Review

High Throughput Screening for Anti–Trypanosoma cruzi Drug Discovery

Julio Alonso-Padilla, Ana Rodríguez

New York University School of Medicine, Department of Microbiology, Division of Parasitology, New York, New York, United States of America

Abstract: The discovery of new therapeutic options against Trypanosoma cruzi, the causative agent of Chagas disease, stands as a fundamental need. Currently, there are only two drugs available to treat this neglected disease, which represents a major public health problem in Latin America. Both available therapies, benznidazole and nifurtimox, have significant toxic side effects and their efficacy against the life-threatening symptomatic chronic stage of the disease is variable. Thus, there is an urgent need for new, improved anti–T. cruzi drugs. With the objective to reliably accelerate the drug discovery process against Chagas disease, several advances have been made in the last few years. Availability of engineered reporter gene expressing parasites triggered the development of phenotypic in vitro assays suitable for high throughput screening (HTS) as well as the establishment of new in vivo protocols that allow faster experimental outcomes. Recently, automated high content microscopy approaches have also been used to identify new parasitic inhibitors. These in vitro and in vivo drug discovery approaches, which hopefully will contribute to bring better anti–T. cruzi drug entities in the near future, are reviewed here.

Introduction

Chagas disease, caused by the protozoan parasite Trypanosoma cruzi, is endemic to Central and South America [1,2]. Its impact represents the largest burden due to a parasitic disease in the Western hemisphere [2]. Estimates point at 20,000 deaths per year; over 10 million people currently infected, many of them unknowingly; and another 100 million living in risk-of-transmission areas [3]. The infection is transmitted by hematophagous triatomine insect vectors (Reduviid order) that easily proliferate in settings with poor housing conditions, rendering those with lesser economic resources most vulnerable to the infection [2,4]. The parasitic cycles between insect vectors and mammalian hosts in four distinguished parasitic life-stages: replicative epimastigotes and host infective metacyclic trypomastigotes in the insect, whereas free-swimming trypomastigotes and intracellular replicative amastigotes are found in the host [4].

The disease starts with a short acute phase, during which blood parasite levels are relatively high and sufficient for successful conventional diagnosis. Although mostly asymptomatic, this first stage can cause death in approximately 5% of acutely infected patients [4]. An indeterminate “silent” stage that may span more than 20 years follows, during which, despite pathogen persistence, parasitemia is hardly detectable and no major clinical symptoms are encountered [4]. Many of the indeterminately infected people clear the infection or just never present clinical symptoms; however, about 30% of the chronically infected patients suffer, at some point, severe heart and gut tissue damage that, if untreated, ultimately leads to total disability and death [2,4]. With no vaccine in view, the treatment of Chagas disease relies on chemotherapy with benznidazole (BNZ; LAFEPE, Pernambuco) and nifurtimox (NFX; Lampit, Bayer Healthcare, Leverkusen), which have been in use for the last 40 years and have failed to control the disease [5]. Long-term dosage regimes are required for effective cure; however, severe side effects often prompt the discontinuation of the treatment [5]. Both drugs show good efficacy against the short and mostly asymptomatic acute stage and are also indicated for treatment during the early chronic phase and to prevent congenital transmission [6]. In the chronic phase, the efficacy of these drugs can vary because of reasons like patient age, the time the patient has had the disease, and the criteria used to define parasite clearance. Although there is not yet agreement about the efficacy of these drugs in the chronic phase [2], recent reports demonstrate significant efficacy and propose that treatment should be administered at all disease stages [7–11]. Changes in the long dosages regimes and/or combinatorial therapies have also been suggested to avoid the severe side effects of the current available drugs [12]. In any case, new therapeutic options are much awaited, not only due to toxicity but also because susceptibility to BNZ and NFX among different T. cruzi strains/isolates is variable [5]. However, the current clinical trials portfolio against Chagas disease is populated with modifications on the use of the already available BNZ and NFX (Table 1) and two re-purposed antifungal azoles. The latter compounds held great promise as potent inhibitors of the parasite fundamental ergosterol biosynthetic pathway [21]. Unfortunately, despite sporadic successful outcomes [22], the Phase II clinical trials performance of posaconazole and the ravuconazole derivative E1224 have been disappointing [23–25].

Search efforts for new anti–T. cruzi drugs stand upon the advancements achieved in the past years with the use of engineered reporter parasites for in vitro phenotypic assays [26–28]; reliable, quick in vivo protocols [28,29]; as well as the application of cutting-edge digital imaging technology like high-content microscopy [30–32]. In the search of new chemotherapeutics against neglected diseases, in which very few validated targets exist, non-reductionist, whole cell phenotypic approaches hold significant advantages [33]. Phenotypic screening represents

Citation: Alonso-Padilla J, Rodríguez A (2014) High Throughput Screening for Anti–Trypanosoma cruzi Drug Discovery. PLoS Negl Trop Dis 8(12): e3259. doi:10.1371/journal.pntd.0003259

Editor: Eric Dumonteil, Universidad Autónoma de Yucatán, Mexico

Published December 4, 2014

Copyright: © 2014 Alonso-Padilla, Rodríguez. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Funding: The authors received no specific funding for this study.

Competing Interests: The authors have declared that no competing interests exist.

* Email: ana.rodriguez@nyumc.org
Methods

Literature searches for this work were made in PubMed under the terms “anti-Trypanosoma cruzi assays,” “image-based parasites,” and further following the citations in related references. Information on clinical trials was obtained from the website http://clinicaltrials.gov and updated press releases were queried on general web browsers. Performance and outcome of the retrieved hits is of great importance [29,35,36]. The availability of these microorganisms transformed the previously time-consuming, labor-intensive, observer-biased visual microscopic counting, into faster, unbiased colorimetric, fluorometric, or luminescence-based measurements, which have made feasible the screening of large compound libraries. Transgenic T. cruzi parasites that express bacterial β-galactosidase reporter enzyme [26], firefly luciferase [28,38], or the tandem tomato fluorescent protein [28] have been made available to the scientific community. As part of their development as tools for anti-T. cruzi inhibitory assays, these genetically engineered microorganisms were shown to biologically perform like their wild-type counterparts, in terms of growth, cycle stage transitions, mice infection, and in vitro sensitivity to known drugs [26,28]. For the recombinant parasites carrying either the β-galactosidase reporter gene or the firefly luciferase gene, a substrate addition step is required and a single end-point measurement of the reporter enzyme activity can be obtained. To detect β-galactosidase activity, colorimetric and luminescent substrates have been used (respectively, chlorophenol-red-β-D-galactopyranoside, Sigma-Aldrich [26], and GalScreen, Life Technologies [39]). It must be noted that colored compounds, which are frequently found in chemical libraries, might interfere with the readout when using colorimetric reporters [26]. This issue can be addressed by using luminescent or fluorescent-based readouts as in the first HTS campaign performed against T. cruzi at the Broad Institute [39] and in the more recent HTS made at GSK [40].

In Vitro Phenotypic Assays to Identify New Anti-T. cruzi Drugs

HTS assays based on recombinant reporter parasites

An increasing knowledge of T. cruzi molecular biology and the development of plasmids and efficient parasite transfection protocols [37] permitted the construction of reliable and robust transgenic T. cruzi parasites of multiple genetic backgrounds [26,28,37,38]. The availability of these microorganisms transformed the previously time-consuming, labor-intensive, observer-biased visual microscopic counting, into faster, unbiased colorimetric, fluorometric, or luminescence-based measurements, which have made feasible the screening of large compound libraries. Transgenic T. cruzi parasites that express bacterial β-galactosidase reporter enzyme [26], firefly luciferase [28,38], or the tandem tomato fluorescent protein [28] have been made available to the scientific community. As part of their development as tools for anti-T. cruzi inhibitory assays, these genetically engineered microorganisms were shown to biologically perform like their wild-type counterparts, in terms of growth, cycle stage transitions, mice infection, and in vitro sensitivity to known drugs [26,28]. For the recombinant parasites carrying either the β-galactosidase reporter gene or the firefly luciferase gene, a substrate addition step is required and a single end-point measurement of the reporter enzyme activity can be obtained. To detect β-galactosidase activity, colorimetric and luminescent substrates have been used (respectively, chlorophenol-red-β-D-galactopyranoside, Sigma-Aldrich [26], and GalScreen, Life Technologies [39]). It must be noted that colored compounds, which are frequently found in chemical libraries, might interfere with the readout when using colorimetric reporters [26]. This issue can be addressed by using luminescent or fluorescent-based readouts as in the first HTS campaign performed against T. cruzi at the Broad Institute [39] and in the more recent HTS made at GSK [40].

Another reason
for preferring luminescence and/or fluorescence detection methodologies is their higher sensitivity, which allows for greater miniaturization of assays. An important issue to be considered in the use of transgenic parasites in the screening of chemical libraries is that reporter activity inhibitors (i.e., chemical compounds that inhibit luciferase or β-galactosidase) are selected as hits. Thus, secondary assays, such as image-based assays that rely on non-recombinant strains, are needed to identify these false hits.

Firefly luciferase-expressing parasites require the addition of luciferin before performing the readout [28,30]. In contrast, the important advantage offered by the self-fluorescent recombinant parasite strains developed in Prof. Tarleton’s lab resides on their substrate-independent detection, which allows a continuous activity measurement [28]. This property appears particularly important to investigate parasite growth kinetics, which could provide valuable insights of compounds’ mechanisms of action and, for example, aid to prioritize those with high parasite lytic activity. However, the higher sensitivity of detection in vivo of methods such as PCR of specific tissues compared to luminescent or fluorescent parasites would make these methods more useful in the detection of small populations of parasites, for example, in chronically infected animals.

Recombinant T. cruzi parasites have already been used in a series of in vitro assays designed to detect drug susceptibility mainly of the intracellular amastigote life-stage [26–28], the parasite replicative form in the mammalian host and the preferred parasite target stage. When screening a molecular library against intracellular amastigotes, parallel host cell toxicity assays on the retrieved hits or “active” compounds are necessary to determine whether the activity is specifically antiparasitic or either totally or partially due to a disruption of host-cell biology [27,41,42]. The selectivity index is then calculated using the relative IC₅₀ values of the host cell and the parasite.

The most important impact of the developed in vitro assays based on transgenic parasites has been their amenability to HTS [39,40]. In terms of impact, the adaptation of the T. cruzi β-galactosidase–based assay [26] to HTS standards established a milestone in the anti-Chagas early drug discovery process. After a protocol improvement to reduce the assay time, but still at its original 96-well plate format, the assay showed its reliability in the screening of a 2,000 compounds chemical library [27]. The β-galactosidase reporter assay was developed into a 384-well format with a luminescent readout allowing higher throughput for the HTS campaign at the Broad Institute, where screening of 303,224 compounds (the NIH library) yielded 4,394 hits. After the confirmation of activity and determination of efficacy and cytotoxicity of the hits, 3,005 compounds with an IC₅₀ <10 μM and >10-fold selectivity against the parasite versus the host cells were selected [39]. All information, including the chemical structures of the hits, is publicly available at Pubchem (AID 1885). Further analysis of the hits selected by this HTS revealed that a high percentage of them were not reproducible in a similar assay. It is likely that the reason for such discrepancies resided in the quality of the chemical compounds used for the original HTS [41]. Another HTS of a diversity-oriented synthesis chemical library of 77,312 compounds has been performed [42]. The most attractive confirmed hits in terms of potency, selectivity, and predicted drug-like physicochemical properties between these two HTSs were picked up for further development and the data obtained from them have served as the basis for different initiatives (Figure 1) [43–46]. The access to high-quality large chemical libraries will be of fundamental importance for the advance of drug discovery against neglected diseases, as it will be to get the resources and know-how to mine and test them. In this perspective, collaboration between pharmaceutical companies and academic laboratories becomes a key aspect for the successful development of future drugs.

Image-based HTS assays

High content screening (HCS) allows the automated acquisition and analysis of multiple cellular features simultaneously [30,46]. HCS is a powerful screening tool, the use of which has expanded in the last years because of high assay throughput, automatic image acquisition and analysis provided by user-friendly software interfaces and the possibility of robotization [30,46]. The availability of this technology has already permitted the development of image-based assays to identify active compounds against Leishmania donovani [34,47–49]. Similarly to HTS based on recombinant reporter parasites, automatic quantification is not biased by the observer visual parasite counting, as it is performed by an image analysis script specifically developed for this purpose [31,32], which once set up has been shown to be extremely useful, providing very accurate readings and saving valuable time [50]. In the drug development process against T. cruzi, this methodology holds a clear advantage compared to reporter-based assays as it offers more versatility, which may allow the use, upon assay adaptation, of any parasite strain that is adapted for infection in vitro (including difficult-to-engineer strains and relevant clinical specimens). Additionally, this assay provides the information on host cell cytotoxicity of each compound in a single experiment, delivering immediate results on the selectivity index. An image-based assay developed by Engel and coworkers that has already been used to screen a FDA-approved drugs library (>900 compounds) where 55 hits were identified [31], and at least two other image-based assays have been referenced in perspective articles [33,36]. More recently, scientists from Institute Pasteur in Korea have developed an algorithm to aid in the image analysis process of a T. cruzi infection drug inhibitory assay [32].

There are yet some disadvantages related to HCS that need to be improved, like the storage and handling of the high amount of data generated and the speed of both picture acquisition and computing analysis that can have an impact on the developed assays HTS scale up [46,51]. A drawback specific to the report anti-T. cruzi image-based assay is that it does not allow serial readouts, since cells must be fixed and stained at a single endpoint. This is similar to β-galactosidase and luciferase reporter-based assays; however, in the imaging assays, the possibility to measure different cell biology features (high-content microscopes offer up to four laser channels and there is a large spectrum of available fluorophores and biological probes) in a single experiment should grant its further development. Besides the identification of new hits, HCS could also provide a major input in the hit-to-lead-optimization process and play a key part in deciphering new mechanisms of action [30,51]. In the future, fluorescently tagged ligands or specific staining with antibodies could be incorporated in the assays to provide mechanistic information of compounds.

In vivo intermediate throughput screening by image acquisition and shortened protocols

A fundamental step in the drug discovery path is the selection and qualification of hits to be progressed into leads [52]. A requirement for this upgrade is consistent efficacy in animal models [29,52]. Although various animal infection models have been studied in Chagas disease research, by far the most used has been the mouse model [53]. The mouse is the most preferred in vivo model because it adequately reproduces the main pathogenic
features of human *T. cruzi* infection, there is accumulated knowledge about its genetics and physiology, a large amount of reagents and tools developed to work with it, and the ease of handling and storage of colonies. Transgenic parasites have also been used with success for in vivo studies. Bio-imaging of mouse models infected with transgenic *T. cruzi* strains (either luciferase or fluorescent recombinant parasites) allows a rapid assessment of the infectious disease process within the animals [28,38,41]. Before the implementation of transgenic parasites and in vivo imaging techniques, animals had to be killed to obtain end-point data on parasite detection and organ dissemination, which was very labor-intensive, presented risk for human infection, and required the use of large animal groups, all contributing to increased costs [53,54]. Furthermore, the techniques used (PCR, in situ hybridization, and microscopic blood or tissue sections parasitic counting) were cumbersome and presented certain limitations [38]. By means of luminescent or fluorescent transgenic parasites and the appropriate detection equipment, it is now possible to accurately evaluate in vivo the infection process and the drug responses that might inhibit it. In comparison with measurements relying on sacrifice, bio-imaging has the great advantage of providing continuous readouts on anesthetized animals, which greatly contributes to better understanding of infection dynamics and parasite tropism [55,56]. Interestingly, *T. cruzi* chronic infections in mice were focal but moved within the body from week to week. Parasites did not localize to the heart, despite these mice developing miocarditis and cardiac fibrosis. *T. cruzi* was found in other organs and consistently in the gastrointestinal tract [56].

Another advantage of these experimental infections for drug development is that, since progression of infection can be followed in live animals, the total number of mice required for the experiments can be drastically reduced, therefore following the 3R principle (Replacement, Reduction, and Refinement) [57], and contributing to reducing research costs [38,54]. With such advantages, in vivo protocols have already been developed [28,29,38,41,54] and are being used to test in vitro–selected hits with the best drug-like characteristics [28,41]. Early tagging of hits in the drug development path as *in vivo*–active, i.e., prior to medicinal chemistry modifications, represents considerable savings in time and funding, which are much needed to further develop the selected compounds into drugs for a disease with serious funding shortages [29]. Some of these assays are available to the community through a fee-for-service, non-profit core facility [58].

Bio-imaging assays normally model the initial acute-phase infection of Chagas disease and would select for fast-acting compounds [54]. Since the parasite detection threshold for *T. cruzi* in bio-imaging assays normally falls around 1,000 parasites [28,41,56], to determine whether a particular treatment, administered either in acute or chronic phase, results in sterile clearance, chemically induced immune suppression [39] should be used after treatment to confirm it [54].

The throughput of some in vivo assays [28] can be considered intermediate, since rapid two-day assays in low number of animals

---

**Figure 1. Anti-*Trypanosoma cruzi* compounds.** Chemical structures of anti-*T. cruzi* compounds currently in clinical use (*) and currently or recently in clinical trials (N), and of the most promising compounds retrieved out of the Broad Institute HTS campaign [44] and the screening of a smaller, diversity-oriented chemical library [39,43] (#). doi:10.1371/journal.pntd.0003259.g001
Key Learning Points
- Genetically engineered parasites have had a revolutionary impact in the development of both high-throughput amenable assays to screen large chemical collections for drug discovery in vitro and in vivo screening protocols to rapidly progress hit compounds.
- High technology equipment, such as automated plate readers, high-content microscopy, and luminescence imagers for live animals, has been applied to accelerate new anti-parasitic drugs search, providing high-quality readouts and reducing drug development costs.
- Collaborative frameworks between academia and the private pharmaceutical industry are necessary to bring together parasite biology and drug discovery expertise to allow the translation of academic research into medicines.

(fewer than 3 per group) can be used for screening hundreds of compounds in short periods of time [29].

Concluding Remarks
The last few years have seen a revolutionary improvement of the in vitro and in vivo screening methodologies to search for new anti-Chagas therapies based on the development of transgenic T. cruzi parasites engineered to constitutively express reporter genes and on the access to imaging technology that can largely ease up the process. These advances have granted the availability of robust in vitro assays suitable for high-throughput screening, as well as providing the necessary tools to design fast and reliable in vivo testing protocols. Their synergy is aimed at accelerating the early drug discovery path against a disease that has waited a long time for better therapeutics.

Very scarce economical resources are dedicated to find new drugs against neglected diseases, and therefore discovery campaigns must necessarily be cost-effective. However, the prospects are promising because the necessary tools are ready to be used and big pharmaceutical and public–private partnerships have shown their interest in the screening of large compound libraries [60], some of which are already under development [40]. It is expected that a substantial number of hits will sprout from these screens in the near future. Solid funding commitments at national and international levels need to be sustained to achieve the goal of a nontoxic, short-course, effective cure for chronic Chagas disease.

References

1. Coura JR, Vinas PA (2010) Chagas disease: a new worldwide challenge. Nature 465: 86–7.
2. Tarleton RL, Reithinger R, Urbina JA, Kirton U, Gurrler RE (2007) The challenges of Chagas Disease—grim outlook or glimmer of hope? PLoS Med 4: e332.
3. Hotze PJ DE, Woe-Coburn L, Serpa JA, Bezek S, et al. (2012) Chagas Disease: ‘The New HIV/AIDS of the Americas’. PLoS Negl Trop Dis 6: e1498.
4. Rassi A Jr, Rassi A, Marin-Neto JA (2010) Chagas disease. Lancet 375: 1388–1402.
5. Wilkinson SR, Kelly JM (2009) Trypanocidal drugs: mechanisms, resistance and new targets. Expert Rev Med Med 11: e31.
6. WHO (2014) Chagas disease (American trypanosomiasis). Available: http://www.who.int/mediacentre/factsheets/fs340/en/. Accessed 10 November 2014.
7. Machado-de-Assis GF, Diniz GA, Montoya RA, Dias JC, Coura JR, et al. (2013) A serological, patho-parasitological and clinical evaluation of untreated Chagas disease patients and those treated with benznidazole before and thirteen years after intervention. Mem Inst Oswaldo Cruz 108: 873–880.
8. Rodriguez-Guirrieneau L, Posay-Barbe KM, Monsonis-Cabedo M, Juncosa-Morros T, Diana A (2014) Pediatric Chagas disease in Europe: 45 cases from Spain and Switzerland. Pediatr Infect Dis J 33: 450–462.
9. Andrade MC, Oliveira Mle F, Nagaos-Dias AT, Coelho IC, Cândido Dda, S. et al. (2013) Clinical and serological evolution in chronic Chagas disease patients in a 4-year pharmacotherapy follow-up: a preliminary study. Rev Soc Bras Med Trop 46: 766–770.
10. Bortoluci GL, Viviano CA, Lococo BG, Petti MA, Viotti RJ (2013) Clinical characteristics and outcome of 107 adult patients with chronic Chagas disease and parasitological cure criteria. Trans R Soc Trop Med Hyg 107: 372–376.
11. Viotti R, Alarcén de Noya B, Araujo-Jorge T, Grijalva SJ, Guhl F, et al. (2014) Towards a paradigm shift in the treatment of chronic Chagas disease. Antimicrob Agents Chemother 58: 635–639.
12. Bustamante JM, Craft JM, Crowe BD, Ketchie SA, Tarleton RL (2014) New, combined, and reduced dosing treatment protocols cure Trypanosoma cruzi infection on mice. J Infect Dis 209: 150–162.
13. Marin-Neto JA, Rassi A Jr, Morillo CA, Aveam A, Connolly SJ, et al. (2008) Rationale and design of a randomized placebo-controlled trial assessing the effects of etiological treatment in Chagas’ cardiomyopathy: the benznidazole evaluation for interrupting trypanosomiasis (BENEFIT). Am Heart J 156: 37–43.
14. Bayer (2014) Study to Assess Bioequivalence of 30 and 120 mg Nifurtimox Tablets in Chronic Chagas’ Patients. ClinicalTrials.gov identifier: NCT01927224. Available: http://clinicaltrials.gov/show/NCT01927224. Accessed 5 November 2014.
15. Hospital Universitari Vall d’Hebron Research Institute (2013) Clinical Trial For The Treatment Of Chronic Chagas Disease With Posaconazole And Benznidazole (CHAGASAZOL). ClinicalTrials.gov identifier: NCT01162967. Available: http://clinicaltrials.gov/show/NCT01162967. Accessed 5 November 2014.
16. Merck Sharp & Dohme Corp. (2014) A Study of the Use of Oral Posaconazole (POS) in the Treatment of Asymptomatic Chronic Chagas Disease (POS267). ClinicalTrials.gov identifier: NCT01377480. Available: http://clinicaltrials.gov/show/NCT01377480. Accessed 5 November 2014.
17. Drugs for Neglected Diseases (2014) Proof-of-Concept Study of E1224 to Treat Adult Patients With Chagas Disease. ClinicalTrials.gov identifier: NCT01489228. Available: http://clinicaltrials.gov/show/NCT01489228. Accessed 5 November 2014.
18. Martinelli M, Rassi A Jr, Marin-Neto JA, de Paola AA, Berwanger O, et al. (2013) Chronic use of amiodarone against implantable cardioverter-defibrillator therapy for primary prevention of death in patients with Chagas cardiomyopathy: study rational and design of a randomized clinical trial. Am Heart J 166: 976–982.

Top Five Papers
1. Buckner FS, Verlinde CL, La Flamme AC, Van Voorhis WC (1996) Efficient technique for screening drugs for activity against Trypanosoma cruzi using parasites expressing beta-galactosidase. Antimicrob Agents Chemother 40: 2592–2597.
2. Bettiol E, Samanovic M, Murkin AS, Raper J, Buckner F, et al. (2009) Identification of three classes of heteroaromatic compounds with activity against intracellular Trypanosoma cruzi by chemical library screening. PLoS Negl Trop Dis 3: e384.
3. Broad Institute (2009) Luminescence cell-based/microorganism primary HTS to identify inhibitors of Trypanosoma cruzi replication. PubChem BioAssay AID 1885. Available: http://pubchem.ncbi.nlm.nih.gov/assay/assay.cgi?aid=1885. Accessed 5 November 2014.
4. Canavaci AM, Bustamante JM, Padilla AM, Perez Brandon CM, Simpson LJ, et al. (2010) In vitro and in vivo high-throughput assays for the testing of anti-Trypanosoma cruzi compounds. PLoS Negl Trop Dis 4: e740.
5. Engel JC, Ang KK, Chen S, Arkin MR, McKerrow JH, et al. (2010) Image-based high-throughput drug screening targeting the intracellular stage of Trypanosoma cruzi, the agent of Chagas disease. Antimicrob Agents Chemother 54: 3326–3334.
22. Pinazo MJ, Espinosa G, Gallego M, Lopez-Chejade PL, Urbina JA, et al. (2010) Successful treatment with posaconazole of a patient with chronic Chagas disease and systemic lupus erythematosus. Am J Trop Med Hyg 82: 363–367.

23. Andriani G, Chessler AD, Courtemanche G, Burleigh BA, Rodriguez A (2011) Activity in vivo of anti-Trypanosoma cruzi compounds selected from a high throughput screening. PLoS Negl Trop Dis 5: e1298.

24. Germain AR, Carmody LA, Dockendorff C, Galan-Rodriguez C, Rodriguez A, et al. (2011) Identification of small-molecule inhibitors of Trypanosoma cruzi replication. Bioorg Med Chem Lett 21: 7197–7200.

25. DNDi (14 November 2013) Drug Trial for Leading Parasitic Killer of the World. Available: http://ictmm2012.ioc.fiocruz.br/program_25_sept.html. Accessed 5 November 2014.

26. Buckner FS, Verlinde CL, La Flamme AC, Van Voorhis WC (2008) Efficient technique for screening drugs for activity against Trypanosoma cruzi using parasites expressing beta-galactosidase. Antimicrob Agents Chemother 40: 2592–2597.

27. Betiti E, Samanovic M, Murkin AS, Raper J, Buckner F, et al. (2009) Identification of three classes of heteroaromatic compounds with activity against intracellular Trypanosoma cruzi by chemical library screening. PLoS Negl Trop Dis 3: e384.

28. Canavaci AM, Bustamante JM, Padilla AM, Perez Brandan CM, Simpson LJ, et al. (2010) In vitro and in vivo high-throughput assays for the testing of anti-Chagas drug compounds. PLoS Negl Trop Dis 4: e740.

29. Rodriguez A, Tarleton RL (2012) Transgenic parasites accelerate drug discovery. Trends Parasitol 28: 90–92.

30. Brodin P, Christophe T (2011) High content screening in infectious diseases. Curr Opin Chem Biol 15: 334–339.

31. Engel JC, Ang KK, Chen S, Arkin MR, McKerrow JH, et al. (2010) Image-based high-throughput drug screening targeting the intracellular stage of Trypanosoma cruzi, the agent of Chagas’ disease. Antimicrob Agents Chemother 45: 3126–3133.

32. Moon S, Siqueira-Neto JL, Mosques CB, Yang G, Kang M, et al. (2014) An image-based algorithm for precise and accurate high throughput assessment of activity against the human parasite Trypanosoma cruzi. PLoS ONE 9(2): e87188.

33. Sykes ML, Avery VM (2013) Approaches to protozoan drug discovery: phenotypic screening. J Med Chem 56: 7727–7740.

34. De Meyder G, Ang KK, Arkin MR, Engel JC, McKerrow JH (2011) A screen against Leishmania intracellular amastigotes: comparison to a promastigote screen and identification of a host cell-specific hit. PLoS Negl Trop Dis 5: e1253.

35. Bustamante JM, Tarleton RL (2011) Methodological advances in drug discovery for Chagas disease. Expert Opin Drug Discov 6: 633–661.

36. Clayton J (2010) Chagas disease: pushing through the pipeline. Nature 465: S12–S15.

37. Taylor MC, Huang H, Kell JM (2011) Genetic techniques in Trypanosoma cruzi. Adv Parasitol 75: 231–250.

38. Hyland KV, Astfal SH, Olson CL, Daniels MD, Engman DM (2008) Bioluminescent imaging of Trypanosoma cruzi infection. Int J Parasitol 38: 1391–1400.

39. Carmody LG, Germain A, Barker D, Galan-Rodriguez C, Betiti E, et al. (2010) Identification of Small-Molecule Inhibitors of Trypanosoma cruzi Infection. In: (2010) Probe Reports from the NIH Molecular Libraries Program [Internet]. Bethesda, MD: National Center for Biotechnology Information (US). Available: http://www.ncbi.nlm.nih.gov/books/NBK55071/. Accessed 10 November 2014.

40. Rodriguez A (2013) HTS for T. cruzi infection of mammalian cells through GSK-Open Lab. 62nd Annual Meeting of the American Society of Tropical Medicine and Hygiene [Abstract]. Available: http://www.abstractsonline.com/PPlans/ViewAbstract.aspx?mID=3356&Key=57479e55bf-2171-43ef-94a6-1e8c7d474e22&Key=947e90c153c4-3e7e-27539bf8af&download=1&f=pdf&Key=cafe0f1a-9b33-4623-a1bb-85f311088f49#. Accessed 5 November 2014.