

1 *Trypanosoma cruzi* produces the specialized proresolving mediators Resolvin D1, Resolvin
2 D5 and Resolvin E2

3

4 **Running Title:** *T. cruzi* and Resolvins

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37 **ABSTRACT**

38 *Trypanosoma cruzi* is a protozoan parasite that causes Chagas disease (CD). CD is a persistent,
39 life-long infection affecting many organs, most notably the heart where it may result in acute
40 myocarditis and chronic cardiomyopathy. The pathological features include myocardial
41 inflammation and fibrosis. In the Brazil-strain infected CD-1 mouse that recapitulates many of
42 the features of human infection, we found increased plasma levels of resolvin D1 (RvD1), a
43 specialized pro-resolving mediator of inflammation, both during the acute and chronic phases of
44 the infection (>100 days post infection) as determined by ELISA. Additionally, ELISA on
45 lysates of trypomastigotes of both the Tuliahan and Brazil strains revealed elevated levels of
46 RvD1 when compared with lysates of cultured epimastigotes of *T. cruzi*, tachyzoites of
47 *Toxoplasma gondii*, and trypomastigotes of *T. brucei*, cultured L₆E₉ myoblasts and culture media
48 containing no cells. Lysates of *T. cruzi* –infected myoblasts also displayed increased levels of
49 RvD1. Lipid mediator metabolomics confirmed that the trypomastigotes of *T. cruzi* produced
50 RvD1, RvD5 and RvE2, which have been demonstrated to modulate the host response to
51 bacterial infections. Plasma levels of RvD1 maybe both host and parasite derived. Since *T. cruzi*
52 synthesizes specialized pro-resolving mediators of inflammation as well as pro-inflammatory
53 eicosanoids, such as thromboxane A₂, one may speculate that by using these lipid mediators to
54 modulate its microenvironment, the parasite is able to survive.

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58 **Key words:** *Trypanosoma cruzi*, Chagas disease, Resolvins, Resolvin D1, Resolvin D5,
59 Resolvin E2, inflammation, immune modulation

60 **List of Abbreviations:**

61 ELISA, Enzyme Linked ImmunoSobrant Assay; FFA, free fatty acid; AA, arachidonic acid;
62 DHA, docosahexenoic acid; EPA, eicosapentaenoic acid; HDHA, hydroxyl-docosahexenoic
63 acid; HETE, hydroxy-eicosatetraenoic acid; HEPE, hydroxy-eicosapentaenoic acid; PG,
64 prostaglandin; TXA₂, thromboxane A₂; TP, Receptor for TXA₂; FP, receptor for PGF_{2α}; LT,
65 leukotriene; LX, lipoxin; SPM, specialized pro-resolving mediator; RvD, resolvins derived from
66 DHA; RvE, resolvins derived from EPA; PD, protectin; MaR, maresin; EFA, essential fatty
67 acid; LO, lipoxygenase; IFN-γ, interferon -γ; PBMC, peripheral blood mononuclear cell; H&E,
68 Hematoxylin and Eosin; MRI, magnetic resonance imaging; ECG, electrocardiogram; LM, lipid
69 metabolite; LC, Liquid chromatography; MS, mass spectrometry; dpi, days post-inoculation;
70 NDGA, nordihydroguaiaretic acid; TcOYE, *T. cruzi* old yellow enzyme; TbPGFS, *T. brucei*
71 PGF_{2α} synthase.

72 **INTRODUCTION**

73 The parasite *Trypanosoma cruzi* is the etiologic agent of Chagas disease. It is endemic in
74 Mexico, Central and South America where millions of persons are infected or at risk (1). In
75 recent years, due to migration of individuals from endemic areas, there has been increased
76 recognition of Chagas disease in the United States, Europe and other non-endemic areas (2-4).
77 Additionally, vector transmission of this parasite has been documented in the United States (5).
78 Although any nucleated cell in the body can be infected, the cardiovascular system is among the
79 organs targeted by this parasite causing acute myocarditis and chronic cardiomyopathy.

80 Acute infection with *T. cruzi* results in an intense inflammatory response associated with an
81 increased expression of pro-inflammatory mediators including cytokines, chemokines and
82 endothelin-1(6). We have also previously described the release of the bioactive lipid mediators
83 prostaglandin $F_{2\alpha}$ ($PGF_{2\alpha}$) and thromboxane A_2 (TXA_2) from *T. cruzi*, although TXA_2 is
84 preferentially synthesized (7). TXA_2 is a pro-inflammatory, vasoconstrictor with pro-thrombotic
85 activity through enhanced platelet activation-aggregation. $PGF_{2\alpha}$ is a pro-inflammatory
86 vasoconstrictor substance like TXA_2 (8); however, it opposes platelet activation by TXA_2 and
87 platelet activating factor (9). Both receptors (TP and FP) couple to $G_{\alpha q}$ and $G_{\alpha 13}$, activate the
88 small GTPase Rho, and promote cellular activation; however, FP expression on leukocytes is not
89 well documented suggesting the effects of FP on inflammation are likely indirect while the
90 effects of TP are direct (8). A previous report indicates that TP can be activated by $PGF_{2\alpha}$ (10),
91 suggesting a single focal point for the action of pro-inflammatory prostaglandins in Chagas
92 disease and re-enforcing the prominence of TP to the pathogenesis of disease.

93 Acute inflammation is a natural host protective mechanism mounted by the body in response to

94 injury or invading pathogens. When self-limited, it restores homeostasis (11); however, if left
95 uncontrolled, excessive inflammation can lead to chronic tissue damage (12). Resolution of
96 inflammation is an active process orchestrated by a genus of potent molecules known as
97 specialized pro-resolving mediators (SPM)(13) . SPM, that include the resolvins, protectins and
98 maresins, are bioactive autacoids with both anti-inflammatory and pro-resolving properties. They
99 are enzymatically produced from essential fatty acids (EFA), with distinct stereochemistries.
100 RvD1 biosynthesis involves sequential oxygenations of DHA by 15-lipoxygenase (LO) and 5-
101 LO (14). Using a total organic synthetic approach the complete stereochemistry of RvD1 has
102 been established as 7S,8R,17S-trihydroxy- 4Z,9E,11E,13Z,15E,19Z-docosahexaenoic acid (15).
103 RvD1 displays potent leukocyte-directed actions in pico-nanogram concentration range *in vivo*
104 (14). In *E. coli*-induced peritonitis, RvD1 reduces bacterial titers and hypothermia, increases
105 survival and enhances microbial containment and killing by phagocytes (16). The aspirin
106 triggered epimer of RvD1, has been found to regulate the host immune response in Chagas
107 disease patients by decreasing IFN- γ and the percentage of necrotic cells in the peripheral blood
108 mononuclear cell (PBMC) pool, and reducing the rate of *T. cruzi* antigen-stimulated PBMC
109 proliferation in cultured cells (17). The source and relevance of resolvins to *T. cruzi* infection is
110 currently uninvestigated.

111 The full complement of the *T. cruzi* lipidome is yet to be determined. We asked whether *T. cruzi*
112 infection resulted in up regulation of SPM expression and whether there were contributions to
113 the pro-resolution pathway from both host and parasite. We hypothesized that infection with *T.*
114 *cruzi* results in an increased expression of resolvins both *in vitro* and *in vivo*. Indeed, herein we
115 demonstrate that mice infected with *T. cruzi* display increased levels of resolvin D1 (RvD1) and
116 that the trypomastigote form of the parasite contains RvD1, RvD5 and RvE2. Thus, it is of

117 interest that *T. cruzi* synthesized both pro-inflammatory (TXA₂) and SPM (resolvins) that are
118 essential regulators of host response that promote transition to chronic infection. This
119 observation may have new implications for the pathogenesis of Chagas disease.

120 **Materials and Methods**

121 **Animal Ethics Statement:** All animal experiment were performed with approval of the
122 Institutional Animal Care and Use Committee of the Albert Einstein College of Medicine.

123 **Parasitology and Pathology:** The Brazil and Tulahuen strains of *T. cruzi* were maintained by
124 serial passage in C3H mice (Jackson Laboratories, Bar Harbor, ME). For studies with the Brazil
125 strain 8-week old male CD-1 mice were infected with 5×10^4 parasites. Parasitemia was
126 determined by counting in an hemocytometer chamber as previously described (18).
127 Trypomastigotes of both the Brazil and Tulahuen strains of *T. cruzi* were propagated in a
128 myoblast cell line (L₆E₉) as previously described(19). The RH strain of *Toxoplasma gondii* was
129 maintained in cultured human foreskin fibroblasts. The *T. brucei brucei* Lister 427 strain was
130 provided to us by Dr. George Cross (The Rockefeller University, New York, NY). Hearts were
131 fixed in 10% (v/v) neutral buffered formalin and sections (5 μ m) were stained with Hematoxylin
132 and Eosin (H&E).

133 **Cardiac Magnetic Resonance Imaging (MRI):** For magnetic resonance imaging (MRI), mice
134 were anesthetized with isoflurane inhalation (2% (v/v) in medical air administered via a nose
135 cone). MRI compatible electrocardiogram (ECG) electrodes were inserted in the left front and
136 right rear paws and the ECG signal was used as a trigger signal with a Small Animal Instruments
137 physiological monitoring system (Stony Brook, NY). Mice were positioned in a 35-mm ID 1H
138 volume coil [Molecules2Man Imaging Co., Cleveland, OH(20)]. Body temperature was
139 maintained at 34~35°C using warm air with feedback from a body surface thermocouple and a
140 small respiratory sensor balloon, taped onto the abdomen, provided for respiratory gating.
141 Images were acquired using a 9.4 T Varian Direct Drive animal magnetic resonance imaging and
142 spectroscopic system (Agilent Technologies, Inc. Santa Clara, CA). One mm thick slices were

143 acquired in short-axis orientation using an ECG-triggered and respiratory gated multi-frame
144 tagged cine sequence. The imaging parameters used were field of view (FOV) of 40×40 mm²,
145 matrix size of 256×256, TE of 2.6 ms, TR of 5.5 ms, flip angle of 25°, number of averages of 2.

146 **General considerations regarding ELISA and LC-MS/MS assays:** *T. cruzi* epimastigotes
147 were cultured in LIT medium which we have assayed and contains no resolvins (by both ELISA
148 and LC-MS-MS). They were washed 3 times in PBS (containing no resolvins) and lysates were
149 made from these epimastigotes. Trypomastigotes of *T. cruzi* were obtained from infected
150 myoblast cultures which we confirmed contained no resolvins (by both ELISA and LC-MS-MS).
151 Lysates of *T. brucei* trypomastigotes and tachyzoites of the RH strain of *Toxoplasma gondii*
152 prepared in similar fashion contained no resolvins indicating the resolvins content of *T. cruzi* was
153 not an artifact of culture conditions. Positive controls for the LC-MS/MS were the known
154 authentic standards for the resolvins. Negative controls were the L₆E₉ myoblasts which we found
155 to have no detectable amounts of resolvins. The limits of detection of this LC-MS-MS system ~
156 10-15 picomoles. Resolvin (Rv)D1 was determined in lysates of 1×10⁶ trypomastigote of both
157 the Tulahuén and Brazil strains of *T. cruzi* and trypomastigotes of *T.b. brucei* according to
158 manufacturer's specifications (Cayman Chemical Company, Ann Arbor, MI, USA). Blood was
159 collected from infected CD-1 mice by retro-orbital bleed, centrifuged and frozen at -80°C until
160 use. RvD1 levels for all experiments were determined according to absorbance measured
161 between 405 - 420 nm via a microplate reader.

162 **Lipid mediator (LM) metabololipidomics:** LM metabololipidomics was performed as
163 described (Colas et al, 2014). After hypotonic lysis, parasites were further sonicated and snap
164 frozen (stored at -80°C until analyzed). Briefly, parasite lysates were placed in ice-cold methanol
165 (containing deuterium labeled internal standard (d₅-RVD2, d₄-PGE₂, d₅-LXA₄ and d₄-LTB₄,

166 500pg each) to facilitate quantification and recovery. Samples were held at -20°C for 45 minutes
167 to allow protein precipitation and centrifuged. Supernatants were collected and placed onto an
168 automated extraction system (Biotage) using a C18 column. Bioactive LMs were collected and
169 injected into an LC system (Shimadzu) coupled to a Qtrap 5500 (ABSciex). Identification was
170 conducted using published criteria including matching retention time to synthetic standards and a
171 minimum of 6 diagnostic ions in the MS-MS mode as reported (21).

172 **Statistical analysis:** All other data are expressed as the mean (\pm SEM) and were analyzed using
173 GraphPad Prism statistics software (GraphPad Software Inc., San Diego, CA). For analysis of
174 differences between groups the Student's *t* test was performed. A level of significance of 5% was
175 chosen to denote differences between means.

176 **Results**

177 **Animal studies**

178 **Parasitology, Pathology and Cardiac Imaging**

179 Parasitemia was first detected at 15 days post infection (dpi), peaked at $4-5 \times 10^5$
180 trypanomastigotes/ml at 35 to 40 dpi and waned by 60 dpi when parasitemia was undetectable
181 (Figure 1A). Mortality was 50% during acute infection (Figure 1B). At 100 dpi cardiac imaging
182 demonstrated an enlarged heart and a significant increase in the right ventricular internal
183 diameter consistent with previous reports (20) (Figure 1C and D). Histopathological examination
184 of the heart during the acute phase revealed myonecrosis and parasite pseudocysts (Figure 1E).
185 During the chronic phase there was cardiac myocyte hypertrophy, chronic inflammation and
186 fibrosis, consistent with previous reports (6).

187

188 **Plasma Resolvin D1 increases during acute *T. cruzi* infection**

189 There was a rapid rise in RvD1 ($\mu\text{g/ml}$ of plasma) in infected mice that was evident at 20 dpi and
190 remained elevated out to 140 dpi during the chronic phase of infection (Figure 2A). This profile
191 suggests that RvD1 is derived both from parasite and host. To assess this hypothesis we
192 examined RvD1 levels in parasite lysates.

193

194 **Resolvin D1 levels in parasite extracts as determined by ELISA**

195 Trypanomastigotes of both the Brazil and Tulahuen strains of *T. cruzi* had significantly greater
196 amounts of RvD1 compared to epimastigotes of the Tulahuen (Figure 2B) and Brazil (data not
197 shown) strains. RvD1 was not detected at significant levels in trypanomastigotes of *T. brucei*,
198 tachyzoites of *T. gondii* or from uninfected L₆E₉ myoblasts. In contrast, myoblasts infected with

199 the Tulahuen strain of *T. cruzi* (which contained intracellular amastigotes) produced significant
200 amounts of RvD1 compared to uninfected myoblasts. Although production was lower than in
201 free trypomastigotes these data suggest that intracellular amastigotes synthesize RvD1 or that
202 infection stimulates production by myoblasts. To unequivocally determine that RvD1 was being
203 produced we performed LC-MS-MS analysis, the gold standard for identification of bioactive
204 lipid mediators.

205

206 **LM metabololipidomics**

207 The above results clearly indicated that RvD1 is present in trypomastigotes of *T. cruzi*. Figure 3
208 demonstrates the characteristics LC-MS-MS spectra used for the identification of RvD1, RvD5
209 and RvE2 in parasite extracts. The resulting lipidomic profiles are representative of 3 separate
210 parasite lysates from both Tulahuen and Brazil strains. Extracts of uninfected L₆E₉ myoblasts
211 and LIT media were negative for lipid species suggesting all identified species were parasite
212 derived (data not shown). Both strains had the precursor lipids docosahexenoic acid (DHA),
213 eicosapentaenoic acid (EPA) and arachidonic acid (Table 1) indicating that biosynthesis of all
214 pro-inflammatory and pro-resolving mediators were possible. DHA and its bioactive
215 metabolome were present in significantly higher quantities indicating a potential preference for
216 biosynthesis of mediators derived from this fatty acid. Consistent with earlier reports from our
217 laboratory and others(8) we identified the arachidonic acid metabolites PGE₂, PGD₂ and PGF_{2 α}
218 in *T. cruzi* lysates (Table 2). The presence of 5/12/15-HETE and 5/12/15/18-HEPE in extracts
219 (Table 3) suggests active lipoxygenase pathways are present in the parasite with arachidonic acid
220 and EPA as the respective precursor FFAs(22). Potential downstream pro-inflammatory end
221 products might also include leukotrienes; however, no leukotriene B₄ production was identified

222 (Table 3). Thus, the primary class of pro-inflammatory lipid species produced by *T. cruzi* is
223 prostaglandins.
224
225 An active lipoxygenase pathway would also initiate and promote the biosynthesis of pro-
226 resolving lipid mediators such as lipoxins, resolvins, maresins and protectins. When we
227 examined lysates by LC-MS-MS no evidence of lipoxins, protectins or maresins were observed
228 (Table 4). However, their pathway markers were identified indicating that trypomastigotes of *T.*
229 *cruzi* have the enzymatic capability to produce these mediators. Of interest, RvD1 and RvD5
230 were biosynthesized by both the Tulahuén and Brazil strains of *T. cruzi* (Table 4). Moreover, the
231 Brazil strain produced significant levels of RvE2 (Table 4) indicating that only the Brazil strain
232 utilized EPA to produce pro-resolving mediators. The precursor of RvE2 (18-HEPE) is lower in
233 the Brazil strain perhaps reflecting increased use of this precursor for RvE2 synthesis. These
234 studies clearly indicate that RvD1 is present in trypomastigotes of *T. cruzi* and that D and E
235 series resolvins are the likely primary lipid mediators through which *T. cruzi* dampen the host
236 response to infection.
237

238 **Discussion**

239 The heart is an important focus of infection by *T. cruzi*, although other tissues and organs are
240 infected as well. Acute *T. cruzi* infection is characterized by an upregulation of pro-inflammatory
241 cytokines and chemokines and an influx of inflammatory cells. In the setting of Chagas disease,
242 as in other disease states, the acute inflammatory response is a “double-edged” sword, in that, it
243 is needed to control the infection, but also leads to tissue injury. The chronic phase of Chagas
244 disease is accompanied by the appearance of chronic inflammatory cells and cardiac remodeling
245 (i.e., fibrosis). The present study clearly demonstrates, for the first time, that *T. cruzi*
246 trypomastigotes and amastigotes produce RvD1 which likely contributes to the resolution of
247 inflammation. This was confirmed by both ELISA and LM metabolomic profiling. In addition,
248 *T. cruzi*-infected mice display elevated circulating RvD1 levels during the acute phase of
249 infection and well into the chronic phase when peripheral parasitemia has waned and, in the
250 mouse model, cardiomyopathy is present. Lysates of trypomastigotes of the Brazil and Tulahuen
251 strains contained RvD5 while trypomastigotes of the Brazil strain contained RvE2. We did not
252 identify RvD1 in a variety of other protozoan parasites (*T. gondii* and *T. brucei*) and non-infected
253 mammalian cells.

254 Historically, the resolution of inflammation was considered a passive process resulting from the
255 loss or dilution of pro-inflammatory mediators from the extracellular milieu. It is now
256 understood, however, that the resolution of inflammation is instead, an active and programmed
257 event (13). The mediators of the resolution of inflammation, SPM (include the resolvins,
258 protectins and maresins), are produced enzymatically from essential fatty acids (EFA). RvD1 is
259 one of these bioactive mediators and as noted in *E. coli*-induced peritonitis, RvD1 reduces
260 bacterial titers and hypothermia, increases survival and enhances microbial containment and

261 killing by phagocytic cells (16). RvD5 has also been shown to be involved in the setting of
262 experimental *E. coli* by enhancing phagocytosis. Combination of RvD5 with the antibiotic
263 ciprofloxacin was able to accelerate the antibiotic effect and the same was observed in the setting
264 of experimental Staphylococcal infection (16). RvE2, identified by LM metabololipidomics, has
265 potent leukocyte-directed actions that regulates chemotaxis of human neutrophils, enhancing
266 phagocytosis and anti-inflammatory cytokine production. RvE2 rapidly down-regulates surface
267 expression of human leukocyte integrins in whole blood and dampened responses to platelet-
268 activating factor. These actions appear to be mediated by leukocyte G-protein-coupled receptors.
269 Collectively, these observations indicate that RvE2 can stimulate host-protective actions
270 throughout initiation and resolution in the innate inflammatory responses (23).

271 The ability of the parasite and host to liberate pro-resolving mediators is likely to alter the course
272 of disease. The identification of 15- and 5(S),15 (S)-HETE in *T. cruzi* extracts suggests 15-LO
273 activity (which has been previously reported in parasites such as *T. gondii*)(24), however, this
274 has not been previously documented and no data is available comparing disease progression in
275 15LO^{-/-} mice or with pharmacological antagonists of 15LO activity. Conversely, a comparison
276 of 5LO activity has been performed (25). 5LO activity is essential for synthesis of many pro-
277 resolving mediators (such as E-series resolvins and lipoxins). Direct comparison of 5-LO null
278 mice (intact parasite RvD1 synthesis) and infected mice treated with NDGA (1.25
279 mg/mouse/day), a 5LO inhibitor (inhibiting both mouse and parasite RvD1) indicated that
280 NDGA treatment promoted earlier development of parasitemia and greater lethality than the 5LO
281 null mice (25) although number of cardiac amastigote nests and anti-oxidant protection were
282 similar. These differences suggest that NDGA may have inhibited parasite- and host-derived
283 5LO activity and that parasite 5LO derivatives may play a significant role in suppressing early

284 parasite growth during acute infection and promote survival of the host through the acute
285 infection. Our data suggest that of the available mediators only RvE2 production would be
286 affected by such a difference (as no lipoxin nor leukotriene production was observed in the *T.*
287 *cruzi* strains tested). These data indicate the liberation of SPMs from the parasite may be
288 important in mediating the transition from acute to chronic disease and promoting host survival
289 during experimental *T. cruzi* infection.

290 Earlier, our groups reported that *T. cruzi* synthesizes the lipid mediators TXA₂ and PGF_{2α}(7),
291 TXA₂ is both a potent vasoconstrictor and a pro-inflammatory mediator synthesized by both
292 trypomastigotes and amastigotes (7, 22). The majority of TXA₂ in the serum of infected mice is
293 derived from the parasite (7). In the present report, we have demonstrated that trypomastigotes
294 and amastigotes of both the Brazil and Tulahuen strains of *T. cruzi* strains contain RvD1, RvD5
295 and RvE2, important pro-resolving mediators. Plasma RvD1 levels are elevated in *T. cruzi*
296 (Brazil strain) infected mice and it is possible that the source of the RvD1 is, in part, from the
297 parasite itself. The synthesis of pro-inflammatory and pro-resolving lipid mediators suggests
298 that the parasite is able to modulate its microenvironment through these metabolites. If the net
299 result is a damping down of the inflammatory response, this may be important in the
300 perpetuation of the infection into the chronic phase. There are other anti-inflammatory factors
301 present during *T. cruzi* such as interleukin-10. In that regard, it was recently demonstrated that
302 there is an interaction between RvD1 and interleukin-10 in the resolution of inflammation in
303 adipose tissue in the experimental obese state (26). These observations suggest that the addition
304 of such an anti-parasitic regimen may ameliorate some of the consequences of this infection.
305 Resolvins dampen the inflammatory response and promote resolution of infections, which
306 prevents fibrosis, and enables parasite persistence and the perpetuation of the parasite life cycle

307 (27).

308 The biosynthesis of lipid mediators by *T. cruzi* is increasingly complex but may provide
309 advantages for longevity of infection. The parasite genome encodes terminal synthases for some
310 eicosanoid species and the ability of *T. cruzi* to elevate the host plasma levels of lipid mediators
311 in mice knocked out for the respective terminal synthase (such as the TXA₂ synthase null mice)
312 suggest these pathways are active during disease (7). While *T. cruzi* proteins such as PGF_{2α}
313 synthase are highly homologous to other prokaryotic enzymes (such as yeast old yellow enzyme
314 (TcOYE) and *T. brucei* synthase (TbPGFS))(28-30) their primary sequences bare significant
315 differences to mammalian terminal enzymes (29, 30). Moreover, the *T. cruzi* enzymes are
316 resistant to pharmacological inhibitors of mammalian terminal synthases indicating that the
317 active sites are also topographically or structurally different (28). Thus, parasite-derived
318 eicosanoids manipulate host responses during infection but are intractable to current therapies.

319 Most interestingly, the terminal synthase for the primary *T. cruzi* prostaonoid TXA₂ has eluded
320 genomic efforts to identify it prompting suggestions that the parasite may co-opt host synthetic
321 pathways by the process of trogocytosis (31). Similarly, the 15LO and 5LO activities needed for
322 D and E series resolvins biosynthesis are not predicted from the *T. cruzi* genome. It therefore
323 remains to be determined whether the biosynthetic pathways exist in *T. cruzi* or are
324 serendipitously stolen from infected host cells. Only the examination of chronically infected null
325 mice, once parasite numbers have declined, is likely to yield these answers.

326 Previously, we reported that *T. cruzi* produces the pro-inflammatory and pro-thrombotic lipid
327 TXA₂. Herein we now demonstrate that this parasite also produces several pro-resolving lipids.
328 Taken together, this parasite may be able to modulate its microenvironment and enhance

329 persistence. On the other hand, the administration of RvD1 may ameliorate the inflammation and
330 fibrosis in the heart. These experiments are currently underway.

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333

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- 421

FFA Precursor Species			Trypomastigotes of <i>T. cruzi</i> strains (range, pg/mg protein)	
	Q1	Q3	Tulahuen	Brazil
EPA	301	257	1.6---1194 [^]	2.9---1937 [^]
DHA	327	283	113---15111 [^]	62.1---3841 [^]
4S,14S-diHDHA	359	221	*	*
17-HDHA	343	245	9.7---199.3	6.4--242.4
14-HDHA	343	205	15.7---86.8	14.9--141.6
7-HDHA	343	141	16.9--262.4	11.3---88.2
4-HDHA	343	101	94.5---427.4	70.0--471.3
AA	303	259	15.5---2529 [^]	32---5766 [^]

422

423 **Table 1: Quantitation of free fatty acid (FFA) precursors in lysates of trypomastigotes by**424 **LC-MS/MS.** Expression of lipid species are quantified from peak height and expressed a pg/mg

425 protein. Data represent the range of samples documented from 3 independent lysates for each

426 strain. * represents samples with values below the detection limit, ^ = samples are in ng/mg

427 protein. EPA,; DHA,; HDHA, hydroxy Docosaheaxaenoic Acid; AA, .arachidonic acid

428

Prostaglandin Lipid Species			Trypomastigotes of <i>T. cruzi</i> strains (range, pg/mg protein)	
	Q1	Q3	Tulahuen strain	Brazil strain
PGD ₂	351	189	1.4-3.9	2.7-23
PGE ₂	351	189	2.0-21.4	6.0-48.6
PGF _{2α}	353	193	2.6-38.9	2.7-2.6

429

430 **Table 2: Quantitation of prostaglandin species in lysates of trypomastigotes of *T. cruzi* by**
431 **LC-MS/MS.** Expression of lipid species are quantified from peak height and expressed a pg/mg
432 protein. Data represent the range of samples documented from 3 independent lysates for each
433 strain. PG, prostaglandin.

434

Lipoxygenase-derived Lipid Species	Q1	Q3	Trypomastigotes of <i>T. cruzi</i> strains (range, pg/mg protein)	
			Tulahuen strain	Brazil strain
LTB ₄	335	195	*	*
20-OH-LTB ₄	351	195	*	*
20-COOH-LTB ₄	365	195	*	*
5-HETE	319	115	54.6-101.6	26.3-122.1
12-HETE	319	179	11.1-46.9	15.3-143.3
15-HETE	319	219	24.8-166.4	39.7-328.1
5S,15S-diHETE	335	235	2.3-5.2	5.7-8.0
5-HEPE	317	115	5.4-12.9	2.5-31.0
12-HEPE	317	179	0.7-6.1	0.6-1.0
15-HEPE	317	219	2.5-3.3	2.3-3.4
18-HEPE	317	259	3.8-5.4	1.7-2.2
5S,15S-diHEPE	333	115	1.3-27.7	1.7-13.6

435 **Table 3: Quantitation of lipoxygenase-derived lipid species in lysates of *T. cruzi***
436 **trypomastigotes by LC-MS/MS.** Expression of lipid species are quantified from peak height
437 and expressed a pg/mg protein. Data represent the range of samples documented from 3
438 independent lysates for each strain. * represents samples with values below the detection limit.
439 LT, leukotriene; HETE,; hydroxy-eicosatetraenoic acid; diHETE, di-hydroxy-eicosatetraenoic
440 acid; HEPE, hydroxy-eicosapentaenoic acid; diHEPE, di-hydroxy- eicosapentaenoic acid

Pro-Resolving Lipid Species	Q1	Q3	Trypomastigotes of <i>T. cruzi</i> (range, pg/mg protein)	
			Tulahuen strain	Brazil strain
RvD1	375	141	1.2-1.4	1.8-7.0
RvD2	375	215	*	*
RvD3	375	147	*	*
RvD5	359	199	1.4-1.6	0.7-1.9
RvD6	359	159	*	*
RvE1	349	161	*	*
RvE2	333	253	*	9.5-23.6
RvE3	333	201	*	*
PD1	359	153	*	*
22-OH-PD1	375	153	*	*
22-COOH-PD1	389	153	*	*
MaR1	359	250	*	*
LXA ₄	351	115	*	*
LXB ₄	351	115	*	*
LXA ₅	349	215	*	*
LXB ₅	349	221	*	*

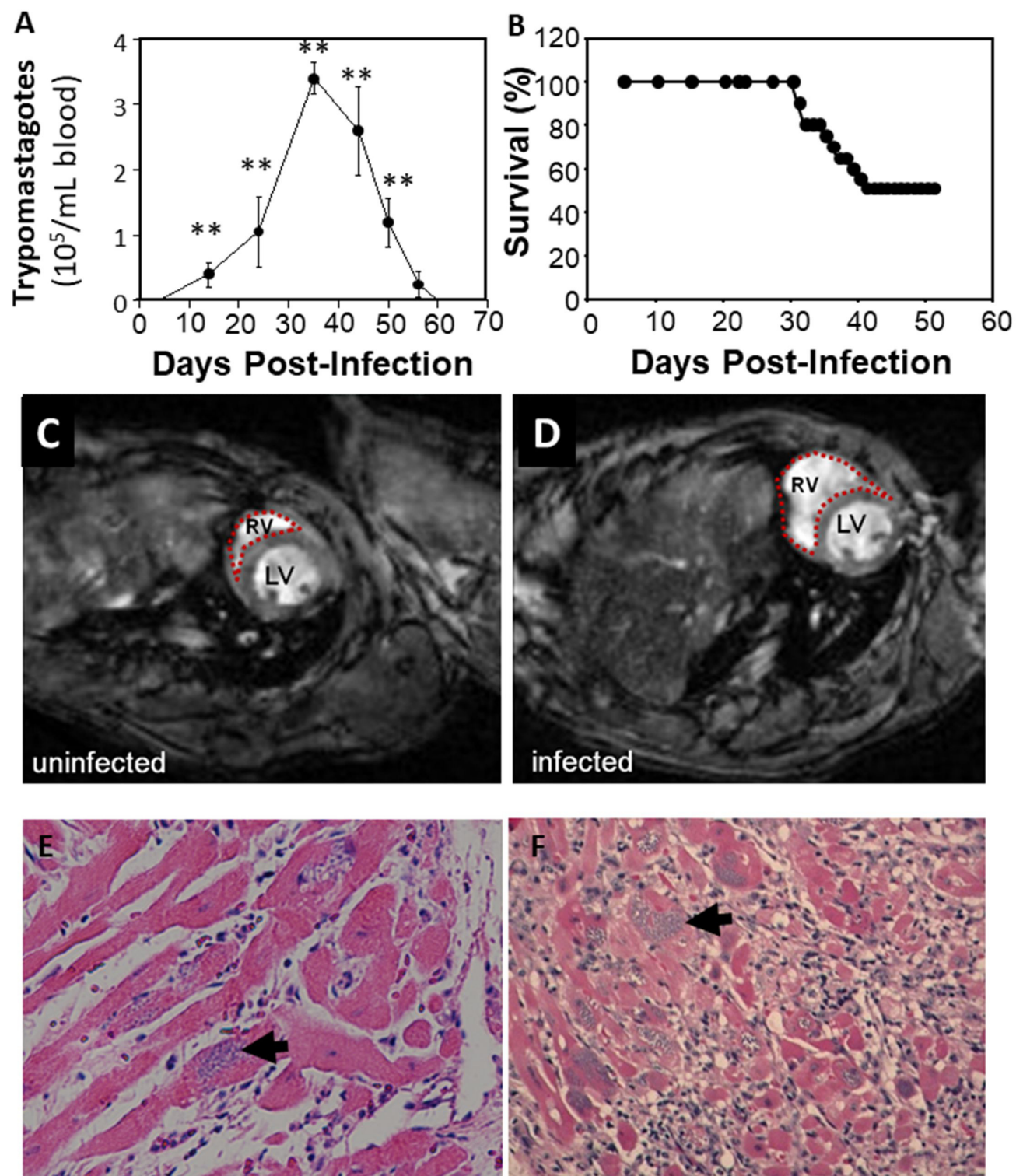
441 **Table 4: Quantitation of pro-resolving lipid mediators in lysates of trypomastigotes of *T.***
442 ***cruzi* by LC-MS/MS.** Expression of lipid species are quantified from peak height and expressed
443 a pg/mg protein. Data represent the range of samples documented from 3 independent lysates for
444 each strain.
445 * represents samples with values below the detection limit. RvD, resolvins derived from
446 docosahexenoic acid; RvE, resolvins derived from eicosapentaenoic acid; PD, protectin; MaR,
447 maresin; LX, lipoxin.
448

449 **FIGURES**

450 **Figure 1. Kinetics of cardiomyopathy in *T. cruzi* infection.** Parasitemia (A) and mortality (B)
451 in CD-1 mice infected with 5×10^4 trypomastigotes of the Brazil strain of *T. cruzi*. **C-D.**
452 Representative cardiac MRI of control (C) and infected (D) mouse heart revealing an enlarged
453 right ventricle. **E-F.** Representative histopathology of the heart during the acute and chronic
454 infection showing pseudocysts of amastigotes (◀) and inflammation. Images are representative
455 of n=5 mice in each group.

456 **Figure 2. *T. cruzi* are a source of RvD1 during infection.** A. Serum RvD1 levels in mice
457 inoculated with 5×10^4 Brazil strain trypomastigotes. Data are mean \pm SD (n=5). ** represents
458 significance ($p \leq 0.05$) from uninfected mice. B. Release of RvD1 from *T. cruzi*, related protists,
459 and infected L₆E₉ cells. RvD1 release was measured by ELISA (pg/mL) from conditioned
460 media. Data are mean \pm SD (n=3). ** represents significance ($p \leq 0.05$) from epimastigotes and
461 uninfected L6E9 cells. epis= epimastigotes forms of Tulahuen strain, trypos=trypomastigotes of
462 the Brazil and Tulahuen strains, Toxo= lysates of the RH strain of *Toxoplasma gondii*, T.
463 *brucei*= *Trypanosma. brucei*. L₆E₉ myoblasts infected (Inf) or uninfected (Uninf) with
464 trypomastigotes of the Tulahuen strain of *T. cruzi*.

465 **Figure 3. Representative spectra used to identify lipid mediators of the resolving class.** LC-
466 MS-MS fragmentation spectra employed for the identification of RvD1, RvD5 and RVE2 in
467 lysates of trypomastigotes of *T. cruzi* from Brazil and Tulahuen strains.



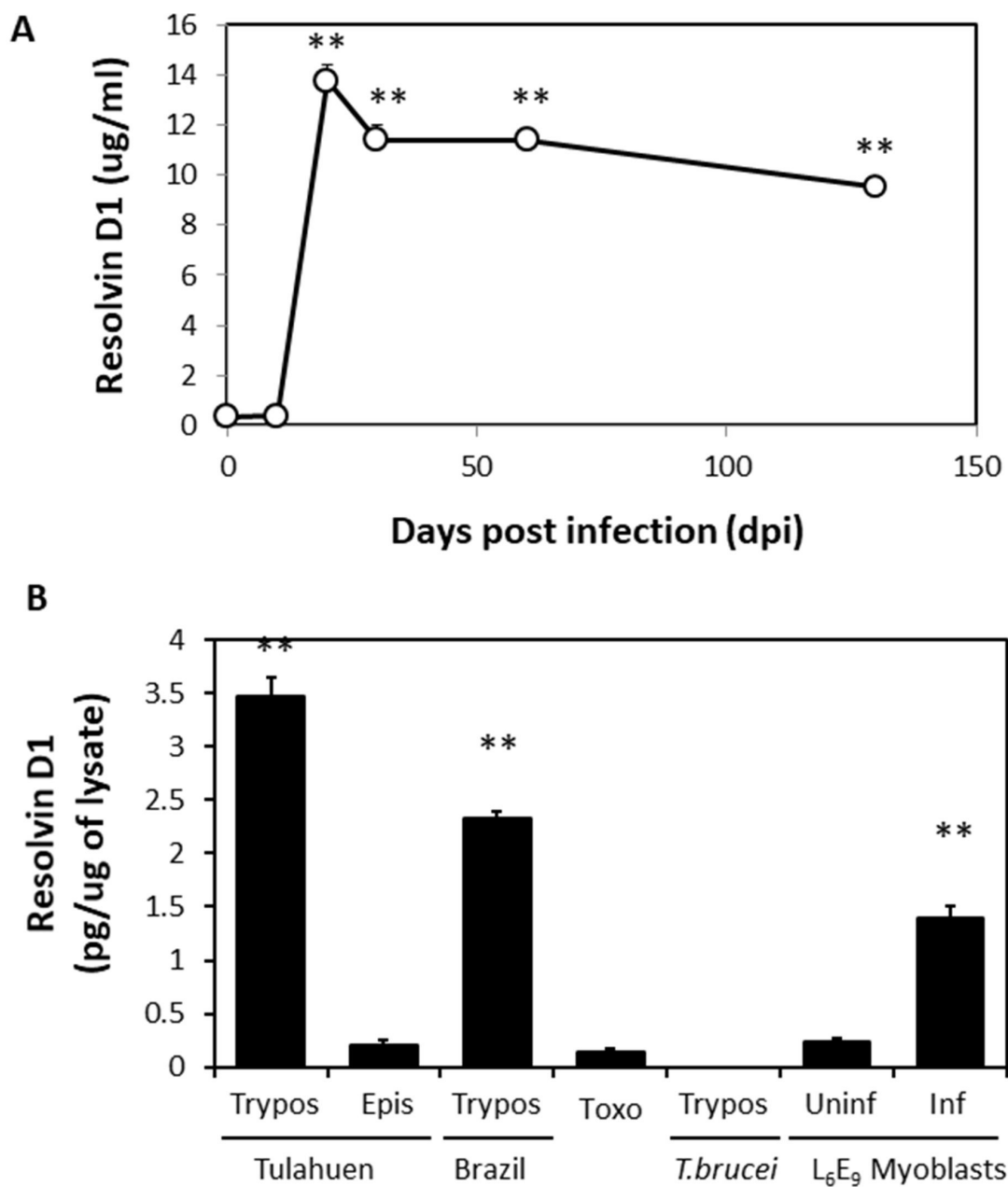


Figure 2

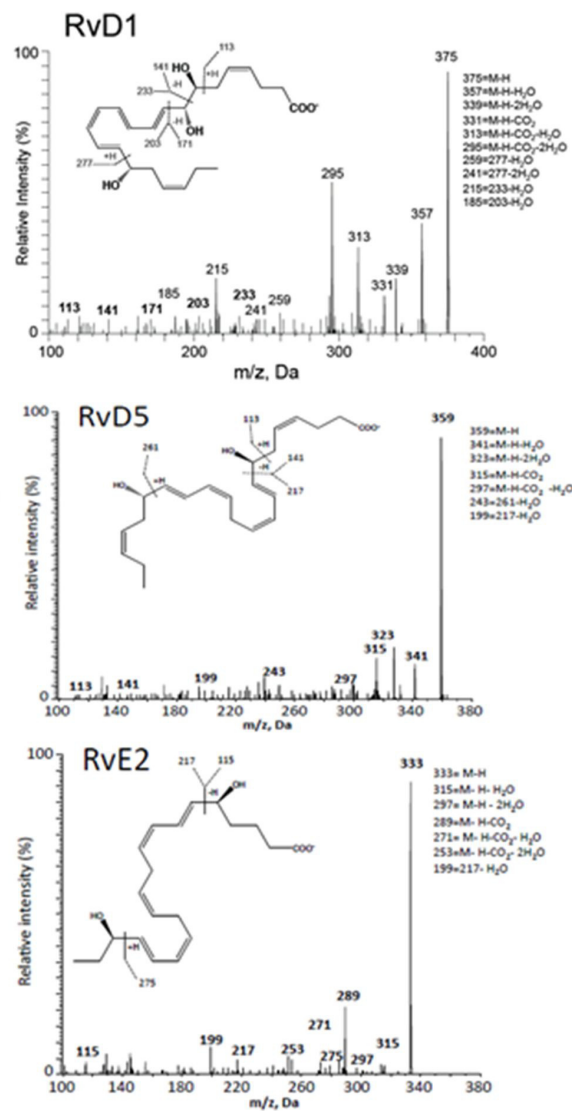
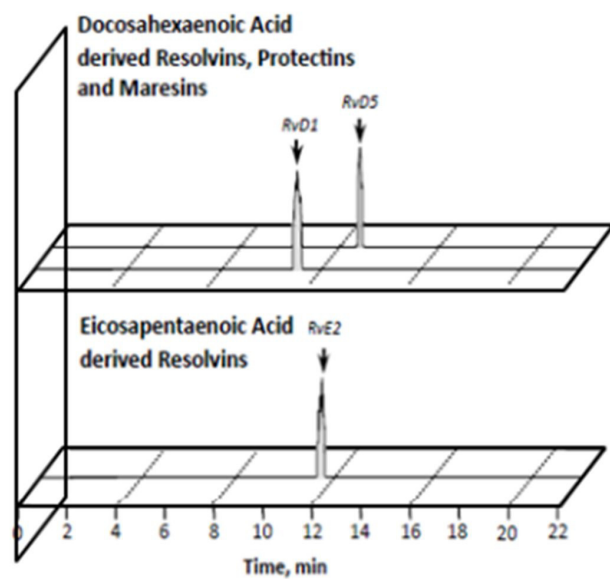


Figure 3