highly sensitive and specific diagnostic field and laboratory tools to determine active *T. cruzi* infection is a crucial requirement.

In the last two years, investigators from Latin America recognized the need to standardize and validate PCR-based assays for Chagas disease diagnosis and disease management across laboratories and countries. This necessity was due to the fact that each laboratory applies their own protocols and technologies, making comparison of PCR-based findings among groups unreliable. As an integrated effort of the WHO/TDR and PAHO, a workshop and symposium were held in Buenos Aires (November 2008) supported by TDR, INGEBI-Conicet UBA and the United Nations University UNU-BIOLAC, in which investigators representing different countries established a "Consortium for standardization and validation the clinical use of PCR for *T. cruzi* DNA detection in Chagas disease". The meeting allowed the definition of the best PCR practices, as well as the applicability of these in a clinical setting and their interpretation (special attention was given to diverse parasite

lineages); it also allowed establishing the limitations of this technology. All of the obtained information led to the definition of a Standard Operating Procedure (SOP) that could be used by those interested in applying PCR as a diagnostic tool, particularly in some special circumstances (e.g., post transplantation, HIV co-infection, congenital infection) and also for follow-up patient treatment outcome. The special value of this SOP is foreseen in the context of evaluating new drug candidates in clinical studies. This SOP, if validated through use, could also become the standard reference method for evaluating new PCR methodologies.

The implementation of quantitative PCR assays to determine bloodstream parasite load and follow its evolution during treatment could be particularly useful as an indicator of response in prolonged therapeutic regimens (Urbina 2001), as it is mandatory in patient management of certain viral infections (Piatak et al. 1993, Berger & Preiser 2000). Until now, it has not been well defined if chronic or asymptomatic patients presenting extremely low or even undetectable circulating parasite levels should present a better response to treatment than those with patent parasitaemia. This hypothesis underlines the urgent need to join efforts for standardization of both conventional and real-time PCR protocols, which will probably be the basis for the future establishment of more reliable healing cure criteria for patients submitted to Chagas disease chemotherapy. Moreover, the parasitic load might be a useful epidemiological tool to estimate patients' infectivity concerning the risk of transmission (Schijman et al. 2003).

Studies have been conducted aimed at the development of real-time PCR strategies using both SybrGreen and TaqMan technologies for the diagnosis of *T. cruzi* infection (Cummings & Tarleton 2003, Virreira et al. 2006, Piron et al. 2007, Duffy et al. 2009), and for molecular typing of the parasite genotypes (Freitas et al. 2005, Burgos et al. 2007, Duffy et al. 2009). Cummings and Tarleton (2003) were pioneers in developing a real-time PCR for accurate and sensitive quantification of tissue parasite burden in mice infected with *T. cruzi*. The method was able to confirm the higher parasite load in mice with acute infections in comparison with chronically infected mice, the detection of tissue-restricted parasite persistence in different parasite: host strain combinations and the observation of increased tissue parasite burden with higher infective inoculation doses. Virreira et al. (2006) compared two PCR methods with primers for both nuclear and kDNA sequences for the diagnosis of congenital infections in amniotic fluids of *T. cruzi*-infected mothers, and also determined the number of parasites in positive samples by real-time PCR (SybrGreen system) targeting the kDNA, following the protocol first described by Cummings and Tarleton (2003), with slight modifications. Piron et al. (2007) developed a real-time PCR based on the use of a TaqMan fluorogenic probe directed to *T. cruzi* satellite DNA sequences for the detection of parasite DNA in the blood of chagasic patients (adult chronic individuals and one child with an acute congenital infection). A 90% concordance was

achieved using a conventional nested PCR designed for the same parasite target, suggesting that real-time PCR provides an optimal alternative to nested PCR with similar sensitivity and higher throughput and can facilitate in determining ongoing parasitaemia in chagasic patients.

An additional application of real-time PCR was first described by Freitas et al. (2005) and concerns its use for genetically typing major *T. cruzi* lineages directly in chronically infected human tissues (oesophagus, heart and colon) from patients presenting gastrointestinal or cardiac forms of Chagas disease. The high sensitivity of PCR-based typing provides direct assessment of parasite genetic diversity in clinical specimens without the need of culture isolation. Following PCR amplification with the SybrGreen system targeting the parasite ribosomal DNA, the products were precisely identified using thermal denaturation curves in real-time. Burgos et al. (2007) used a similar approach with primers directed to nuclear genomic markers to identify bloodstream *T. cruzi* lineages in the blood of congenitally infected children and their mothers from endemic localities of Argentina and Bolivia. The results indicated no association between a particular *T. cruzi* genotype and vertical transmission of Chagas disease in these countries.

More recently, Duffy et al. (2009) developed an accurate real-time PCR strategy using SybrGreen targeted to conserved motifs within the *T. cruzi* repetitive satellite sequence for monitoring Chagas disease reactivation in heart-transplanted patients who received immunosuppressive therapy, and also for etiological treatment outcome in paediatric patients. The approach was able to identify parasites according to the number of satellite repeats detected per genome for different *T. cruzi* lineages. In this work, the addition of a standardized amount of plasmid containing a heterologous sequence allows normalization of the DNA extraction yields and detection of false-negatives due to inhibition under any clinical situation.

In general, the true potential of real-time PCR will be recognized in situations for which PCR-based techniques have been promoted, such as congenital infections (Schijman et al. 2003, Virreira et al. 2003, Mora et al. 2005), monitoring parasitaemia during and after treatment (Russomando et al. 1998, Britto et al. 2001, Schijman et al. 2003, Apt et al. 2005, Sánchez et al. 2005), early detection of relapses after heart transplantation (Maldonado et al. 2004) and other immunosuppressive circumstances. To deal with more precise results and make them comparable between laboratories, the main constraint until now has been the lack of a universal reagent presenting accurately quantified *T. cruzi* DNA samples to be used as standards in all quantification assays (Piron et al. 2007). International standards for various infectious agents (human immunodeficiency virus, hepatitis C virus, etc.) are commercially available for molecular biology assays in order to standardize and compare methods and laboratory performances. In this sense, agencies providing such materials should be strongly encouraged to make properly characterized *T. cruzi* reagents available for this purpose.

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