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Monica Cal, Jean-Robert Ioset, Matthias Fügen, Pascal Mäser, Marcel Kaiser

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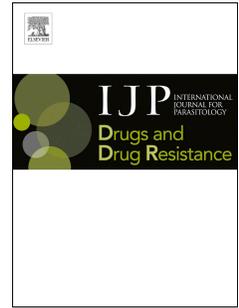
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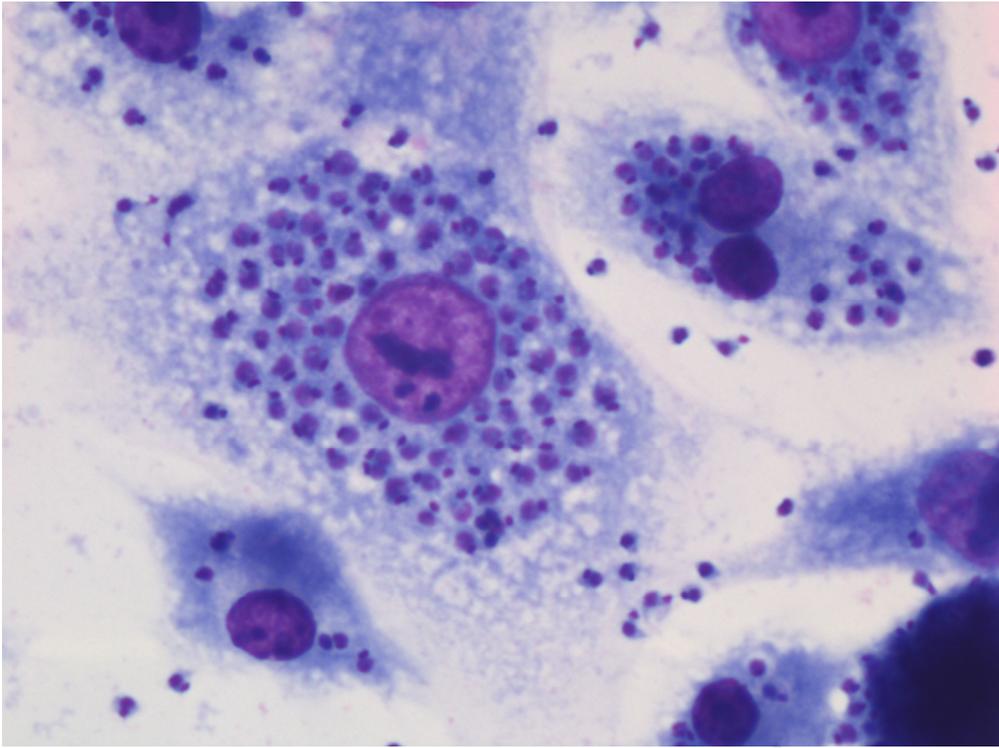
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ACCEPTED MANUSCRIPT

1 **Assessing anti-*T.cruzi* candidates *in vitro* for sterile cidalty**

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5 Monica Cal^{1,2}, Jean-Robert Ioset³, Matthias Fügi^{1,2}, Pascal Mäser^{1,2}, Marcel Kaiser^{1,2*}

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9 ¹ Swiss Tropical and Public Health Institute, CH-4051 Basel, Switzerland10 ² University of Basel, CH-4003 Basel, Switzerland11 ³ Drugs for Neglected Diseases *initiative*, CH-1202 Geneva, Switzerland

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15 *Corresponding author Swiss Tropical and Public Health Institute

16 Socinstrasse 57, 4051 Basel, Switzerland

17 E-mail marcel.kaiser@unibas.ch

18 Tel +41 61 284 8205

19 Fax +41 61 284 8101

20

21 **Abstract**

22 Total clearance of the *T. cruzi* infection – referred to herein as “sterile cure” – seems to be a critical
23 prerequisite for new drug candidates for Chagas disease, ensuring long-term beneficial effects for
24 patients in the chronic indeterminate stage. This requirement is notably supported by the recent
25 findings of clinical studies involving posaconazole and fosravuconazole, where the majority of
26 patients treated eventually relapsed after an apparent clearance of parasitaemia at the end of
27 treatment. We have adapted an *in vitro* system to predict the ability of a compound to deliver sterile
28 cure. It relies on mouse peritoneal macrophages as host cells for *Trypanosoma cruzi* amastigotes.
29 The macrophages do not proliferate, allowing for long-term testing and wash-out experiments.
30 Giemsa staining followed by microscopy provides a highly sensitive and specific tool to quantify
31 the numbers of infected host cells. Combining macrophages as host cells and Giemsa staining as the
32 read-out, we demonstrate that posaconazole and other CYP51 inhibitors are unable to achieve
33 complete clearance of an established *T. cruzi* infection *in vitro* in spite of the fact that these
34 compounds are active at significantly lower concentrations than the reference drugs benznidazole
35 and nifurtimox. Indeed, a few macrophages remained infected after 96 h of drug incubation in the
36 presence of CYP51 inhibitors– albeit at a very low parasite load. These residual *T. cruzi*
37 amastigotes were shown to be viable and infective, as demonstrated by wash-out experiments. We
38 advocate characterizing any new anti-*T. cruzi* early stage candidates for sterile cidal activity early in the
39 discovery cascade, as a surrogate for delivery of sterile cure *in vivo*.

40

41 **Keywords**42 *Trypanosoma cruzi*; *in vitro* assay; posaconazole; macrophage; Giemsa; Chagas disease

43 **1. Introduction**

44 *Trypanosoma cruzi* is the causative agent of Chagas disease, endemic in 21 countries of Latin
45 America (WHO fact sheet 340). Migration and travel have additionally contributed to the spread of
46 Chagas disease to other continents, including North America and Europe. The World Health
47 Organization estimates that 6-7 million people are infected world-wide, leading to more than 10,000
48 deaths annually (WHO fact sheet 340). In the absence of a vaccine, the only treatment option is
49 chemotherapy. However, the existing drugs benznidazole and nifurtimox have several limitations,
50 notably in relation to their severe adverse effects and contraindications (Andrade et al., 1992;
51 Bahia-Oliveira et al., 2000; Cancado, 2002; Urbina, 2010). Safer drugs are urgently needed. Over
52 the last few years, the development of new anti-*T. cruzi* agents has focused on azoles as inhibitors
53 of CYP450-dependent lanosterol demethylase (CYP51; 1.14.13.70) that act by blocking
54 trypanosomatid ergosterol synthesis (Buckner and Urbina, 2012; Chatelain, 2015). Azoles display
55 remarkable nanomolar range *in vitro* potency against *T. cruzi* as well as a good safety profile in
56 humans (Buckner and Urbina, 2012; Soeiro Mde et al., 2013). Azoles have been in use as antifungal
57 agents for decades, which has greatly facilitated their preclinical development for Chagas disease.
58 Recently, the two triazoles posaconazole and fosravuconazole, a prodrug of ravuconazole, were
59 tested in controlled clinical phase II studies (Urbina, 2015). However, both molecules failed to
60 show sufficient levels of efficacy in chronic Chagas patients; after an initial phase of apparent
61 clearance of parasitaemia following the end of treatment, 80% of the patients relapsed 10 months
62 after the end of treatment in the posaconazole (CHAGAZASOL) study (Molina et al., 2014), while
63 71% relapsed 12 months after the end of treatment in the fosravuconazole study (Torricco, 2013), as
64 determined by real time qPCR detection of *T. cruzi* DNA. These disappointing clinical results for
65 azoles contrast with the relatively low treatment failures observed with benznidazole, which showed
66 80% and 94% sustained clearance of parasites at the same endpoint in the posaconazole and
67 fosravuconazole trials respectively. The outcome of a clinical trial depends on several factors,
68 pharmacokinetics and host immune system play an important role. The obtained posaconazole

69 levels in patients treated with 100-400 mg b.i.d. were clearly below the plasma levels in mice
70 (20mg/kg/day) (Urbina, 2015). A prolonged treatment duration and a higher dose or an improved
71 formulation to get higher plasma levels could lead to a better clinical outcome. While the full
72 potential of azoles for combination chemotherapy has not yet been realized (Fügi et al., 2015), our
73 working hypothesis, derived from the disappointing outcome of the clinical trials, is that any novel
74 anti-*T. cruzi* hit or chemistry starting point emerging from the discovery pipeline should be assessed
75 at an early stage for its potential to deliver sterile cidal activity against different *T. cruzi* genotypes
76 (Moraes et al., 2014; Chatelain, 2015). Our aim is to develop an *in vitro* test for sterile cidal activity
77 towards *T. cruzi* amastigotes. Posaconazole and benznidazole can serve as benchmarks for such a
78 test (Fortes Francisco et al., 2015). Here we report the adaptation of *in vitro* assay protocols
79 (McCabe et al., 1983) that can be performed on any *T. cruzi* strain without requiring sophisticated
80 laboratory equipment, and the activity profiling of a small panel of CYP51 inhibitors in these assays
81 to investigate whether the clinical relapses observed following posaconazole and fosravuconazole
82 therapies could have been predicted, at least partly, from these simple *in vitro* surrogate assays.

83 **2. Materials and methods**

84 *2.1 Cells and media*

85 A *T. cruzi* Tulahuen C2C4 strain that expresses the β -galactosidase gene (LacZ) (Buckner et al.,
86 1996) was cultured in RPMI-1640 supplemented with 10% inactivated FBS (iFBS) and 2 μ M L-
87 glutamine at 37°C and 5% CO₂. L6 rat skeletal myoblast cells (ATCC CRL-1458) were used as host
88 cells for infection with transgenic *T. cruzi* trypomastigotes.

89 *2.2 Drugs*

90 Nifurtimox, posaconazole, fenarimol, clotrimazole, econazole, itraconazole, ketoconazole and
91 tioconazole were purchased from Sigma-Aldrich. Benznidazole was synthesized by Epichem Pty
92 Ltd, Murdoch, Australia. Stock solutions of these drugs were prepared in DMSO 100% at 10 mg/ml
93 and 1 mg/ml.

94 2.3 *LacZ/CPRG assay*

95 L-6 cells or murine peritoneal macrophages (MPM) were seeded, 1,000 cells per well (L6) or 4,000
96 cells per well (MPM), in 96-well microtiter plates (Costar) in 100 μ l RPMI 1640 with 10% heat-
97 inactivated FBS (Connectorate AG) and 2 μ M L-glutamine at 37°C and 5% CO₂. At 24 h the
98 medium was replaced by 100 μ l fresh medium containing 5,000 (L6) or 4,000 (MPM)
99 trypanomastigote *T. cruzi*. At 72 h the medium was replaced by 100 μ l (L6) or 200 μ l (MPM) fresh
100 medium with serial drug dilution from 30,000 to 0.5 ng/ml in 3-fold steps. After 96 h incubation,
101 the plates were inspected microscopically, followed by the addition of CPRG/Nonidet solution
102 (0.25 μ M Chlorophenol red- β -D-galactopyranoside and 0.25% Nonidet in PBS; 50 μ l per well).
103 After 5 h further incubation, the plates were read photometrically at 540 nm; IC₅₀ and IC₉₀ values as
104 well as the Hill factor were calculated by the four parameter nonlinear regression model using the
105 software Softmax Pro (Molecular Devices Cooperation, Sunnyvale, CA, USA). All values are
106 means from at least three independent assays.

107 2.4 *Giemsa assay*

108 4,000 MPM were seeded into 16-chamber slides (Lab-tek) (McCabe et al., 1983). Incubation,
109 infection with trypanomastigote *T. cruzi* at MOI of 1, and drug exposure were performed as described
110 above. After the 96 h drug exposure the medium was removed and the slides were fixed with
111 methanol for 10 min, followed by staining with 10% Giemsa solution (Sigma). The infection status
112 (infected/non-infected) of at least 400 cells was determined microscopically. If possible, the number
113 of intracellular amastigotes was counted for 100 infected cells. At low infection rates, the number of
114 intracellular amastigotes was determined for all infected cells. The infection rates of the untreated
115 controls were between 85% and 95%. The results were expressed either as a percentage of infected
116 host cells compared to untreated controls, or as mean numbers of amastigotes per infected
117 macrophage. The *T. cruzi* population size was calculated as the number of intracellular amastigotes
118 per 100 macrophages (% infection rate \times number of amastigotes per infected cell). IC₅₀ and IC₉₀
119 values, as well as the Hill factor, were calculated from the sigmoidal dose-response curve using the

120 four parameter nonlinear regression model of the software Softmax Pro (Softmax Pro Molecular
121 Devices Cooperation, Sunnyvale, CA, USA). All values are means from at least three independent
122 assays. For assessment of cidality, the medium was removed after 96 h drug exposure and the
123 adherent MPM were washed four times with 200 μ l fresh medium. 200 μ l fresh medium was added,
124 followed by a further 168 h incubation. The medium was then removed, and the slides were fixed
125 with methanol and stained with 10% Giemsa as described above.

126 3. Results and Discussion

127 3.1 Myoblasts vs. macrophages as host cells

128 *Trypanosoma cruzi* can infect practically every type of nucleated mammalian cell. Muscle cells, and
129 cardiomyocytes in particular, are among the cells targeted by *T. cruzi* that contribute to the
130 pathology of Chagas disease (Nagajyothi et al., 2012). Rat L6 are widely used as convenient and
131 relevant host cells for testing compounds against the intracellular amastigote stage of *T. cruzi*. The
132 standard assay requires 96 h exposure to a test compound followed by quantification of the parasites
133 (Buckner et al., 1996). Longer incubation times with this specific cell line are not possible because
134 L6 cells multiply with a population doubling time of about 15 h and the cultures would overgrow.
135 Instead, we used mouse peritoneal macrophages as a non-dividing type of host cell. The infection
136 rates of the macrophages with *T. cruzi* trypomastigotes were over 80%. The replication time of
137 intracellular *T. cruzi* amastigotes is 18-20 hours and was determined in previous experiments. A
138 selection of antifungal CYP51 inhibitors, 6 azoles plus the non-azole fungicide fenarimol, were
139 tested against *T. cruzi*, along with the standard drugs benznidazole and nifurtimox, in parallel in L6
140 cells and macrophages. A *LacZ* transfected Tuluahen strain (C2C4) was used for ease of read-out
141 with the chromogenic β -galactosidase substrate CPRG (Chlorophenol red- β -D-galactopyranoside).
142 All CYP51 inhibitors had IC_{50} values in the low nanomolar range, whereas the reference drugs
143 benznidazole and nifurtimox were clearly less potent, with IC_{50} values in the micromolar range
144 (Table 1, left two columns). Posaconazole was over 1,000-fold more potent than benznidazole in
145 both systems (Table 1). The IC_{50} values of all compounds tested were lower in macrophages than in

146 L6 cells ($p=0.004$; two-tailed Wilcoxon matched pairs test). IC_{90} values could not be determined
147 accurately (Table 1) as the variance at the tail of the dose-response curve was too high. Thus the
148 macrophages provided a highly sensitive test system, but the colorimetric signal from turn-over of
149 CPRG was a suboptimal read-out for accurately measuring signals close to background from small
150 numbers of residual *T. cruzi* amastigotes.

151 3.2 β -galactosidase vs. Giemsa staining as read-out

152 As a more accurate alternative to CPRG-based colorimetric readout, we tested Giemsa staining
153 followed by microscopic read-out to determine numbers of infected macrophages and, for selected
154 compounds, the parasite load of infected cells. This more labor-intensive method of detection
155 allowed the specific identification and accurate counting of *T. cruzi* amastigotes at the single cell
156 level (see the Graphical Abstract for a sample image). The resulting IC_{50} values based on the
157 infection rate were more than an order of magnitude higher than those determined by CPRG (Table
158 1). This can be explained by the fact that Giemsa staining determined the percentage of infected
159 macrophages (regardless of the number of intracellular amastigotes), whereas the fluorescence
160 signal from CPRG turn-over is proportional to the total number of amastigotes (regardless of the
161 percentage of infected host cells). The IC values derived by Giemsa were thus not proportional to
162 the total number of parasites, and the resulting IC_{50} values were comparably higher. The IC_{90} values
163 obtained by microscopic determination of the infection rate following Giemsa staining are,
164 however, in our opinion more meaningful than those obtained with CPRG, because single surviving
165 amastigotes can be detected by microscopic counting. Figures 1 & 2A compare the two different
166 kinds of dose-response curves. The reference drugs benznidazole and nifurtimox exhibited a smaller
167 shift in IC_{50} from a decrease in overall parasitaemia (CPRG read-out) to a decrease in infected host
168 cells (Giemsa read-out) than the seven antifungals. Furthermore, the Hill factors (HF) of the dose-
169 response curves for the reference drugs were higher for the Giemsa read-out while the CYP51
170 inhibitors exhibited flatter curves for the Giemsa read-out than for CPRG (mean Hill factors of 0.90
171 vs. 1.8; $p=0.047$; two-tailed Wilcoxon matched pairs test). Importantly, the Giemsa dose-response

172 curves of the reference drugs (Figure 2A) reached base-line, while those of the antifungals (Figure
173 2A) did not. Thus none of the CYP51 inhibitors tested led to a 100% clearance of parasitaemia from
174 the infected cell population-. A small percentage of host cells remained infected, although with a
175 very low parasite load of 1-4 parasites per infected cell, compared to >10 parasites in untreated
176 infected control cells. The parasite load of untreated cells was very high 96 h post infection (>10
177 amastigotes/cell). For the purpose of Figure 2B, the average parasite load was counted up to 10
178 amastigotes/cell. The drop in mean parasite load is reflected by the drop in the CPRG signal. The
179 CYP51 inhibitors showed a clear reduction of the parasite load per infected cell, which settled at a
180 low load (Figure 2B). The standard drug benznidazole tends to show a continuous reduction of the
181 parasite load. All compounds were tested to the maximum non-cytotoxic concentration. In Figure
182 2C, the calculated total number of intracellular *T. cruzi* amastigotes in 100 macrophages is plotted
183 against drug concentration. This clearly shows the flat dose-response of CYP51 inhibitors and their
184 inability to kill all parasites within 96 h of drug exposure. The drug exposure period of 96 h
185 corresponds to 4-5 parasite generations. This should be long enough for a complete depletion of the
186 preformed pool of sterols, which is a prerequisite for a cidal effect of the ergosterol biosynthesis
187 inhibitors (Urbina et al 1998). However, we cannot exclude that an even longer exposure would kill
188 all the intracellular *T. cruzi* amastigotes. We calculated that >10 parasites per 100 macrophages
189 survive at the highest concentrations of CYP51 inhibitors. Benznidazole showed a good dose-
190 response curve and at a concentration of 40 μ M, only <1 parasite per 100 macrophages survived
191 (Figure 2C). These results correlate with the previous observations made using high content
192 imaging technology in a panel of *T. cruzi* I-VI genotype strains (Moraes et al., 2014), validating the
193 inability of CYP51 inhibitors to lead to full clearance of *T. cruzi* amastigote populations following
194 drug exposure, independent of the drug concentration tested.

195 3.3 A long-term *in vitro* assay for sterile cidalty

196 To assess whether these few residual intracellular *T. cruzi* amastigotes were viable, we prolonged
197 the *in vitro* test to a total duration of 11 days. The drug-containing medium was removed after 96 h

198 incubation, the adherent macrophages were washed twice, and fresh, drug free medium was added.
199 After another 168 h incubation, the numbers of infected macrophages, and for selected compounds
200 the parasite load of the infected cells, were determined by Giemsa staining. For benznidazole and
201 nifurtimox, the resulting IC_{50} values only increased slightly (10-20%) compared to those after 96 h
202 (Table 1, right). In contrast, the majority of the CYP51 inhibitors tested exhibited a strong shift
203 ($>200\%$) in IC_{50} from 96 h to 96+168 h tests. IC_{90} could not be determined, because the dose-
204 response curves of the CYP51 inhibitors did not reach background levels. Figure 3A illustrates the
205 shift in IC values based on the percentage of infected cells. Figures 3B and C show the
206 corresponding mean number of intracellular amastigotes per infected cell and the estimated total
207 number of parasites per 100 macrophages, respectively. The dose-response curves of the antifungals
208 are flat in the 96 h drug exposure experiment (Figure 2A-C) whereas the curves are steeper in the
209 wash-out experiment (Figure 3A-C). The wash-out experiments confirmed that the few residual
210 amastigotes observed after 96 h incubation were viable and infective to other macrophages. Thus
211 the tested antifungals posaconazole, clotrimazole, econazole, ticonazole and itraconazole were not
212 able to deliver sterile cidalty *in vitro*.

213 4. Conclusion

214 Mouse macrophages provide a highly sensitive system for testing molecules *in vitro* against *T. cruzi*
215 intracellular amastigotes. Combined with Giemsa staining and microscopic read-out, this system
216 allows drug sensitivity tests over long periods of time with high sensitivity and specificity, detecting
217 single residual parasites. Wash-out experiments demonstrated that these residual parasites were
218 alive and able to infect new host cells. When the percentage of infected host cells was quantified by
219 Giemsa staining, all the CYP51 inhibitors tested displayed lower IC_{50} values than the reference
220 drugs benznidazole and nifurtimox, but the dose-response curves were much flatter and did not
221 reach baseline, showing that the CYP51 inhibitors were unable to clear all *T. cruzi* amastigotes.
222 This indicates that measuring drug potency in terms of IC_{50} based on viability markers is an
223 insufficient readout to predict for clearance of parasitaemia *in vitro* and – by extension – *in vivo*, in

224 particular if the parasites are persistent and the compound's effect is rather static than cidal. By
225 using macrophages as host cells and Giemsa staining as the read-out, it was indeed possible to
226 demonstrate that the azoles, as well as another non-azole based CYP51 inhibitors, do not deliver
227 sterile cidity defined as full clearance of parasitaemia *in vitro*. To evaluate the anti-*T.cruzi*
228 potential of a compound, in our opinion both the reduction in infection rate as well as the reduction
229 in the total number of parasites is important. The approach presented may provide early stage
230 evidence – and potentially offer a predictive preclinical tool – for the observed parasitological
231 relapse of Chagas patients treated with azoles. The long-term incubation assay, relying on mouse
232 peritoneal macrophages as host cells and Giemsa staining as the read-out, is not amenable to high
233 throughput screening. It can be implemented as a secondary assay to profile and eventually
234 prioritize anti-*T.cruzi* hits identified from screening campaigns, or it may serve as the starting point
235 for developing a high-content assay amenable to higher throughput screening. Wash-out
236 experiments demonstrated that the *T. cruzi* amastigotes that persist following *in vitro* drug exposure
237 with CYP51 inhibitors are viable and replicating. While we cannot exclude that drugs which are
238 unable to deliver sterile cidity *in vitro* might still work *in vivo* thanks to the contribution of the
239 host's immune system or any other factor not captured under the experimental conditions of our
240 assay, we hope that the presented assay will contribute to render the discovery cascade for Chagas
241 disease more predictable.

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254

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311 **Figure 1. *In vitro* drug sensitivity of *T. cruzi* amastigotes in mouse macrophages after 96 h**
312 **drug exposure.** The solid lines represent the azoles and the dashed lines represent the reference
313 drugs benznidazole and nifurtimox. The Hill factors (HF) of the dose response curves are indicated
314 in the legend. Percentage growth of the parasites as quantified by colorimetric read-out of CPRG
315 turn-over.

316
317 **Figure 2. *In vitro* drug sensitivity of *T. cruzi* amastigotes in mouse macrophages after 96 h**
318 **drug exposure.** The solid lines represent the azoles and the dashed lines represent the reference
319 drugs benznidazole and nifurtimox. Hill factors (HF) of the dose response curves are indicated in
320 the legend. A) Percentage infected host cells as quantified by microscopic read-out of Giemsa
321 stains. B) Mean numbers of amastigotes per infected macrophage as quantified by microscopic
322 read-out of Giemsa stains. C) Estimated numbers of intracellular amastigotes in 100 macrophages.
323 Calculation: % infection rate x mean number of amastigotes per infected cell.

324
325 **Figure 3. *In vitro* drug sensitivity of *T. cruzi* amastigotes in mouse macrophages after drug**
326 **washout (96 h) and 168 h recovery period.** The solid lines represent the azoles and the dashed
327 lines represent the reference drugs benznidazole and nifurtimox. Hill factors (HF) of the dose
328 response curves are indicated in the legend. A) Percentage infected host cells as quantified by
329 microscopic read-out of Giemsa stains. B) Mean numbers of amastigotes per infected macrophage
330 as quantified by microscopic read-out of Giemsa stains. C) Estimated numbers of intracellular
331 amastigotes in 100 macrophages. Calculation: % infection rate x mean number of amastigotes per
332 infected cell.

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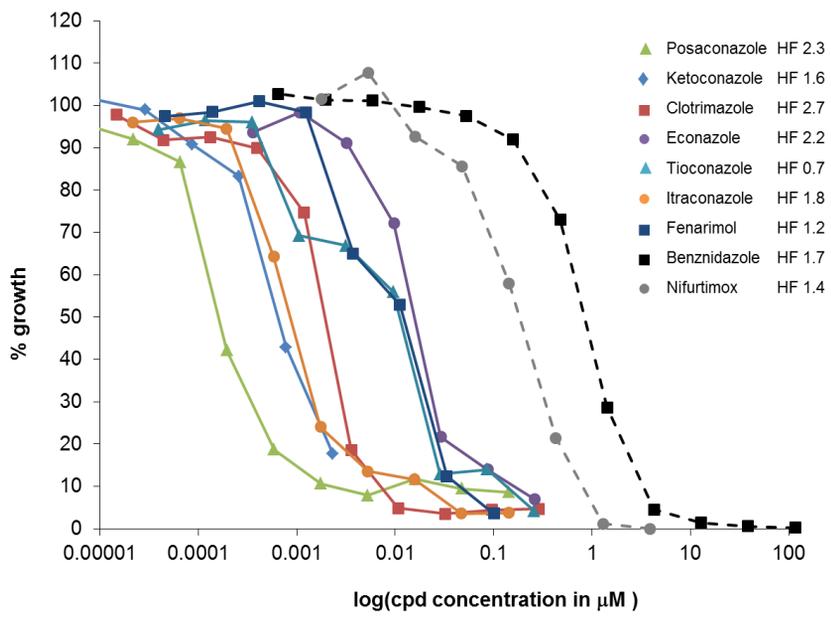
337 **Table 1. *In vitro* drug sensitivity tests for *T. cruzi*.** Values are the means of at least 3 replicates \pm
 338 S.D. in nM (L6, rat L6 myoblasts; MPM, murine peritoneal macrophages; CPRG, Chlorophenol
 339 red- β -D-galactopyranoside).

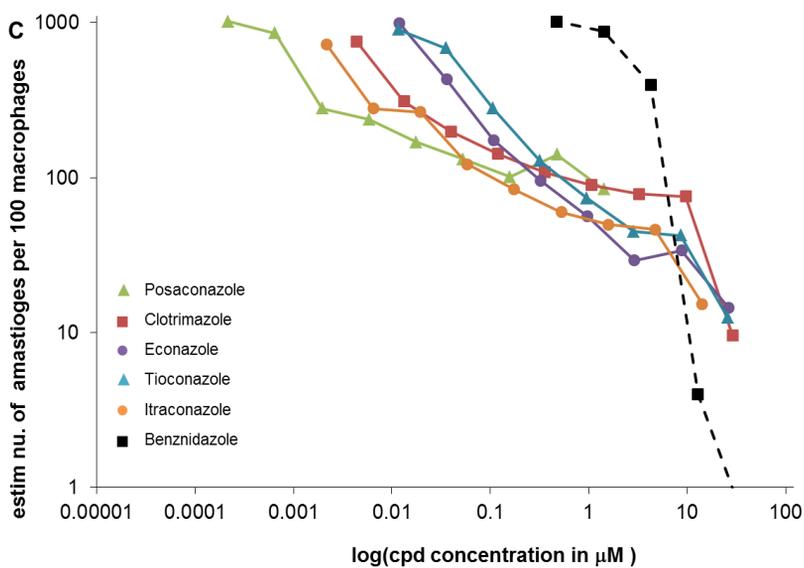
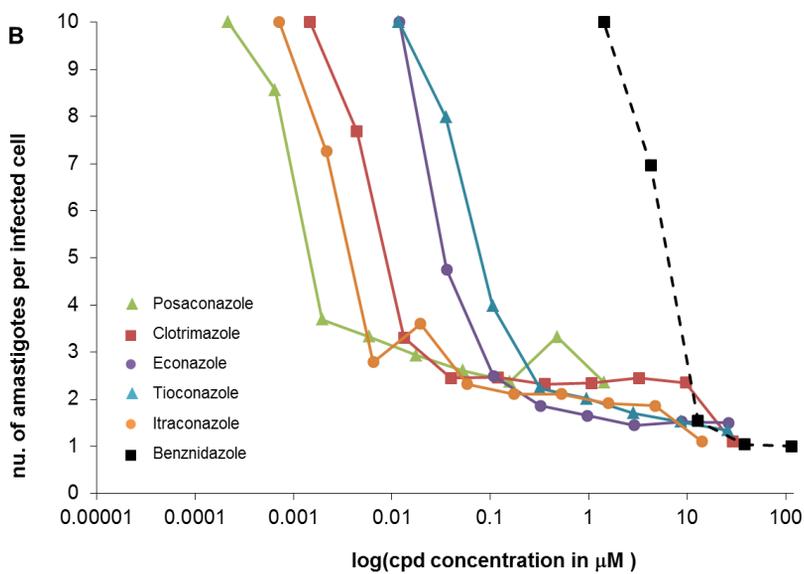
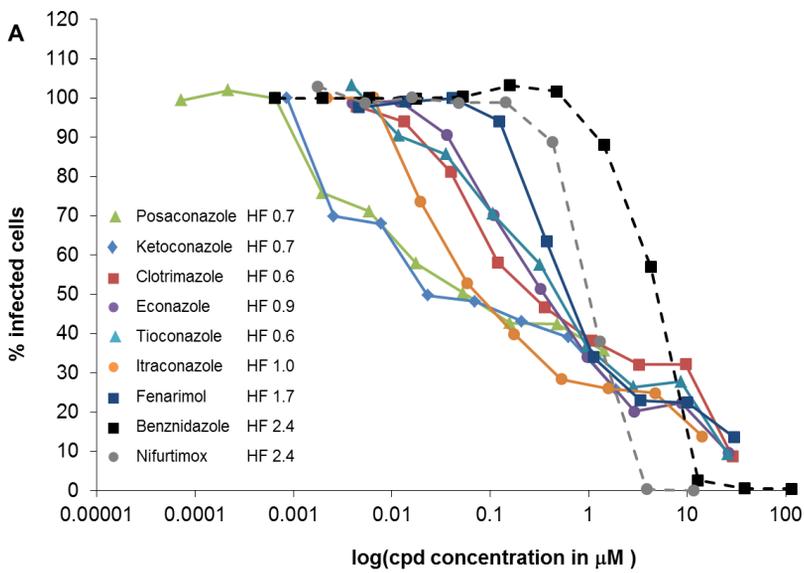
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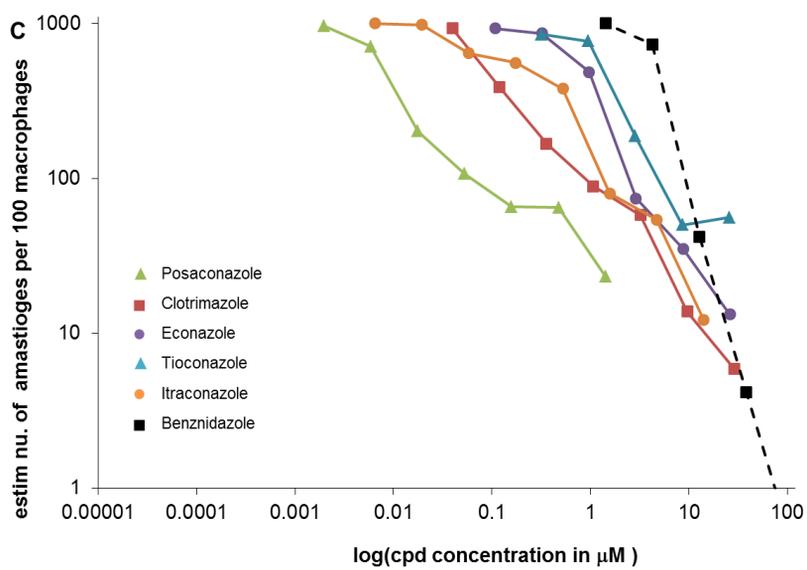
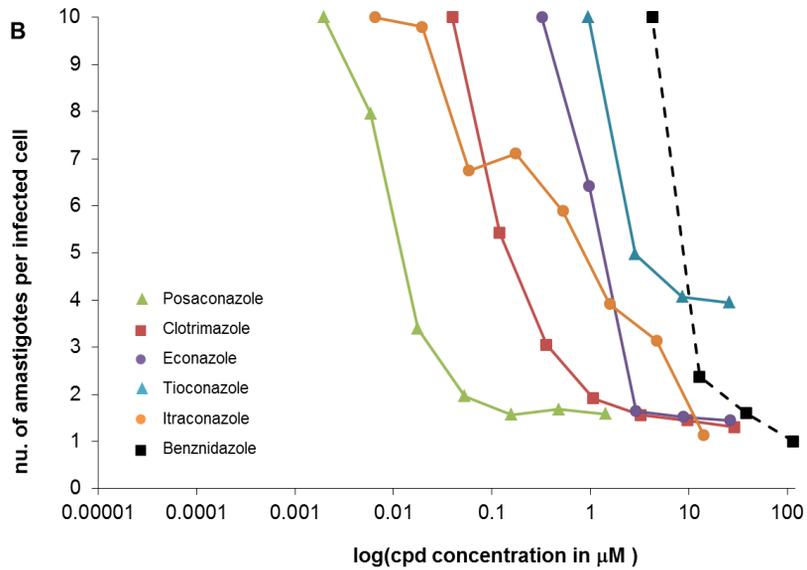
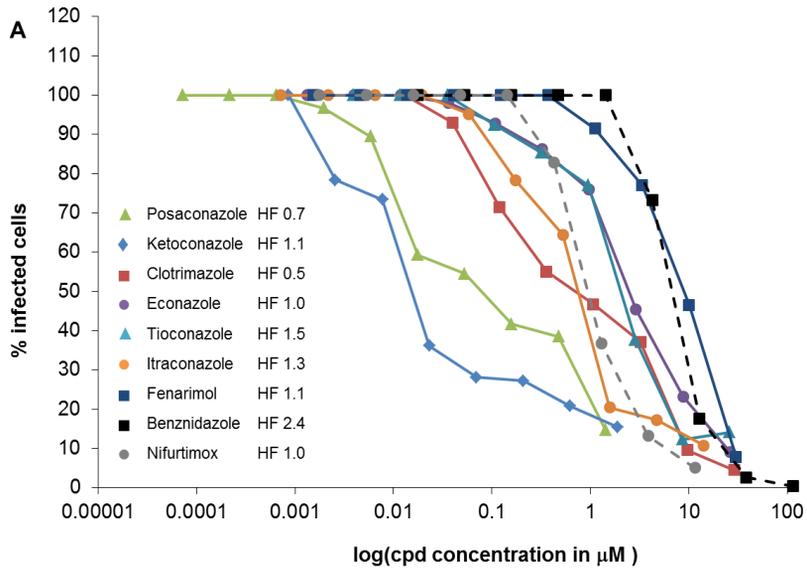
		L6	MPM	MPM	MPM
		96h, CPRG	96h, CPRG	96h, Giemsa	96+168h, Giemsa
Benznidazole	IC50	1,800 \pm 540	850 \pm 190	4,800 \pm 1,100	5,800 \pm 2,900
	IC90	6,600 \pm 2,800	3,100 \pm 790	11,000 \pm 650	18,000 \pm 9,700
Nifurtimox	IC50	470 \pm 160	180 \pm 45	990 \pm 200	1,100 \pm 720
	IC90	2,200 \pm 760	770 \pm 7	2,700 \pm 290	4,300 \pm 3,900
Posaconazole	IC50	0.7 \pm 0.3	0.3 \pm 0.07	46 \pm 2.8	280 \pm 460
	IC90	57 \pm 49	1.4 \pm 1.0	>1,400	>4,300
Ketoconazole	IC50	1.7 \pm 1.5	0.4 \pm 0.4	34 \pm 35	n/a
	IC90	67 \pm 64	1.7 \pm 0.4	>1,900	n/a
Clotrimazole	IC50	20 \pm 15	1.7 \pm 0.6	470 \pm 440	4,200 \pm 520
	IC90	310 \pm 190	7.3 \pm 2.9	>29,000	>29,000
Econazole	IC50	60 \pm 11	16 \pm 5.2	320 \pm 210	1,100 \pm 130
	IC90	1,300 \pm 530	170 \pm 24	18,000 \pm 9500	>26,000
Tioconazole	IC50	72 \pm 10	10 \pm 7.8	500 \pm 410	1,400 \pm 110
	IC90	970 \pm 360	21 \pm 26	>26,000	>26,000
Itraconazole	IC50	5.7 \pm 4.3	0.9 \pm 0.4	99 \pm 58	1,200 \pm 150
	IC90	69 \pm 51	5.7 \pm 4.3	>1,400	>14,000
Fenarimol	IC50	160 \pm 6.0	12 \pm 9.1	540 \pm 57	5,700 \pm 5,000
	IC90	>300	33 \pm 39	>30,000	>30,000

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Highlights (max 85 char.)

- A highly sensitive *Trypanosoma cruzi* in vitro drug test for sterile cidalty
- This test demonstrates failure of CYP51 inhibitors to deliver sterile cure

ACCEPTED MANUSCRIPT