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Identification of Pyrazolo[3,4-e][1,4]thiazepin based CYP51 inhibitors as potential Chagas disease therapeutic alternative: *In vitro* and *in vivo* evaluation, binding mode prediction and SAR exploration

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1 **Identification of Pyrazolo[3,4-e][1,4]thiazepin based CYP51 inhibitors as**
2 **potential Chagas disease therapeutic alternative: *In vitro* and *in vivo* evaluation,**
3 **binding mode prediction and SAR exploration**

4
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20 **Running title:** Pyrazolo[3,4-e][1,4]thiazepin activity against *T. cruzi*

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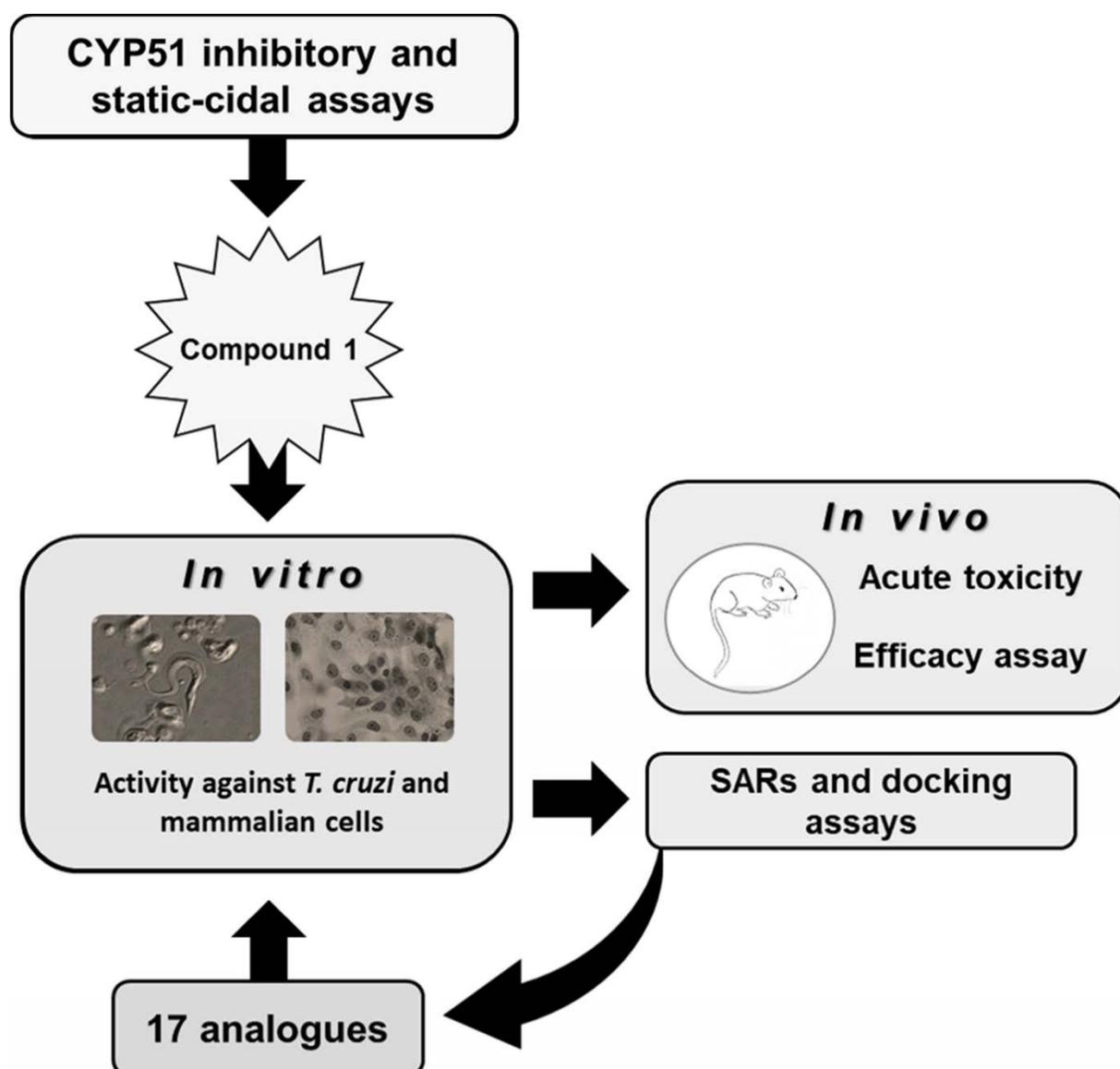
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23 **Highlights**

- 24 • Structure based studies predicted the binding of novel CYP51 inhibitor with CYP51Tc.
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26 • The compounds **1** and **1f** displayed trypanocidal effect upon *T. cruzi in vitro*.
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28 • Compound **1f** gave no effect in vivo, while compound **1** reduced the parasitemia peak.

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30

31 **Graphical abstract**

32

33 **Abstract**

34 American trypanosomiasis or Chagas disease (CD) is a vector borne pathology caused by the
35 parasite *Trypanosoma cruzi* (*T. cruzi*), which remains a serious global health problem. The
36 current available treatment for CD is limited to two nitroderivatives with limited efficacy and
37 several side effects. The rational design of ergosterol synthetic route inhibitors (e.g. CYP51
38 inhibitors) represents a promising strategy for fungi and trypanosomatids, exhibiting excellent
39 anti-*T.cruzi* activity in pre-clinical assays. In the present work, we evaluate through different
40 approaches (molecular docking, structure activity relationships, CYP51 inhibitory assay, and
41 phenotypic screenings *in vitro* and *in vivo*) the potency and selectivity of a novel CYP51
42 inhibitor (compound **1**) and its analogues against *T.cruzi* infection. Regarding anti-parasitic
43 effect, compound **1** was active *in vitro* with EC₅₀ 3.86 and 4.00 μM upon intracellular
44 (Tulahuen strain) and bloodstream forms (Y strain), respectively. *In vivo* assays showed that
45 compound **1** reduced in 43 % the parasitemia peak but, unfortunately failed to promote animal
46 survival. In order to promote an enhancement at the potency and pharmacological properties,
47 17 new analogues were purchased and screened *in vitro*. Our findings demonstrated that five
48 compounds were active against intracellular forms, highlighting compounds **1e** and **1f**, with
49 EC₅₀ 2.20 and 2.70 μM, respectively, and selectivity indices (SI) = 50 and 36, respectively.
50 Against bloodstream trypomastigotes, compound **1f** reached an EC₅₀ value of 20.62 μM, in a
51 similar range to Benznidazole, but with low SI (3). Although improved the solubility of
52 compound **1**, the analogue **1f** did not enhance the potency *in vitro* neither promote better *in*
53 *vivo* efficacy against mouse model of acute *T.cruzi* infection arguing for the synthesis of novel
54 pyrazolo[3,4-e][1,4]thiazepin derivatives aiming to contribute for alternative therapies for CD

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57 **Key-words:** Chagas disease, *Trypanosoma cruzi*, CYP51 inhibitors and SAR studies.

58

59 1. Introduction

60 Chagas disease (CD) is a parasitic infection caused by the protozoan *Trypanosoma*
61 *cruzi* (*T. cruzi*). Considered, as a worldwide public health problem due to migration
62 flows and non-vector transmission routes, is estimates that about 6 to 7 million people
63 are carries of *T. cruzi* and an average of over 12,000 people die every year [1, 2]. The
64 main mechanism of transmission includes: classical vectorial transmission by
65 triatomine bugs also known as blood sucking bugs, via congenital when the parasite is
66 transmitted from mother to child, by oral contamination through ingesting the parasite
67 on contaminated food or drink and through iatrogenic that comprehend blood
68 transfusion or organ transplantation [3]. In many endemic countries like Brazil, vector
69 and blood bank control measures lead to a drastic decreased in the number of new
70 cases via classical vectorial route however, in last decades the oral transmission has
71 emerged with epidemiological relevance in the Amazon region [2]. The CD has two
72 phases, the acute phase appears soon after infection and is characterizing by the
73 patent parasitemia and mostly displaying oligosymptoms ranging from flu-like to
74 intense myocarditis. Due to immune host response, there is a control of the parasite
75 load and most patients move to a later stage called the chronic phase. Although most
76 stay at the indeterminate stage, about 30-40% may develop symptoms after 10 to 20
77 years post infection mainly related to cardiomyopathy associated or not to
78 gastrointestinal effects [1,4]. The CD available treatment is restricted to two oral
79 nitroheterocyclic compounds, nifurtimox and benznidazole (Bz) both introduced in
80 clinical more than 50 years ago. Even though 25 million people are at risk of infection,
81 according to the WHO, less than 1% of infected patients receive treatment [5]. Thus,
82 the development of new effective drugs for Chagas disease is urgently needed, a

83 neglected pathology that mostly affects poor regions, and with low interest by the
84 pharmaceutical companies due to the low monetary welfares. In addition, other
85 limitations of using the current nitro-derivative drugs are: varying results according to
86 the disease stage, treatment period and dose, age and geographical origin of the
87 patients, side effects, besides a natural resistance profile of some parasite strains
88 against nitro-derivatives [1, 6, 7, 8]. In order to face the limitations of the available
89 therapy, several *in vitro* and *in vivo* studies of potentials new drug candidates for CD
90 have been performed; some of these studies involving inhibitors of the ergosterol
91 biosynthesis pathway. Differently from mammalian cells, in fungi and trypanosomatids,
92 the cytochrome P450 enzyme or sterol 14 α -demethylase (CYP51) pathway leads to
93 formation of ergosterol-like products, which are essential for the survival of these
94 parasites, producing viable membranes and making possible cell growth and division
95 [9,10]. Recently, two CYP51 inhibitors first introduced as antifungal, Posaconazole
96 and E1224 (the prodrug of Ravuconazole) and that later showed *in vitro* and *in vivo*
97 efficacy against *T. cruzi* were moved for clinical trials upon chronic Chagas disease
98 patients. Unsuccessfully, both inhibitors demonstrated therapeutic failure as compared
99 to the reference drug, Bz [3, 11]. Among several possibilities, the disappointing results
100 could be related to the lack of translation from *in vitro* and *in vivo* models as compared
101 to the clinic trials as well as lack of pharmacological ideal conditions for human
102 therapy. In fact, the efficacy of the CYP51 inhibitors may be related to the dose and
103 time of exposure. It has been proposed that in clinical trials the administrated dose
104 was inferior to those reached in pre-clinical analysis required to sustain the
105 trypanocidal effect. Also, longer drug therapy administration could have improved their
106 clinical trial efficacy [12, 13, 14]. Then, this class of compounds should be more

107 studied in pre-clinical studies as the parasite is dependent on endogenous sterols and

108 their products, reinforcing the purpose to study CYP51 inhibitors as potential treatment
109 to CD [15, 16, 17]. Also, it must be considering others CYP51 inhibitors molecules, like
110 imidazolic compounds VNI and VFV, that revealed very promising activity against a
111 variety of *T. cruzi* strains *in vitro* and *in vivo* including a high stringent male mouse
112 experimental model [10, 18 19, 20]. These findings reinforces the need to test the
113 potential activity of novel inhibitors of CYP51 more potent and selective and that could
114 be designed with optimized pharmacological properties, besides presenting reduced
115 production cost, and which would allow treatment to be carried out for extended
116 periods of time [10, 16, 20]. Thus, the aim of this study was to evaluate the anti-
117 parasitic activity of a novel pyrazolo[3,4-e][1,4]thiazepin based CYP51 inhibitor
118 (compound **1**) and its 17 analogues through different *in silico*, *in vitro* and *in vivo*
119 analysis.

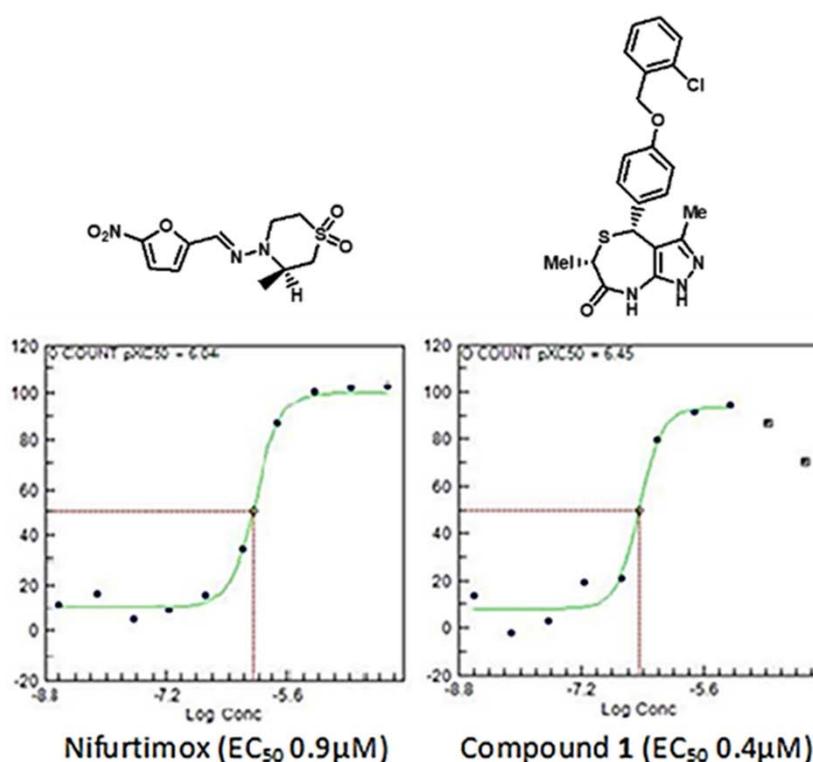
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121 **2. Results and discussion**

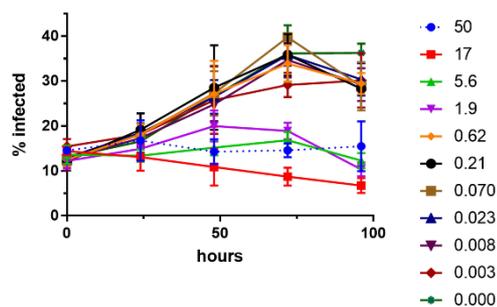
122 Compound dataset was prepared using commercially available compounds, which
123 cover the broad chemical landscape from the Asinex supplier of ZINC database
124 (release 6). A dataset of 240,000 compounds was extracted (smiles format) based on
125 “lead-like” properties such as Molecular weight between 150 and 500, H-bond donor

126 ≤ 5 , H-bond acceptors ≤ 10 , XlogP ≤ 5 , Number of rotatable bonds ≤ 8 . The minimum
127 cut off for the molecular weight was chosen as 150 to avoid the selection of reagent
128 like molecules. 2D-fingerprints in combination with tanimoto distances were used to
129 select a diverse set of 100 compounds. These compounds were screened for their *in*
130 *vitro* efficacy against intracellular *T.cruzi* amastigotes (strain: Silvio X10/7) in a single
131 replicate at 5 μ M. Five compounds that showed >80 % growth inhibition of intracellular

132 amastigotes were selected and a 10-point dose-response curves was carried out to
 133 determine their potency. We next assessed these compounds in the static-cidal and
 134 CYP51_{Tc} assays. Here, we discuss an interesting compound **1** (MW 413.93; cLogP
 135 4.84; H-bond acceptors 4; H-bond donors 2; Number of rotatable bonds 4) with a
 136 pyrazolo[3,4-e][1,4] thiazepin scaffold that showed strong CYP51_{Tc} inhibition, with
 137 activity comparable to Nifurtimox (Figure 1). This compound demonstrated an EC₅₀ of
 138 0.4 μM against intracellular Silvio X10/7 parasites, cidal nature (rate of kill assay) with
 139 minimum cidal concentration (MCC) of 17 μM (Figure 2) and CYP51_{Tc} inhibition with
 140 an IC₅₀ of 0.1 μM (Figure 3). Both potency and cidal nature of this compound are
 141 compared to Nifurtimox in Table 1.



142
 143 **Figure 1.** Molecular structures and ten point dose response curves of Nifurtimox and
 144 compound **1** with calculated potency values. The X-axis shows log of compound molar
 145 concentrations (M) and Y-axis shows the normalized activity based on the
 146 measurement of number of amastigotes per host cell.



147 **Figure 2.** Rate-of-kill assay for compound 1. Ten concentrations of compound were
 148 tested. Growth curves are shown for the compound at different concentrations
 149 indicated on the right hand side of each growth-line (μM). The level of infection was
 150 assessed every 24 hours for 96 hours. Dotted line represents concentration at which
 151 host cell toxicity was observed. All measurements are the average of three replicates.
 152

153 **Table 1.** Potency assay and Static-cidal (SC) assay results for control compound
 154 Nifurtimox and compound 1

Compound	MAX PI * (Static Cidal)	Activity against Silvio X10/7 strain EC ₅₀ (μM)
Nifurtimox	100	no
Compound 1		

155 *Maximum Percent Inhibition in the static cidal assay, only derived from compound concentrations that
 156 are not toxic to the host cells.

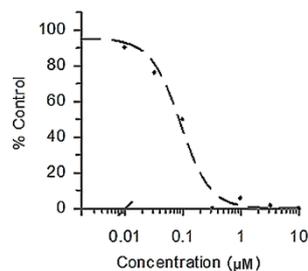
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162 **Inhibitory effect of compound 1 on CYP51_{Tc} activity**

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164 **Figure 3.** *T. cruzi* CYP51 Inhibition Assay results for the compound **1** showing an IC_{50} value of
165 0.1 μ M. 'Y' axis range 95.02 (std. error 4.94), slope factor 1.70 (std. error 0.39). Refer
166 supporting information for test concentrations and associated data.
167

168 Compound **1** was tested in a fluorescence based functional assay [21] using
169 recombinant expressed *T. cruzi* CYP51 (Tulahuen strain). Posaconazole was used as
170 a reference compound. Compound **1** showed potent inhibition of *T. cruzi* CYP51 with
171 IC_{50} value of 0.1 μ M (Figure 3), whereas Posaconazole displayed an IC_{50} value of
172 0.048 μ M (refer supporting information). In a similar assay, Riley et al. [21] reported an
173 IC_{50} value of 0.880 μ M for Fluconazole. Overall compound **1** displayed over 8-fold
174 more inhibitory activity than Fluconazole and 2-fold less inhibitory activity than
175 Posaconazole.

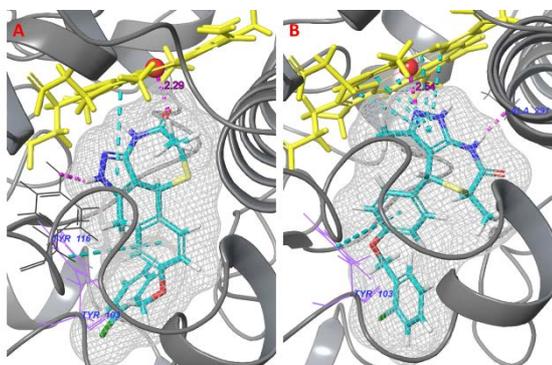
176 As compound **1** showed good potency, cidal nature and strong CYP51_{Tc} inhibition, we
177 have selected this compound for further development. Initially we have assessed the

178 binding interactions of compound **1** with CYP51 using structure based drug design in
179 order to gain deeper understanding of the interactions at the molecular level. Further,

180 we have tested this compound against other strains of *T.cruzi* representatives of
181 distinct parasite discrete type units - DTUs (Tulahuen – DTU VI and Y, DTU II).

182 **Binding Mode Prediction and Molecular interactions analysis of compound 1**

183 The ligand present in the CYP51_{Tc} (PDB ID: 4C27) crystal structure was re-docked to
184 validate the docking protocol, which was able to successfully reproduce the binding
185 mode observed in the crystal structure. The RMSD value between the heavy atoms of
186 the GLIDE-predicted pose and the crystallographic binding pose is 0.41 Å. We used
187 this docking protocol to dock compound **1** to CYP51_{Tc}.



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F CYP51 of *T. cruzi* is represented as a transparent surface with cartoon secondary structure.
i Heme in yellow sticks, Fe in red CPK model. Compound **1** representation: Van der Waals
g surface around the compound **1** in mesh format, nitrogen in blue, oxygen in red, sulfur in yellow,
u chlorine in green. Heme iron co-ordination for both binding poses (A and B) in magenta
r dotted lines. π - π stacking interactions between pyrazole ring and heme macromolecule in
e teal dashed lines. Side chains of the amino acid residues (wire format) involved in interactions
4 with compound **1** (A, B) through either π - π stacking (Teal dashed lines) or hydrogen bonding
: (magenta dashed lines). All structure based drug design Images in this article are generated
P using Schrödinger drug design software.

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201 Docking studies predicted two binding poses for compound **1** (Figure 4). In both the
202 poses, the benzyloxy phenyl moiety attached to the basic scaffold pyrazolo-thiazepin
203 mainly occupies the hydrophobic cleft making good van der Waals and hydrophobic
204 interactions with the surrounding amino acids, but the basic scaffold is in different
205 orientation, and the heme iron coordination is different. In the top scoring **pose 1** (XP
206 GScore = - 9.44), the heme iron coordination is predicted with the carbonyl oxygen of
207 the basic scaffold at a distance of 2.29 Å. The Pyrazole ring of the basic scaffold
208 engages in π - π stacking interactions with the heme macrocycle and hydrogen bonds
209 through N-H group with a hydroxyl oxygen of the B'/C loop residue Tyr116. In addition,
210 the phenyl group undergoes a π - π stacking interaction with the phenyl ring of the B'
211 helix residue Tyr103. In the slightly lower scoring **pose 2** (XP GScore = - 8.66), the
212 heme iron coordination is predicted with the nitrogen atom of the pyrazole ring at a
213 distance of 2.54 Å. Similar to pose 1, the Pyrazole ring undergoes π - π stacking
214 interactions with the heme macrocycle and the phenyl group engages in π - π stacking
215 interaction with the phenyl ring of the B' helix residue Tyr103. In addition, the N-H
216 group of the thiazepin ring hydrogen bonded to the carbonyl oxygen of the I-helix
217 residue Ala291. The proposed hydrogen bond formation with Tyr116, π - π stacking
218 interactions with the heme macrocycle and Tyr103 appear to mimic the
219 posaconazole/fluconazole interactions with the CYP51_{TC}. Overall, both the poses are

220 stabilized in the binding pocket with a network of interactions including Fe-
221 coordination, π - π stacking and hydrogen bonding interactions. It appears that these
222 interactions play a dominant role in the binding affinity of compound **1**.

223 **Evaluation of *in vitro* activity of compound 1 against Tulahuen and Y strains**

224 Compound **1** was further evaluated against intracellular forms of *T. cruzi* (Tulahuen
 225 strain transfected with the β -galactosidase). The anti-parasitic activity of this
 226 compound was measured to determine the EC₅₀ values after 96 h of compound
 227 incubation. The data showed that compound **1** present considerable reduction in the
 228 number of parasite population, with an EC₅₀ of 3.86 μ M, which is comparable to the
 229 activity of reference compound (Bz) that reached an EC₅₀ of ~2.63 μ M (Table 2). Our
 230 data confirmed that this CYP51 inhibitor is able to act upon different parasite strains
 231 (Sylvio-X10/7 and Tulahuen), belonging also to different *T. cruzi* DTUs (I and VI,
 232 respectively), which is a very promising characteristic for a novel drug for CD [8].

233 **Table 2.** Activity (EC₅₀ - Mean \pm SD) against intracellular forms (Tulahuen B-Galactosidase
 234 transfected strain) and bloodstream trypomastigotes of *T. cruzi* (2 and 24h - Y strain).

Compound	Activity against intracellular forms (EC ₅₀ μ M)	Activity against bloodstream trypomastigotes (EC ₅₀ μ M)	
		2 h	24 h
Benznidazole	2.63 \pm 0.49	>10	6.02 \pm 1.47
Compound 1	3.86 \pm 0.26	>10	4.00 \pm 0.35

235
 236 Next, as compound **1** displayed a potential activity *in vitro* against intracellular forms of
 237 *T. cruzi* (Tulahuen and Sylvio X10/7 strains), this compound was assayed against
 238 other relevant forms for mammalian infection, the bloodstream trypomastigotes, using
 239 also another parasite strain and DTU (Y strain, DTU II), under a time drug exposure
 240 incubation (2 and 24 h). The findings demonstrated that although both compound **1**

241 and Bz were not active after 2 h of incubation (EC_{50} of $>10 \mu\text{M}$), the CYP51 inhibitor
242 showed trypanocidal effect after longer periods of incubation of BT, with an EC_{50} of ~ 4
243 μM , and as found for intracellular parasites (Tulahuen strain), with comparable
244 potency as Bz (Table 2) and Nifurtimox (Table 1). We have therefore evaluated a set
245 of selected analogues (**1a-1q**, Table 3) with various functional groups that differ in

246 electronic properties, position and steric properties as follows. The phenotypic effect of
247 these compounds was first screened regarding their ability to reduce the infection
248 levels against intracellular forms of *T. cruzi* (Tulahuen strain). Compounds with > 50 %
249 reduction on infection levels using a single concentration (10-12 μ M, corresponding
250 the EC₉₀ value of Bz, Table 3) were further subjected to potency assays (activity
251 expressed as EC₅₀ in Table 3). Toxicity profiles were determined against L929 cell
252 cultures by incubating for 96 h with different concentrations of these compounds and
253 then cell viability evaluated by both light microscopy and colorimetric assay
254 (AlamarBlue tests) and expressing the respective LC₅₀ values (Table 3). The ratio of
255 LC₅₀ and EC₅₀ values is presented in the same table as Selectivity Index (SI),
256 indicating the quantity of compound that is active against the *T.cruzi* but is not toxic

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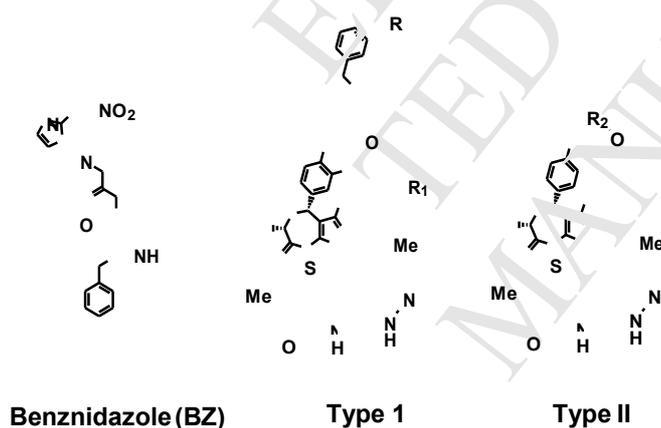
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269 **Table 3.** Activity (percentage reduction of the L929 culture infection) using a fixed
 270 concentration (10-12 μM), and anti-parasitic potency (EC_{50} - Mean \pm SD – 96 h), mammalian
 271 host cell toxicity (L929 – 96 h) and selectivity against intracellular forms of *T.cruzi* (Tulahuen
 272 *B*-Galactosidase transfected strain).

273



ID	Type	R			R ₁	R ₂	% redt 1 infecti (si conce
		Ortho	Meta	Para			
Bz	-	-	-	-	-	-	89.3
1	I	Cl	H	H	H	-	14.88
1a	I	H	H	Cl	H	-	63.64
1b	I	F	H	H	H	-	70.68
1c	I	H	F	H	H	-	32.42
1d	I	H	H	F	H	-	69.14
1e	I	Me	H	H	H	-	89.2
1f	I	H	Me	H	H	-	

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277 The *in vitro* analysis revealed that five compounds (**1b**, **1c**, **1e**, **1f** and **1k**, Table 3)
278 were able to decrease the intracellular parasite load in the infected L929 cell cultures,
279 ranging 53 up to 89 % of reduction when concentration equivalent of EC₉₀ of Bz was
280 used (Table 3). These compounds also displayed low toxicity profile against
281 mammalian host cells (LC₅₀ ≥ 50 μM, table 3). Compounds **1c**, **1e** and **1f** did not
282 induce loss of cellular viability after incubation for 96 h with doses up to 98 μM.
283 Compound **1b** maintained mammalian cell viability up to 200 μM. However, compound
284 **1k** exhibited moderate toxicity with LC₅₀ of ~50 μM. Compound **1e** displayed high SI
285 value (> 50), comparable to benznidazole. Compounds **1b**, **1c**, **1f** displayed
286 considerable selectivity with SI values >38, >19 and >36 respectively. Although
287 compound **1k** showed moderate potency against intracellular parasites, it showed
288 poor selectivity (SI value = 4.7). An interesting observation from the above results
289 (Table 3) is that compounds with *ortho*-substitutions displayed better selectivity and
290 toxicity profiles when compared with *meta*-substituted compounds (compare SI and
291 LC₅₀ values of **1b** with **1c** and **1e** with **1f**) when assayed against intracellular forms of
292 *T. cruzi* (Tulahuen strain). Compounds (**1b**, **1c**, **1e**, **1f** and **1k**, Table 3) showed good
293 *in vitro* potency against intracellular forms of *T. cruzi* (Tulahuen β-Galactosidase
294 transfected strain) were further tested for their activity against bloodstream

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295 trypomastigotes (Y strain). The results are shown in Table 4. After 2 h of incubation, all
296 compounds showed an EC₅₀ of >50 μM, similarly as benznidazole. After 24 h of
297 incubation, compounds **1b**, **1e** and **1k** did not show much improvement in the potency.
298 Compound **1c** showed large variations in potency possibly due to compound
299 instability. Interestingly, compound **1f** showed (EC₅₀ of ~20 μM) comparable activity to
300 reference compound (EC₅₀ of ~16 μM). The predicted binding pose of compound **1f** is

301 shown in Figure 5 (A). The orientation of the binding pose and the receptor
 302 interactions are similar to compound **1**. We have further evaluated all five compounds
 303 for their toxicity on primary cell cultures. Uninfected cardiac cells were incubated for 24
 304 h with different doses of these compounds and then cell viability evaluated by both
 305 light microscopy and colorimetric assay Prestoblue.

306 **Table 4.** Activity (EC_{50} μ M - Mean \pm SD – 2 and 24h) against bloodstream trypomastigotes of
 307 *T. cruzi* (Y strain) besides toxicity upon cardiac cells (24h) and selectivity profile.

Compound	EC_{50} μ M		LC μ M 50	SI (LC /EC – 24 h) μ M	
	2 h	24 h		50	50
Benznidazole	>50	16.86 \pm 3.34	>200		>12
1b	>50	>50	>200		>4
1c	>50	46.98 \pm 10.93	93.97 \pm 11.73		3.71
1e	>50	38.6 \pm 6	>200		>5.2
1f	>50	20.62 \pm 6.12	66 \pm 25.82		3.20
1k	>50	39.37 \pm 6.37	ND*		ND*

308 * ND: Not determined

309 Compounds **1b**, **1e** did not induce loss of cellular viability after incubation for 24 h up
 310 to 200 μ M (Table 4). Compounds **1c** and **1f** displayed low toxicity with LC_{50} values of
 311 ~93 and ~66 μ M, respectively.

312 Structure Activity Relationship (SAR) Analysis

313 Next, the structure activity relationships of these compounds (**1a-1q**) is discussed
 314 according to their effect towards the infection level and the EC_{50} values against the
 315 intracellular parasites. Compounds **1a-1g** (Table 3) that differ in the substitution

316 pattern on the benzyl ring showed varied levels of potency. Compound **1** with *ortho*-
317 chloro substitution displayed potency with an EC₅₀ of ~3.86 μM. The switching of
318 electron withdrawing, bulky chlorine from *ortho*- position (**1**) to *para*- position (**1a**)
319 resulted in loss of potency. Replacement of *ortho*- chloro substituent with fluoro group
320 (**1b**) displayed potency with an EC₅₀ of ~5.30 μM, and the switching of fluoro group

321 from *ortho*- (**1b**) to *meta*- position (**1c**) displayed similar potency. However, *para*-
322 fluoro substituted compound (**1d**) showed poor potency. Replacement of bulky,
323 electron withdrawing *ortho*- chloro group (**1**) with isosteric, isolipophilic and electron
324 donating methyl group (**1e**) displayed improved potency with an EC₅₀ of ~2.20 μM, and
325 the switching of methyl group from *ortho*- (**1e**) to *meta*- position (**1f**) retained the
326 potency with an EC₅₀ of ~2.70 μM. However, *para*- methyl substitution (**1g**) resulted in
327 complete loss of potency, suggesting that the substitution pattern on the aromatic ring
328 is important. The general trend of activity being *ortho*- and *meta*- substituted
329 compounds are more potent than *para*- substituted compounds (compare **1** with **1a**;
330 compare **1b**, **1c** with **1d**; compare **1e**, **1f** with **1g**). Overall, *para*- substitutions
331 (chloro/fluoro/methyl; **1a**, **1d**, **1g**) on the benzyl ring generally displayed poor or no
332 potency. Docking studies on these compounds suggest that *para*- substitutions
333 protrude from the hydrophobic cleft towards a more solvent exposed area, while the
334 *ortho*- and *meta*- substitutions allow better interactions with the hydrophobic cleft. This
335 may explain why all substitutions at the *para*- position (**1a**, **1d**, **1g**) are not tolerated.
336 The results suggest that the electronic properties of the substituents (chlorine moiety
337 being electron withdrawing and methyl moiety being electron donating) did not
338 influence the activity; however, the position specific and hydrophobic effects of the
339 substituents on the benzyl ring are mainly involved in the variation in potency. We next

340 turned our attention to compounds **1h-1n** with *ortho*- methoxy substituted phenyl ring.

341 Compound **1h** with unsubstituted benzyloxy group showed poor potency. Compounds

342 with *ortho*- (**1i**, **1l**), *meta*- (**1j**, **1m**) and *para*- substitutions (**1n**) on the benzyl ring also

343 displayed poor or no potency, the only exception is *para*-chloro substitution which

344 resulted in a comparatively potent compound **1k**. Our binding mode analysis of

345 compounds **1i**, **1j**, **1l** and **1m** suggested that the introduction of a bulky methoxy group

346 at the *ortho*- position restricts the conformation of the phenyl ring, thereby resulting in
347 a different orientation, losing key interactions such as coordination with the heme iron,
348 π - π stacking with Tyr103 and hydrogen bonding with Tyr116. This appears to have
349 resulted in loss of potency for these compounds. Interestingly, in the top scored
350 binding poses of compounds **1h**, **1k** and **1n** reverse binding modes were observed,
351 where the pyrazolo [3,4-*e*] [1,4] thiazepin-7-one scaffold is placed in the hydrophobic
352 binding cleft, benzyloxy moiety is placed in the B'/C loop and I helix region. This
353 reverse binding mode do not appear to have favoured the potency of compounds **1h**
354 and **1n**. However, for compound **1k**, methoxy phenyl ring is in a different orientation
355 compared to **1h** and **1n**, making strong hydrophobic interactions with the surrounding
356 amino acids, thereby displaying moderate potency with an EC₅₀ of ~10.85 μ M. The

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371 hydrogen bonding (magenta dashed lines) interaction with the NH moiety (pyrazole ring) of
372 compound **1k**.

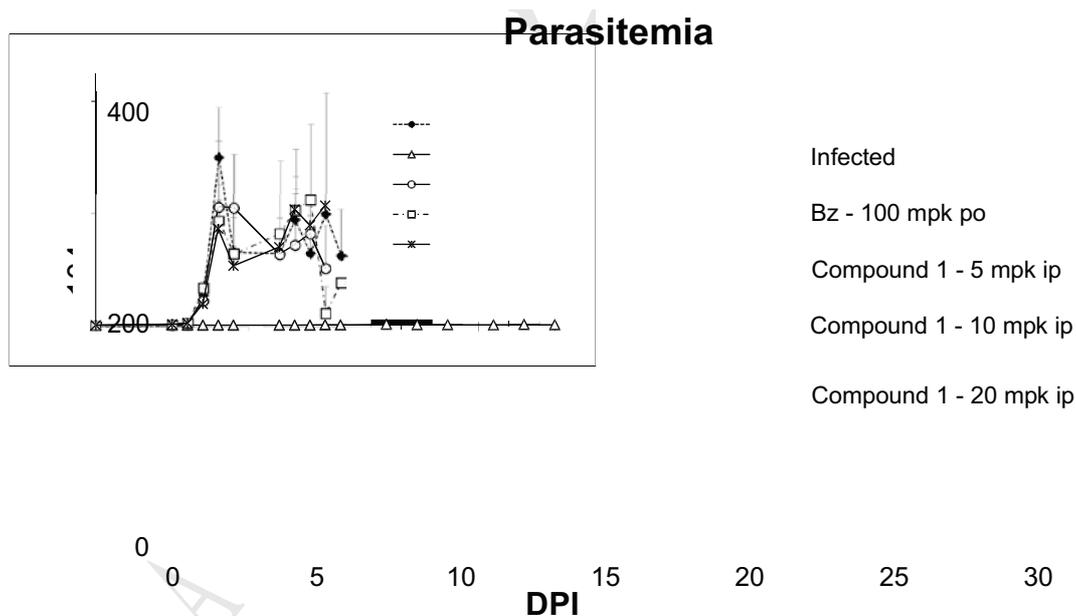
373 We finally probed the contribution of the substituted benzyl group of the pyrazolo [3,4-
374 e] [1,4] thiazepin-7-one ring system towards the potency by screening a small set of
375 analogues with aliphatic substitutions (Type II). Replacement of benzyl ring with
376 isopropyl (**1o**), n-propyl (**1p**) and allyl (**1q**) moieties completely abolished the potency
377 altogether, suggesting that substituted benzyl ring plays an important role in stabilising
378 the complex by forming strong hydrophobic interactions with the hydrophobic cleft.
379 Moreover, perhaps these aliphatic substitutions do not fill the hydrophobic pocket
380 enough, which was needed for potency and stability. Overall, these results suggest
381 that the coordination with the heme iron, π - π stacking and hydrogen bond interactions
382 in the B' helix and B'/C loop region, and the hydrophobic interactions with the residues
383 surrounding the hydrophobic tunnel are crucial for the potency of this series of
384 compounds.

385 Overall, **1e** and **1f** were the more promising compounds regarding potency and
386 selectivity profiles in both the intracellular (Tulahuen) and bloodstream trypomastigote
387 (Y strain) forms of *T. cruzi* when compared to **1b**, **1c**, and **1k**. Hence, as **1e** showed
388 poor solubility profile, we have further evaluated compounds **1** and **1f** for their *in vivo*
389 activity in mouse models of acute *T. cruzi* infection.

390 **Evaluation of compounds *in vivo***

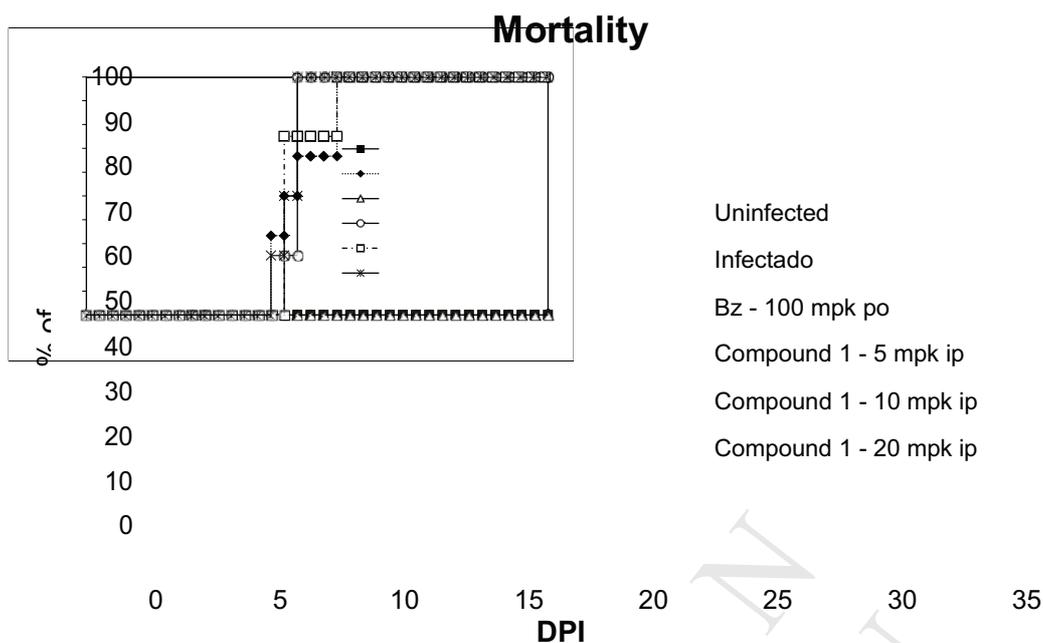
391 The *in vivo* efficacy of the compound **1** was evaluated using non-toxic doses
392 (previously determined in acute mice toxicity tests, showing an NOAEL > 50 mg/kg -
393 i.p., data not shown) using 5, 10 and 20 mg/kg/daily doses. Bz was tested in parallel at
394 its optimal dose of 100 mg/kg [22]. The administration of compound **1** for five days

395 resulted in a mild reduction in the parasitemia peak levels, being much lower than Bz
 396 which reached 100 % of peak suppression. The CYP51 inhibitor reached 30, 38 and
 397 43% of reduction in respective doses of 5, 10 and 20 mg/kg/day at the peak of
 398 parasitemia, (**Figure 6**). Regarding the mortality, while Bz guaranteed 100 % animal
 399 survival, all treated animals with compound **1**, died (**Figure 7**). Unfortunately, using the
 400 same mouse experimental model of *T.cruzi* infection, the administration for five
 401 consecutive days of 5-20 mg/kg of compound **1f** (non-toxic doses based on previous
 402 acute toxicity assays using female Swiss mice) did not induce parasitemia decrease
 403 neither protect against animal mortality. Bz was able to completely suppress
 404 parasitemia peak (8 dpi) and gave 100 % mice survival (data not shown).



405

406 **Figure 6.** *In vivo* effect of Compound **1** (5-20 mg/kg) and Bz (100 mg/kg) upon *T.cruzi*
 407 parasitemia levels using an mouse model of acute infection (Y strain).mpk = mg/kg; DPI: Days
 408 post-infection.



409

410 **Figure 7.** *In vivo* effect of Compound 1 (5-20 mg/kg) and Bz (100 mg/kg) upon animal
 411 mortality induced by *T. cruzi* infection using an mouse model of acute infection (Y strain). mpk
 412 = mg/kg; DPI: Days post-infection.

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416 3. Conclusion

417 Only two drugs are available for the treatment of Chagas disease, and both have

418 limited efficacy in addition to several adverse effects. So the search for more effective

419 therapeutic alternatives is urgently needed, to face these limitations CYP51 inhibitors

420 were suggested due to its specific and selective mechanism of action. In this study we

421 have focused on the initial lead optimization of pyrazolo[3,4-e][1,4]thiazepin based

422 lead compound 1. This compound displayed comparable activity with nifurtimox and

423 benznidazole, being potent against multiple strains (Sylvio-X10/7, Tulahuen and Y

424 strain) and forms of *T. cruzi*. Fluorescence based functional assay confirmed that the

425 mode of action of lead compound **1** is via inhibition of CYP51_{Tc}. Structure based drug
426 design studies predicted the binding interactions of lead compound **1** with CYP51_{Tc} to
427 be similar to Posaconazole and Fluconazole. A small library of analogues of

428 compound **1** were evaluated for their potency and selectivity profiles against both the
429 intracellular (Tulahuen) and bloodstream trypomastigote (Y strain) forms of *T. cruzi*.
430 This resulted in interesting SARs that are in good agreement with our predicted
431 binding modes. Compound **1f** showed most promising anti-*T.cruzi* activity. Lead
432 compound **1** and compound **1f** were further evaluated for their *in vivo* activity in mouse
433 models of acute *T.cruzi* infection. Compound **1** is only partially effective in the
434 reduction of parasitemia possibly due to limited permeability and solubility
435 characteristics. Compound **1f** failed to reduce the parasitemia possibly due to poor
436 permeability. Solubility and permeability are important physicochemical characteristics
437 of drug like compounds that reflect their bioavailability. The CYP51_{Tc} inhibitors
438 described here provide an excellent template for further lead optimization of
439 pyrazolo[3,4-e][1,4]thiazepin based analogues with improved solubility and
440 permeability properties that may facilitate the anti-*T.cruzi* drug development.

441 **Acknowledges:** The authors are in debt to Dr. Jennifer Riley, Dr. John Thomas and
442 Dr. Manu De Rycker (University of Dundee) for help with generating data associated
443 with CYP51 inhibition assay, potency and rate of kill assays (Silvio X10/7 strain). We
444 would also like to thank Dr. Rhodri Owen (University of East London) for providing
445 HPLC support and for all technical support offering by Marcos Meuser

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450 Cruz, PDTIS, PAEF/CNPq/Fiocruz, CAPES and University of East London. MNCS is
451 research fellow of CNPq and CNE research.

452 **4. Experiment**

453 **Compound database preparation:** All the compounds in the dataset were processed
454 through concord for the three-dimensional structure generation (mol2 files).
455 Compounds with inappropriate isomeric specifications, compounds containing
456 disallowed atoms in terms of valency and compounds with errors in ring definition are
457 excluded from the co-ordinate generation. These compounds were converted into SLN
458 format (SYBYL Line Notation) using UNITY translate module in SYBYL Version 7.3
459 (Tripos) [23]. The selector module was used to generate two dimensional fingerprints
460 and tanimoto distances. All the compounds in this study were purchased from
461 ASINEX.

462 **Molecular docking:** Molecular modelling studies were performed primarily using
463 Schrödinger drug design software suite (Schrödinger Release 2017-2) [24].

464 **Protein preparation:** The CYP51_{Tc} (PDB: 4C27 with a resolution of 1.95 Å) protein
465 structure was retrieved from Protein Data Bank. The protein was initially prepared
466 using the Protein Preparation Wizard module of Schrödinger suite that prepares the
467 protein by adding hydrogens, assigning correct bond orders, creating zero bond
468 orders to metals, fixing errors like missing side chains and adjusting the ionisation and
469 tautomeric states (via Epik). We have deleted all the water molecules, chloride ions

470 and solvents (ethylene glycol) from the protein. Optimisation of the hydrogen bonding
471 network and the orientation of the hydroxyl/thiol groups, terminal amide groups in Asn
472 and Gln, and His states was carried out using the ProtAssign algorithm. Finally,
473 restrained minimization was carried out using all atoms OPLS3 force fields, with
474 converge heavy atoms to RMSD set to 0.3 Å (default).

475 **Ligand Preparation:** All the compounds were prepared using Ligprep module, which
476 performed addition of hydrogens, 2D to 3D conversion, generation of ionization and
477 tautomeric states including metal binding states (via Epik) at physiological pH 7.0 ±
478 2.0, and also generated possible stereoisomers and ring conformations using default
479 settings. All the compounds were energy minimised using OPLS3 force fields.

480 **Docking studies:** The Receptor Grid generation was carried out by identifying the
481 ligand in the 4C27 crystal structure and excluding it from the grid generation. The grid
482 covers the entire binding site, which includes the heme iron macromolecule. Metal
483 coordination constraint was applied at the Fe atom of the heme that defines an
484 interaction between the ligand atom and the heme iron. Docking of compounds was
485 carried out using Schrödinger GLIDE, in extra precision (XP) mode.

486 **CYP51 inhibitory assay:** The CYP51 assay was carried out as described in [21]. All
487 the potency assays and rate of kill assays with *T. cruzi* Silvio X10/7 strain were carried
488 out as described in [25].

489 **Spectroscopic data analysis and compound purity assessment by HPLC**

490 All 18 compounds were purchased directly from Asinex and used as supplied unless
491 otherwise stated. Accurate mass and nominal mass measurements were performed
492 using Bruker micrOTOF mass spectrometer. All ¹HNMR spectra were recorded in

493 deuterio-DMSO in 5 mm tubes, with tetramethylsilane as an internal standard, using a
494 Bruker instrument. Chemical shifts (δ) are reported in ppm, and coupling constants (J)
495 are given in Hz. Signals are represented by s (singlet), d (doublet), t (triplet), q
496 (quartet), dq (double quartet) m (multiplet), bs (broad singlet), and dd (double doublet).
497 HPLC was carried out using Agilent-1200 instrument. Column: Agilent eclipse plus
498 C18 (150mm \times 4.6mm, 5 μ m particle size), mobile phase 0.01% TFA in

499 acetonitrile/water (5% to 95% organic over 20min at 1mL/min), 10 μ L injection.
500 Detection was at 254nm, runtime 25 min. All the samples were prepared by dissolving
501 the 5mg of compound in 0.7 mL of d^6 -DMSO (Sample solution prepared for HNMR
502 was analyzed by HPLC). Blank (without the compound) d^6 -DMSO was used as a
503 reference standard. Most of the compounds displayed >90% purity except compounds
504 **1h** (77%), **1j** (88%), **1l** (87%) and **1n** (84%). Refer to the supporting information for
505 complete data on HPLC purity analysis. Isomer ratios of the compounds were
506 calculated from $^1\text{H-NMR}$ spectra based on the CH_3 signal (s) from substituted pyrazole
507 ring of the one of the isomer against CH_3 signal (s) from substituted pyrazole ring of
508 the other isomer.

509 *4-[4-(2-Chloro-benzyloxy)-phenyl]-3,6-dimethyl-4,8-dihydro-1H-pyrazolo[3,4-*
510 *e][1,4]thiazepin-7-one (Compound 1)*. Mixture of two isomers (98:2), δ_{H} /ppm (400
511 MHz, d^6 -DMSO): Major isomer peaks are at 12.32 (1H, bs, NH from 3-pyrazole ring),
512 9.66 (1H, s, NH from sec.amide), 7.62-7.55 (1H, m, Ar-H), 7.54-7.46 (1H, m, Ar-H),
513 7.43-7.34 (2H, m, Ar-H), 7.21 (2H, d, $J = 8.7$ Hz, Ar-H), 6.94 (2H, d, $J = 8.7$ Hz, Ar-H),
514 5.56 (1H, s, methine from [1,4] thiazepin ring), 5.11 (2H, s, CH_2), 3.68 (1H, q, $J = 7.0$,
515 methine from [1, 4] thiazepin ring), 1.91 (3H, s, CH_3 from pyrazole ring), 1.19 (3H, d, J
516 = 7.0 Hz, CH_3 from [1, 4] thiazepin ring); Observable minor isomer peaks are at 9.86
517 (1H, s, NH from sec.amide), 7.0 (2H, d, $J = 8.8$ Hz, Ar-H), 5.33 (1H, s, methine from
518 [1, 4] thiazepin ring), 5.14 (2H, s, CH_2), 1.69 (3H, s, CH_3 from pyrazole ring), 1.08 (3H,

519 d, $J = 7.1$ Hz, CH_3 from [1, 4] thiazepin ring). HRMS- m/z (ESI): found 414.1046
520 ($\text{C}_{21}\text{H}_{22}\text{N}_3\text{O}_2\text{S} [\text{M} + \text{H}]^+$) requires 414.0965.

521 4-[4-(4-Chloro-benzyloxy)-phenyl]-3,6-dimethyl-4,8-dihydro-1H-pyrazolo[3,4-
522 e][1,4]thiazepin-7-one (**1a**). Mixture of two isomers (96:4), δ_{H} /ppm (400 MHz, d^6 -

523 DMSO): Major isomer peaks are at 12.31 (1H, s, NH from 3-pyrazole ring), 9.65 (1H,
524 s, NH from sec.amide), 7.48-7.41 (4H, m, Ar-H), 7.19 (2H, d, J = 8.7 Hz, Ar-H), 6.91
525 (2H, d, J = 8.7 Hz, Ar-H), 5.55 (1H, s, methine from [1, 4] thiazepin ring), 5.05 (2H, s,
526 CH₂), 3.67 (1H, q, J = 7.04 Hz, methine from [1, 4] thiazepin ring), 1.90 (3H, s, CH₃
527 from pyrazole ring), 1.18 (3H, d, J = 7.08 Hz, CH₃ from [1,4] thiazepin ring);
528 Observable minor isomer peaks are at 9.86 (1H, s, NH from sec.amide), 6.97 (2H, d, J
529 = 8.7 Hz, Ar-H), 5.32 (1H, s, methine from [1, 4] thiazepin ring), 5.08 (2H, s, CH₂), 1.68
530 (3H, s, CH₃ from pyrazole ring), 1.07 (3H, d, J = 7.2 Hz, CH₃ from [1, 4] thiazepin ring).
531 HRMS-m/z (ESI): found 414.1042 (C₂₁H₂₂N₃O₂S [M + H]⁺), requires 414.0965.

532 *4-[4-(2-Fluoro-benzyloxy)-phenyl]-3,6-dimethyl-4,8-dihydro-1H-pyrazolo[3,4-*
533 *e][1,4]thiazepin-7-one (1b)*. Mixture of two isomers (95:5), δ_{H} /ppm (400 MHz, d⁶-
534 DMSO): Major isomer peaks are at 12.31 (1H, bs, NH from 3-pyrazole ring), 9.66 (1H,
535 s, NH from sec.amide), 7.58-7.50 (1H, m, Ar-H), 7.46-7.37 (1H, m, Ar-H), 7.29-7.15
536 (2H, m, Ar-H) this multiplet is overlapped with doublet from the aromatic ring that is
537 directly attached to the pyrazolo[3, 4-e][1, 4]thiazepin basic scaffold, 7.20 (2H, d, J =
538 8.7 Hz, Ar-H), 6.94 (2H, d, J = 8.7 Hz, Ar-H), 5.56 (1H, s, methine from [1, 4] thiazepin
539 ring), 5.09 (2H, s, CH₂), 3.68 (1H, q, J = 7.0 Hz, methine from [1, 4] thiazepin ring),
540 1.91 (3H, s, CH₃ from pyrazole ring), 1.19 (3H, d, J = 7.0 Hz, CH₃ from [1,4] thiazepin
541 ring); Observable minor isomer peaks are at 9.85 (1H, s, NH from sec.amide), 6.94

542 (2H, d, J = 8.7 Hz, Ar-H), 5.56 (1H, s, methine from [1, 4] thiazepin ring), 5.11 (2H, s,
543 CH₂), 1.68 (3H, s, CH₃ from pyrazole ring), 1.07 (3H, d, J = 7.2 Hz, CH₃ from [1, 4]
544 thiazepin ring). ¹⁹FNMR: 118.41 ppm. HRMS-m/z (ESI): found 398.1341
545 (C₂₁H₂₁FN₃O₂S [M + H]⁺), requires 398.1260.

546 4-[4-(3-Fluoro-benzyloxy)-phenyl]-3,6-dimethyl-4,8-dihydro-1H-pyrazolo[3,4-
547 e][1,4]thiazepin-7-one (**1c**). Mixture of two isomers (86:14), δ H/ppm (400 MHz, d^6 -
548 DMSO): Major isomer peaks are at 12.30 (1H, bs, NH from 3-pyrazole ring), 9.65 (1H,
549 s, NH from sec.amide), 7.47-7.38 (1H, m, Ar-H), 7.31-7.23 (2H, m, Ar-H), 7.16-7.12
550 (1H, m, Ar-H) this multiplet is overlapped with doublet from the aromatic ring that is
551 directly attached to the pyrazolo[3, 4-e][1, 4]thiazepin basic scaffold, 7.19 (2H, d, J =
552 8.7 Hz, Ar-H), 6.92 (2H, d, J = 8.7 Hz, Ar-H), 5.55 (1H, s, methine from [1, 4] thiazepin
553 ring), 5.08 (2H, s, CH₂), 3.67 (1H, q, J = 7.03 Hz, methine from [1, 4] thiazepin ring),
554 1.90 (3H, s, CH₃ from pyrazole ring), 1.19 (3H, d, J = 7.08 Hz, CH₃ from [1,4] thiazepin
555 ring); Observable minor isomer peaks are at 9.85 (1H, s, NH from sec.amide), 6.97
556 (2H, d, J = 8.7 Hz, Ar-H), 5.32 (1H, s, methine from [1, 4] thiazepin ring), 5.10 (2H, s,
557 CH₂), 3.24 (1H, q, J = 1.67 Hz), 1.67 (3H, s, CH₃ from pyrazole ring), 1.06 (3H, d, J =
558 7.2 Hz, CH₃ from [1, 4] thiazepin ring). ¹⁹FNMR: 113.20 ppm. HRMS-m/z (ESI): found
559 398.1341 (C₂₁H₂₁FN₃O₂S [M + H]⁺), requires 398.1260.

560 4-[4-(4-Fluoro-benzyloxy)-phenyl]-3,6-dimethyl-4,8-dihydro-1H-pyrazolo[3,4-
561 e][1,4]thiazepin-7-one (**1d**). Mixture of two isomers (96:4), δ H/ppm (400 MHz, d^6 -
562 DMSO): Major isomer peaks are at 12.33 (1H, bs, NH from 3-pyrazole ring), 9.65 (1H,
563 s, NH from sec.amide), 7.54-7.44 (2H, m, Ar-H), 7.25-7.16 (4H, m, Ar-H), 6.91 (2H, d,
564 J = 8.7 Hz, Ar-H), 5.55 (1H, s, methine from [1, 4] thiazepin ring), 5.03 (2H, s, CH₂),
565 3.67 (1H, q, J = 7.02 Hz, methine from [1, 4] thiazepin ring), 1.90 (3H, s, CH₃ from

566 pyrazole ring), 1.19 (3H, d, J = 7.08 Hz, CH₃ from [1,4] thiazepin ring); Observable
567 minor isomer peaks are at 9.85 (1H, s, NH from sec.amide), 6.96 (2H, d, J = 8.7 Hz,
568 Ar-H), 5.32 (1H, s, methine from [1, 4] thiazepin ring), 5.05 (2H, s, CH₂), 1.67 (3H, s,
569 CH₃ from pyrazole ring), 1.07 (3H, d, J = 7.1 Hz, CH₃ from [1, 4] thiazepin ring).

570 ^{19}F NMR: 114.49 ppm. HRMS- m/z (ESI): found 398.1340 ($\text{C}_{21}\text{H}_{21}\text{FN}_3\text{O}_2\text{S}$ [$\text{M} + \text{H}$] $^+$),
571 requires 398.1260.

572 *3,6-Dimethyl-4-[4-(2-methyl-benzyloxy)-phenyl]-4,8-dihydro-1H-pyrazolo[3,4-*
573 *e][1,4]thiazepin-7-one (1e)*. Mixture of two isomers (97:3), $\delta\text{H/ppm}$ (400 MHz, d^6 -
574 DMSO): Major isomer peaks are at 12.30 (1H, bs, NH from 3-pyrazole ring), 9.66 (1H,
575 s, NH from sec.amide), 7.38 (1H, d, $J = 7.30$ Hz, Ar-H), 7.28-7.14 (5H, m, Ar-H), 6.94
576 (2H, d, $J = 8.7$ Hz, Ar-H), 5.56 (1H, s, methine from [1, 4] thiazepin ring), 5.03 (2H, s,
577 CH_2), 3.68 (1H, q, $J = 7.0$ Hz, methine from [1, 4] thiazepin ring), 2.30 (3H, s, *ortho*
578 CH_3), 1.91 (3H, s, CH_3 from pyrazole ring), 1.19 (3H, d, $J = 7.0$ Hz, CH_3 from [1,4]
579 thiazepin ring); Observable minor isomer peaks are at 9.85 (1H, s, NH from
580 sec.amide), 6.99 (2H, d, $J = 8.7$ Hz, Ar-H), 5.32 (1H, s, methine from [1, 4] thiazepin
581 ring), 5.06 (2H, s, CH_2), 2.32 (3H, s, *ortho* CH_3), 1.68 (3H, s, CH_3 from pyrazole ring),
582 1.08 (3H, d, $J = 7.2$ Hz, CH_3 from [1, 4] thiazepin ring). HRMS- m/z (ESI): found
583 394.1586 ($\text{C}_{22}\text{H}_{24}\text{N}_3\text{O}_2\text{S}$ [$\text{M} + \text{H}$] $^+$), requires 394.1511.

584 *3,6-Dimethyl-4-[4-(3-methyl-benzyloxy)-phenyl]-4,8-dihydro-1H-pyrazolo[3,4-*
585 *e][1,4]thiazepin-7-one (1f)*. Mixture of two isomers (66:34), $\delta\text{H/ppm}$ (400 MHz, d^6 -
586 DMSO): Major isomer peaks are at 12.31 (1H, bs, NH from 3-pyrazole ring), 9.65 (1H,
587 s, NH from sec.amide), 7.32-7.09 (6H, m, Ar-H), 6.91 (2H, d, $J = 8.8$ Hz, Ar-H), 5.55
588 (1H, s, methine from [1, 4] thiazepin ring), 5.01 (2H, s, CH_2), 3.67 (1H, q, $J = 7.0$ Hz,

589 methine from [1, 4] thiazepin ring), 2.31 (3H, s, *meta* CH₃), 1.90 (3H, s, CH₃ from
590 pyrazole ring), 1.19 (3H, d, J = 7.0 Hz, CH₃ from [1, 4] thiazepin ring); Observable
591 minor isomer peaks are at 9.85 (1H, s, NH from sec.amide), 7.32-7.09 (6H, m, Ar-H),
592 6.96 (2H, d, J = 8.8 Hz, Ar-H), 5.32 (1H, s, methine from [1,4] thiazepin ring), 5.03(2H,
593 s, CH₂), 3.24 (1H, q, J = 7.2 Hz methine from [1, 4] thiazepin ring), 2.31 (3H, s, *meta*

594 CH₃), 1.70 (3H, s, CH₃ from pyrazole ring), 1.01 (3H, d, J = 7.2 Hz, CH₃ from [1,4]
595 thiazepin ring). HRMS; m/z (ES): found 394.1580 (C₂₂H₂₄N₃O₂S [M + H]⁺), requires
596 394.1511.

597 *3,6-Dimethyl-4-[4-(4-methyl-benzyloxy)-phenyl]-4,8-dihydro-1H-pyrazolo[3,4-*
598 *e][1,4]thiazepin-7-one (1g)*. Mixture of two isomers (69:31), δ H/ppm (400 MHz, d⁶-
599 DMSO): Major isomer peaks are at 12.30 (1H, bs, NH from 3-pyrazole ring), 9.65 (1H,
600 s, NH from sec.amide), 7.36-7.27 (2H, m, Ar-H), 7.22-7.12 (4H, m, Ar-H), 6.90 (2H, d,
601 J = 8.7 Hz, Ar-H), 5.55 (1H, s, methine from [1, 4] thiazepin ring), 5.00 (2H, s, CH₂),
602 3.67 (1H, q, J = 7.0 Hz, methine from [1, 4] thiazepin ring), 2.29 (3H, s, *para* CH₃),
603 1.90 (3H, s, CH₃ from pyrazole ring), 1.18 (3H, d, J = 7.08 Hz, CH₃ from [1,4] thiazepin
604 ring); Observable minor isomer peaks are at 9.85 (1H, s, NH from sec.amide), 7.36-
605 7.27 (2H, m, Ar-H), 7.22-7.12 (4H, m, Ar-H), 6.94 (2H, d, J = 8.8 Hz, Ar-H), 5.31 (1H,
606 s, methine from [1, 4] thiazepin ring), 5.01 (2H, s, CH₂), 3.24 (1H, q, J = 7.2 Hz
607 methine from [1, 4] thiazepin ring), 2.30 (3H, s, *para* CH₃), 1.67 (3H, s, CH₃ from
608 pyrazole ring), 1.07 (3H, d, J = 7.2 Hz, CH₃ from [1,4] thiazepin ring). HRMS; m/z (ES):
609 found 394.1582 (C₂₂H₂₄N₃O₂S [M + H]⁺) requires 394.1511.

610 *4-(4-Benzyloxy-3-methoxy-phenyl)-3,6-dimethyl-4,8-dihydro-1H-pyrazolo[3,4-*
611 *e][1,4]thiazepin-7-one (1h)*. Mixture of two isomers (63:37), δ H/ppm (400 MHz, d⁶-
612 DMSO): First isomer peaks are at 12.30 (1H, bs, NH from 3-pyrazole ring), 9.66 (1H,
613 s, NH from sec.amide), 7.50-7.27 (5H, m, Ar-H), 7.01-6.89 (2H, m, Ar-H), 6.75 (1H, dd,

614 J = 8.3 + 2.0 Hz, Ar-H), 5.51 (1H, s, methine from [1, 4] thiazepin ring), 5.02 (2H, s,
615 CH₂), 3.71 (3H, s, OCH₃), 3.67 (1H, q, J = 7.0 Hz, methine from [1, 4] thiazepin ring),
616 1.95 (3H, s, CH₃ from pyrazole ring), 1.19 (3H, d, J = 7.04 Hz, CH₃ from [1,4] thiazepin
617 ring); Observable second isomer peaks are at 9.84 (1H, s, NH from sec.amide), 7.50-

618 7.27 (5H, m, Ar-H), 7.01-6.89 (2H, m, Ar-H), 6.63 (1H, dd, $J = 8.3 + 1.8$ Hz, Ar-H), 5.30
619 (1H, s, methine from [1, 4] thiazepin ring), 5.04 (2H, s, CH₂), 3.73 (3H, s, OCH₃), 3.30
620 (1H, q, methine from [1, 4] thiazepin ring, this peak is partly covered by water peak
621 from DMSO), 1.68 (3H, s, CH₃ from pyrazole ring), 1.09 (3H, d, $J = 7.2$ Hz, CH₃ from
622 [1,4] thiazepin ring). HRMS; m/z (ES): found 410.1539 (C₂₂H₂₄N₃O₃S [M + H]⁺)
623 requires 410.1460.

624 *4-[4-(2-Chloro-benzyloxy)-3-methoxy-phenyl]-3,6-dimethyl-4,8-dihydro-1H-*
625 *pyrazolo[3,4-*e*][1,4]thiazepin-7-one (1i)*. Mixture of two isomers (72:28), δ H/ppm (400
626 MHz, d⁶-DMSO): Major isomer peaks are at 12.31 (1H, bs, NH from 3-pyrazole ring),
627 9.67 (1H, s, NH from sec.amide), 7.63-7.54 (1H, m, Ar-H), 7.53-7.46 (1H, m, Ar-H),
628 7.42-7.34 (2H, m, Ar-H), 7.02-6.90 (2H, m, Ar-H), 6.77 (1H, dd, $J = 8.3 + 2.0$ Hz, Ar-
629 H), 5.53 (1H, s, methine from [1, 4] thiazepin ring), 5.10 (2H, s, CH₂), 3.72 (3H, s,
630 OCH₃), 3.67 (1H, q, $J = 7.0$ Hz, methine from [1, 4] thiazepin ring), 1.96 (3H, s, CH₃
631 from pyrazole ring), 1.19 (3H, d, $J = 7.0$ Hz, CH₃ from [1,4] thiazepin ring); Observable
632 minor isomer peaks are at 9.84 (1H, s, NH from sec.amide), 7.63-7.54 (1H, m, Ar-H),
633 7.53-7.46 (1H, m, Ar-H), 7.42-7.34 (2H, m, Ar-H), 7.02-6.90 (2H, m, Ar-H), 6.65 ((1H,
634 dd, $J = 8.3 + 1.80$ Hz, Ar-H), 5.31 (1H, s, methine from [1, 4] thiazepin ring), 5.11 (2H,
635 s, CH₂), 3.74 (3H, s, OCH₃), 3.30 (1H, q, methine from [1, 4] thiazepin ring, this peak
636 is partly covered by water peak from DMSO), 1.69 (3H, s, CH₃ from pyrazole ring),

637 1.09 (3H, d, J = 7.10 Hz, CH₃ from [1,4] thiazepin ring). HRMS; m/z (ES): found
638 444.1159 (C₂₂H₂₃ClN₃O₃S [M + H]⁺) requires 444.1070.

639 *4-[4-(3-Chloro-benzyloxy)-3-methoxy-phenyl]-3,6-dimethyl-4,8-dihydro-1H-*

640 *pyrazolo[3,4-e][1,4]thiazepin-7-one (1j)*. Mixture of two isomers (68:32), δH/ppm (400

641 MHz, d⁶-DMSO): First isomer peaks are at 12.30 (1H, bs, NH from 3-pyrazole ring),

642 9.67 (1H, s, NH from sec.amide), 7.52-7.34 (4H, m, Ar-H), 7.03-6.87 (2H, m, Ar-H),
643 6.76 (1H, dd, J = 8.3 + 2.1 Hz, Ar-H), 5.52 (1H, s, methine from [1, 4] thiazepin ring),
644 5.05 (2H, s, CH₂), 3.72 (3H, s, OCH₃), 3.67 (1H, q, J = 7.0 Hz, methine from [1,4]
645 thiazepin ring), 1.95 (3H, s, CH₃ from pyrazole ring), 1.20 (3H, d, J = 7.04 Hz, CH₃
646 from [1,4] thiazepin ring); Observable second isomer peaks are at 9.84 (1H, s, NH
647 from sec.amide), 7.52-7.34 (4H, m, Ar-H), 7.03-6.89 (2H, m, Ar-H), 6.64 (1H, dd, J =
648 8.2 + 1.9 Hz, Ar-H), 5.30 (1H, s, methine from [1, 4] thiazepin ring), 5.10 (2H, s, CH₂),
649 3.74 (3H, s, OCH₃), 1.68 (3H, s, CH₃ from pyrazole ring), 1.09 (3H, d, J = 7.2 Hz, CH₃
650 from [1, 4] thiazepin ring). HRMS; m/z (ES): found 444.1154 (C₂₂H₂₃ClN₃O₃S [M + H]⁺)
651 requires 444.1070.

652 *4-[4-(4-Chloro-benzyloxy)-3-methoxy-phenyl]-3,6-dimethyl-4,8-dihydro-1H-*
653 *pyrazolo[3,4-e][1,4]thiazepin-7-one (1k)*. Mixture of two isomers (68:32), δH/ppm (400
654 MHz, d⁶-DMSO): major isomer peaks are at 12.31 (1H, bs, NH from 3-pyrazole ring),
655 9.66 (1H, s, NH from sec.amide), 7.51-7.41 (4H, m, Ar-H), 7.00-6.87 (2H, m, Ar-H),
656 6.75 (1H, dd, J = 8.4 + 2.0 Hz, Ar-H), 5.51 (1H, s, methine from [1, 4] thiazepin ring),
657 5.03 (2H, s, CH₂), 3.71 (3H, s, OCH₃), 3.67 (1H, q, J = 7.0 Hz, methine from [1,4]
658 thiazepin ring), 1.95 (3H, s, CH₃ from pyrazole ring), 1.19 (3H, d, J = 7.0 Hz, CH₃ from
659 [1,4] thiazepin ring); Observable minor isomer peaks are at 9.84 (1H, s, NH from
660 sec.amide), 7.51-7.41 (4H, m, Ar-H), 7.00-6.87 (2H, m, Ar-H), 6.63 (1H, dd, J = 8.2 +

661 1.8 Hz, Ar-H), 5.30 (1H, s, methine from [1, 4] thiazepin ring), 5.04 (2H, s, CH₂), 3.73
662 (3H, s, OCH₃), 3.30 (1H, q, methine from [1, 4] thiazepin ring, this peak is partly
663 covered by water peak from DMSO), 1.67 (3H, s, CH₃ from pyrazole ring), 1.09 (3H, d,
664 J = 7.2 Hz, CH₃ from [1,4] thiazepin ring). HRMS; m/z (ES): found 444.1157
665 (C₂₂H₂₃N₃O₃S [M + H]⁺) requires 444.1070.

666 4-[3-Methoxy-4-(2-methyl-benzyloxy)-phenyl]-3,6-dimethyl-4,8-dihydro-1H-
667 pyrazolo[3,4-e][1,4]thiazepin-7-one (**1l**). Mixture of two isomers (66:34), δ H/ppm (400
668 MHz, d⁶-DMSO): major isomer peaks are at 12.30 (1H, bs, NH from 3-pyrazole ring),
669 9.67 (1H, s, NH from sec.amide), 7.43-7.34 (1H, m, Ar-H), 7.28-7.14 (3H, m, Ar-H),
670 7.03-6.91 (2H, m, Ar-H), 6.77 (1H, dd, J = 8.3 + 2.0 Hz, Ar-H), 5.52 (1H, s, methine
671 from [1,4] thiazepin ring), 5.00 (2H, s, CH₂), 3.70 (3H, s, OCH₃), 3.67 (1H, q, J = 7.0
672 Hz, methine from [1,4] thiazepin ring), 2.30 (3H, s, *ortho* CH₃), 1.96 (3H, s, CH₃ from
673 pyrazole ring), 1.19 (3H, d, J = 7.0 Hz, CH₃ from [1, 4] thiazepin ring); Observable
674 minor isomer peaks are at 9.84 (1H, s, NH from sec.amide), 7.43-7.34 (1H, m, Ar-H),
675 7.28-7.14 (3H, m, Ar-H), 7.03-6.91 (2H, m, Ar-H), 6.65 (1H, dd, J = 8.2 + 1.8 Hz, Ar-
676 H), 5.30 (1H, s, methine from [1,4] thiazepin ring), 5.02 (2H, s, CH₂), 3.72 (3H, s,
677 OCH₃), 3.30 (1H, q, methine from [1, 4] thiazepin ring, this peak is partly covered by
678 water peak from DMSO), 2.32 (3H, s, *ortho* CH₃), 1.69 (3H, s, CH₃ from pyrazole ring),
679 1.10 (3H, d, J = 7.12 Hz, CH₃ from [1, 4] thiazepin ring). HRMS; m/z (ES): found
680 424.1688 (C₂₃H₂₆N₃O₃S [M + H]⁺) requires 424.1617.

681 4-[3-Methoxy-4-(3-methyl-benzyloxy)-phenyl]-3,6-dimethyl-4,8-dihydro-1H-
682 pyrazolo[3,4-e][1,4]thiazepin-7-one (**1m**). Mixture of two isomers (65:35), δ H/ppm (400
683 MHz, d⁶-DMSO): major isomer peaks are at 12.30 (1H, bs, NH from 3-pyrazole ring),
684 9.66 (1H, s, NH from sec.amide), 7.30-7.10 (4H, m, Ar-H), 7.00-6.87 (2H, m, Ar-H),

685 6.74 (1H, dd, $J = 8.3 + 2.0$ Hz, Ar-H), 5.51 (1H, s, methine from [1, 4] thiazepin ring),
686 5.00 (2H, s, CH₂), 3.71 (3H, s, OCH₃), 3.66 (1H, q, $J = 7.07$ Hz, methine from [1,4]
687 thiazepin ring), 2.31 (3H, s, *meta* CH₃), 1.95 (3H, s, CH₃ from pyrazole ring), 1.19 (3H,
688 d, $J = 7.04$ Hz, CH₃ from [1, 4] thiazepin ring); Observable minor isomer peaks are at
689 9.84 (1H, s, NH from sec.amide), 7.30-7.10 (4H, m, Ar-H), 7.00-6.87 (2H, m, Ar-H),

690 6.62 (1H, dd, $J = 8.4 + 2.0$ Hz, Ar-H), 5.30 (1H, s, methine from [1,4] thiazepin ring),
691 5.00 (2H, s, CH₂), 3.73 (3H, s, OCH₃), 3.30 (1H, q, methine from [1,4] thiazepin ring,
692 this peak is partly covered by water peak from DMSO), 2.31 (3H, s, *meta* CH₃), 1.68
693 (3H, s, CH₃ from pyrazole ring), 1.09 (3H, d, $J = 7.2$ Hz, CH₃ from [1, 4] thiazepin ring).
694 HRMS; m/z (ES): found 424.1687 (C₂₃H₂₆N₃O₃S [M + H]⁺) requires 424.1617.

695 *4-[3-Methoxy-4-(4-methyl-benzyloxy)-phenyl]-3,6-dimethyl-4,8-dihydro-1H-*
696 *pyrazolo[3,4-e][1,4]thiazepin-7-one (1n)*. Mixture of two isomers (62:38), δ H/ppm (400
697 MHz, d⁶-DMSO): major isomer peaks are at 12.29 (1H, bs, NH from 3-pyrazole ring),
698 9.66 (1H, s, NH from sec.amide), 7.35-7.26 (2H, m Ar-H), 7.24-7.14 (2H, m, Ar-H),
699 7.00-6.87 (2H, m, Ar-H), 6.74 (1H, dd, $J = 8.3 + 2.1$ Hz, Ar-H), 5.51 (1H, s, methine
700 from [1,4] thiazepin ring), 4.97 (2H, s, CH₂), 3.70 (3H, s, OCH₃), 3.66 (1H, q, $J = 7.0$
701 Hz, methine from [1,4] thiazepin ring), 2.29 (3H, s, *para* CH₃), 1.95 (3H, s, CH₃ from
702 pyrazole ring), 1.18 (3H, d, $J = 7.0$ Hz, CH₃ from [1, 4] thiazepin ring); Observable
703 minor isomer peaks are at 9.84 (1H, s, NH from sec.amide), 7.35-7.26 (2H, m Ar-H),
704 7.24-7.14 (2H, m, Ar-H), 7.00-6.87 (2H, m, Ar-H), 6.62 (1H, dd, $J = 8.3 + 1.8$ Hz, Ar-
705 H), 5.30 (1H, s, methine from [1,4] thiazepin ring), 4.98 (2H, s, CH₂), 3.72 (3H, s,
706 OCH₃), 3.29 (1H, q, methine from [1, 4] thiazepin ring, this peak is partly covered by
707 water peak from DMSO), 2.30 (3H, s, *para* CH₃), 1.68 (3H, s, CH₃ from pyrazole ring),

708 709

1.0 (3H, d, J = 7.12 Hz, CH₃ from [1, 4] thiazepin ring). HRMS; m/z (ES): found
9 424.1687 (C₂₃H₂₆N₃O₃S [M + H]⁺) requires 424.1617.

710 *4-(4-Isopropoxy-phenyl)-3,6-dimethyl-4,8-dihydro-1H-pyrazolo[3,4-e][1,4]thiazepin-7-*
711 *one (1o)*. Mixture of two isomers (97:3), δH/ppm (400 MHz, d⁶-DMSO): major isomer
712 peaks are at 12.30 (1H, s, NH from 3-pyrazole ring), 9.64 (1H, s, NH from sec.amide),
713 7.15 (2H, d, J = 8.68 Hz, Ar-H), 6.80 (2H, d, J = 8.76 Hz, Ar-H), 5.54 (1H, s, methine

714 from [1, 4] thiazepin ring), 4.54 (1H, hept, CH of isopropyl), 3.67 (1H, q, J = 7.0 Hz,
715 methine from [1, 4] thiazepin ring), 1.90 (3H, s, CH₃ from pyrazole ring), 1.23 (6H, d, J
716 = 6.0 Hz, (CH₃)₂ of isopropyl), 1.18 (3H, d, J = 7.08 Hz, CH₃ from [1,4] thiazepin ring);
717 Observable minor isomer peaks are at 9.83 (1H, s, NH from sec.amide), 6.85 (2H, d, J
718 = 8.7 Hz, Ar-H), 5.30 (1H, s, methine from [1, 4] thiazepin ring), 1.67 (3H, s, CH₃ from
719 pyrazole ring), 1.07 (3H, d, J = 7.1 Hz, CH₃ from [1, 4] thiazepin ring). HRMS; m/z
720 (ES): found 332.1428 (C₁₇H₂₂N₃O₂S [M + H]⁺) requires 332.1354.

721 *3,6-Dimethyl-4-(4-propoxy-phenyl)-4,8-dihydro-1H-pyrazolo[3,4-e][1,4]thiazepin-7-one*
722 **(1p)**. Mixture of two isomers (85:15), δH/ppm (400 MHz, d⁶-DMSO): major isomer
723 peaks are at 12.30 (1H, s, NH from 3-pyrazole ring), 9.65 (1H, s, NH from sec.amide),
724 7.16 (2H, d, J = 8.7 Hz, Ar-H), 6.82 (2H, d, J = 8.7 Hz, Ar-H), 5.55 (1H, s, methine
725 from [1, 4] thiazepin ring), 3.87 (2H, t, J = 6.52 Hz, OCH₂ of n-propyl), 3.67 (1H, q, J =
726 7.0 Hz, methine from [1, 4] thiazepin ring), 1.90 (3H, s, CH₃ from pyrazole ring), 1.76-
727 1.62 (2H, m, CH₂ of n-propyl), 1.19 (3H, d, J = 7.0 Hz, CH₃ from [1,4] thiazepin ring),
728 0.95 (3H, t, J = 7.45 Hz, CH₃ of n-propyl); Observable minor isomer peaks are at 9.84
729 (1H, s, NH from sec.amide), 7.13 (2H, d, J = 8.64 Hz, Ar-H), 6.86 (2H, d, J = 8.72 Hz,
730 Ar-H), 5.30 (1H, s, methine from [1, 4] thiazepin ring), 3.90 (2H, t, J = 6.56 Hz, OCH₂
731 of n-propyl), 3.23 (1H, q, J = 7.17 Hz this peak is partly covered by water peak from
732 DMSO), 1.76-1.62 (2H, m, CH₂ of n-propyl), 1.06 (3H, d, J = 7.20 Hz, CH₃ from [1, 4]

733 thiazepin ring), 0.97 (3H, t, J = 7.47 Hz, CH₃ of n-propyl); HRMS; m/z (ES): found
734 332.1428 (C₁₇H₂₂N₃O₂S [M + H]⁺) requires 332.1354.

735 *4-(4-Allyloxy-phenyl)-3,6-dimethyl-4,8-dihydro-1H-pyrazolo[3,4-e][1,4]thiazepin-7-one*
736 (**1q**). Mixture of two isomers (97:3), δH/ppm (400 MHz, d⁶-DMSO): major isomer
737 peaks are at 12.31 (1H, s, NH from 3-pyrazole ring), 9.65 (1H, s, NH from sec.amide),

738 7.17 (2H, d, J = 8.7 Hz, Ar-H), 6.85 (2H, d, J = 8.7 Hz, Ar-H), 6.09-5.94 (1H, m, =CH of
739 allyl), 5.55 (1H, s, methine from [1, 4] thiazepin ring), 5.37 [1H, dq, J = 1.71 + 17.25
740 Hz, =CH₂ (H_a-terminal vinyl proton on SP² carbon)], 5.23 [1H, dq, J = 1.4/1.6 + 10.5
741 Hz, =CH₂ (H_b-terminal vinyl proton on SP² carbon)], 4.52 [1H, t, J = 1.48, O-CH₂ (H_a-
742 allyl proton adjacent to SP² carbon)], 4.51 [1H, t, J = 1.48, O-CH₂ (H_b-allyl proton
743 adjacent to SP² carbon)], 3.67 (1H, q, J = 7.1 Hz, methine from [1, 4] thiazepin ring),
744 1.90 (3H, s, CH₃ from pyrazole ring), 1.19 (3H, d, J = 7.1 Hz, CH₃ from [1,4] thiazepin
745 ring), Observable minor isomer peaks are at 9.84 (1H, s, NH from sec.amide), 6.87
746 (2H, d, J = 8.72 Hz, Ar-H), 5.31 (1H, s, methine from [1, 4] thiazepin ring), 1.67 (3H, s,
747 CH₃ from pyrazole ring), 1.07 (2H, d, J = 7.1 Hz, CH₃ from [1, 4] thiazepin ring),
748 HRMS; m/z (ES): found 330.1272 (C₁₇H₂₀N₃O₂S [M + H]⁺) requires 330.4180.

749 **Compounds preparation:** Benznidazole (Bz) (N-benzyl-2-nitroimidazole-1-
750 acetamide) was obtained from the Pharmaceutical Laboratory of the State of
751 Pernambuco (LAFEPE, Brazil) and used in all trials as reference drug. All the CYP51
752 inhibitor compounds were purchased from Asinex commercial vendor, diluted in
753 dimethylsulfoxide (DMSO - *in vitro*) or in trappsol (*in vivo*) not reaching vehicle levels
754 above 0.6 % and 20 % for *in vitro* and *in vivo* assays, respectively.

755 **T. cruzi parasites:** Bloodstream trypomastigotes (BT - Y strain) were obtained from
756 heart puncture of infected male Swiss Webster mice at parasitemia peak. The
757 parasites were purified and resuspended in RPMI 1640 medium (pH 7.2 to 7.4)

758 without phenol red (Gibco BRL) supplemented with 10% fetal bovine serum (FBS) and
759 2 mM glutamine, as reported previously [26, 27] The analysis upon intracellular forms
760 *T. cruzi* (Tulahuen strain transfected with *Escherichia coli* β -galactosidase gene) [28]

761 was conducted using L929 cell lineages that were infected with tissue culture-derived
762 trypomastigotes using a 10:1 parasite/ host cell ratio [27].

763 ***In vitro* assays**

764 **Mammalian cell culture:** The cardiac cells (CC) were obtained from Swiss embryos
765 Webster mice as described by Meirelles and coworkers [29], and seeded 96 well plate.

766 The cardiac cell cultures were stored at 37°C in Du Ibeyco's modified Eagle medium
767 (DMEM; without phenol red; Sigma-Aldrich) supplemented with 5% fetal bovine serum,
768 2.5 mM CaCl₂, 1 mM L-glutamine, streptomycin, and 2% chicken embryo extract [22,
769 29]. Moreover, L929 fibroblastic cells were cultivated (4 x10³ cells/well in 96-well
770 microplates) at 37°C in RPMI 1640 medium (pH 7.2 to 7.4) without phenol red (Gibco
771 BRL) supplemented with 10% fetal bovine serum and 1% glutamine, as reported
772 previously [27, 30].

773 **Cytotoxicity assays:** To evaluate the toxicity profile and determine the selectivity
774 index (SI), cardiac and L929 cells were incubated at 37°C with crescent
775 concentrations of the studied compounds for 24 and 96 h and cellular viability were
776 evaluated by colorimetric analysis using Prestoblue [27] and Alamarblue [30] assays
777 as reported, respectively. After 5 and 6 h of incubation respectively, the absorbance
778 were measured at 570 and 600 nm according to manufactures instructions and the

779 results confirmed by the analysis of morphology and physiology aspects through light
780 microscopy. The results are expressed as the percent of reduced viability in
781 compound-treated and vehicle-treated samples by following the manufacturer's
782 instructions and the EC_{50} values calculated (concentration that reduces the cellular
783 viability by 50%) [26, 27, 30].

784 **Trypanocidal effects:** BT forms of Y strain (5×10^6 /mL) were incubated for 2 and 24 h
785 at 37°C in absence or presence of crescent concentrations (0 – 50 μ M) of each tested
786 compound. The compounds were diluted in RPMI 1640 medium (Roswell Park
787 Memorial Institute - Sigma Aldrich – USA) supplemented with 5% FBS. Subsequently
788 the incubation, death rates were assessed by light microscopy quantification using a
789 Neubauer chamber to determine the EC_{50} that correspond the compound
790 concentration that reduces in 50% of the number of the parasite population [26,27,
791 31]. For the effect against intracellular forms (Tulahuen- β -galactosidase strain), L929
792 cells were infected for 2h, rinsed using saline to remove non-internalized parasites and
793 then incubated for 48 h at 37°C to establish the culture infection. The infected L929
794 cell cultures were exposed first to a fixed concentration of tested compounds (10-12
795 μ M). Those that presented ≥ 50 % of reductions in the parasite load, were further
796 screened using infected cultures submitted increasing non-toxic concentrations of the
797 selected compounds to determine the EC_{50} values. Next, the absorbance was
798 measured at 570 nm and results expressed as the percentage of *T. cruzi* growth
799 inhibition in compound-tested cells compared to untreated cells. In parallel, the
800 therapy using the reference drug (Bz) was always performed as reported [26].

801 ***In vivo* assay**

802 **Acute toxicity:** NOAEL (no-observed-adverse-effect level) was obtained by injecting,
803 via intraperitoneal (ip), increasing concentrations of the tested compound (up to 200
804 mg/kg of mice body weight) in female Swiss Webster mice (weight, 20 to 23 g; *n* - 2
805 mice per assay for two assays). Treated animals were inspected for toxic and subtoxic
806 symptoms according to the Organization for Economic Cooperation and Development

807 (OECD) guidelines. At 48 h after compound injection, the NOAEL values were
808 determined as reported previously [31].

809 **Mice infection and treatment schemes:** Male Swiss Webster mice (18 to 20 g)
810 obtained from the animal facilities of Instituto de Ciência e Tecnologia em Biomodelos
811 (ICTB) from Fundação Oswaldo Cruz (FIOCRUZ) were housed at a maximum of 6
812 animals per cage, and kept in a specific-pathogen-free (SPF) room at 20 to 24°C
813 under a 12- light and 12-h dark cycle, providing sterilized water and chow *ad libitum*.

814 The animals were acclimated for 7 days before starting the experiments. Infection was
815 performed by i.p. injection of 10^4 bloodstream trypomastigotes (Y strain). Compound **1**
816 was first dissolved in DMSO and then freshly diluted with 20% of Trappsol (CTD, Inc.,
817 USA) and Bz was prepared in sterile distilled water with 3% Tween 80 (Sigma-
818 Aldrich). The animals were divided into the following groups: uninfected (noninfected
819 and untreated), untreated (infected with *T.cruzi* but treated only with vehicle), and
820 treated (infected and treated) i.p. with nontoxic doses of 5 to 20 mg/kg/day test
821 compound or with 100 mg/kg/day Bz p.o. The mouse received a 0.1-ml i.p. dose, the
822 treatment started at 5th day post infection (dpi) that represent the parasitemia onset in
823 this experimental model and followed for five consecutive days, until 9th dpi. For Bz
824 treatment, infected mice received a 0.1 mL p.o. following the same therapeutic
825 schemes as described above [19].

826 **Parasitemia and mortality rates:** Parasitemia was individually detected by Pizy-
827 Brener [32] methodology using direct light microscopy to quantify the number of
828 parasites in 5 μ L of blood collected from tail vein, and the mice were checked for
829 mortality daily until 30 days post treatment [31]. Mortality was expressed by the
830 percentage of cumulative mortality (CM) as described by Batista and coworkers [22].

831 **Ethics**

832 All procedures were carried out in accordance with the guidelines established by the
833 FIOCRUZ Committee of Ethics for the Use of Animals (CEUA LW16/14).

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