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Posted Date: 6 December 2023

doi: 10.20944/preprints202312.0371.v1

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Article

Anti-Trypanosomal Bufadienolides from the Oocytes of the Toad *Rhinella alata* (Anura, Bufonidae)

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Abstract: Amphibians are widely known as a prolific source of bioactive metabolites. In this work we isolated and characterized compounds with antiparasitic activity from the oocytes of the toad *Rhinella alata* collected in Panama. Bio-guided isolation and structural elucidation was carried out using chromatographic and spectroscopic techniques. The organic crude extract was subjected to solid phase extraction followed by HPLC purification of the fraction with *in vitro* activity against *Trypanosoma cruzi* trypomastigotes. Seven steroids (1-7) of the bufadienolide family were isolated and their structures determined using NMR and MS analyses; of these 19-formyl-dyscinobufotalin (3) is reported as a new natural product. Compounds 1, 3-7, resulted with a good anti-trypanosomal activity profile, among these 16β-hydroxyl-hellebrigenin (1) and bufalin (7) showed significant selectivity for *T. cruzi*. Furthermore, molecular docking analysis showed compounds 1, 3 and 7 interact through H-bonds with the amino acid residues GLN-19, ASP-158, HIS-159 and TRP-177 from cruzipain, at the catalytic site. Given the lack of therapeutic options to currently treat the American trypanosomiasis, this work can serve as the basis for further studies that aim the development of bufadienolides or their derivatives as drugs against Chagas disease.

Keywords: *Rhinella alata*; toads; Bufonidae; oocytes; bufadienolides; anti-trypanosomal activity; *Trypanosoma cruzi*

1. Introduction

Chagas disease, or American trypanosomiasis, is a zoonotic disease caused by the parasite *Trypanosoma cruzi*. This protozoan is transmitted to humans mainly through the feces of insects of the family Reduviidae [1]. Although Chagas disease is recognized as endemic from Latin America, it has become a worldwide issue with 10,000 deaths per year and eight million people at risk of infection [2]. There are no specific symptoms for diagnostics during its acute phase; however, once the disease advances, chronic pathological features such as cardiomyopathy, megacolon, megaesophagus and stroke are developed [3]. The low effectiveness and side effects of the available drugs used for the

treatment of this disease, during its chronic phase, highlight the need of more efficacious and safer anti-trypanosomal treatments [4].

Bufadienolides are polyhydroxy steroids that contain an unsaturated 2H-pyran-2-one ring at β -17 position [5]. Because of their wide bioactivity against human cancer cells lines and their great chemical diversity, bufadienolides have attracted a considerable attention in recent years [6]. These metabolites are widely distributed in nature and have been reported in fireflies, mammals, plants, snakes and toads [7]. Bioprospecting research has revealed that bufadienolides isolated from some amphibian species, have promising antibiotic and antiparasitic properties [8].

Herein we report the bio-guided isolation of anti-trypanosomal bufadienolides from the oocytes of the toad *Rhinella alata* (Thominot, 1884) from Panama [9]. Seven compounds were isolated and their chemical structures were elucidated by spectroscopic techniques. Compounds were tested against trypomastigotes of *T. cruzi* and evaluated on Vero cells for mammalian cytotoxicity. Furthermore bufadienolides 1, 3 and 7 were analyzed by molecular docking with cruzipain revealing chemical interactions with amino acid residues at the active site.

2. Results and Discussion

2.1. Isolation and structural elucidation of bufadienolides

The crude extract from oocytes of *R. alata* was prepared by maceration with a mixture of methanol and chloroform and then submitted to C-18 SPE purification. The fraction eluted at 80 % CH₃OH showed anti-trypanosomal activity and was further analyzed by semi-preparative RP-HPLC (Figure S1). Compounds 1-7 were isolated and their structures, depicted in Figure 1, were determined based on extensive spectroscopic analysis and comparison of their data with those reported in the literature. While compounds 1, 2 and 4-7 corresponded to known compounds, the bufadienolide 3 is reported as a new natural product.

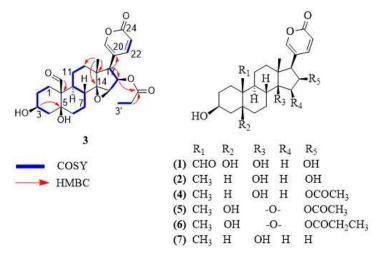


Figure 1. Chemical structures of bufadienolides 1-7.

Compound **3** was isolated as a white solid. The molecular formula was determined as $C_{27}H_{34}O_8$ based on its HR-MALDI-MS data which showed the molecular ion at m/z 487.2283 [M + H]⁺ (calculated for $C_{27}H_{35}O_8$, 487.2326). Eleven degrees of unsaturation were inferred from the molecular formula. The ¹H-NMR and ¹³C-NMR spectra of compound **3** showed the feature signals for a 2H-pyran-2-one moiety [δ_H 6.24 (1H, dd, J = 0.8, 9.7 Hz), 7.36 (1H, m), 8.02 (1H, m); δ_C 114.2, 118.2, 150.9, 153.6, 164.0], a tertiary methyl [δ_H 0.81 (3H, s); δ_C 17.3], a secondary methyl [δ_H 0.94 (3H, t, J = 7.8 Hz); δ_C 9.3], three oxygenated methines [(δ_H 3.75, m; δ_C 60.7), (δ_H 4.15, m; δ_C 68.0), (δ_H 5.47, m; δ_C 76.3)], two oxygenated quaternary carbons [δ_C 75.8, 73.1] and two carbonyl carbons characteristic of aldehyde and ester [δ_C 210.1, 174.9]. Comparison of the NMR signals (Table 1) of compound **3** with those of the reported bufadienolide dyscinobufotalin [10]. revealed the data of the steroidal nucleus

of the two compounds were very similar, indicating that compound 3 had the same bufadienolide core with two carbinols at C-3 and C-5, an epoxy group at C-14 and C-15, and the esterification of the hydroxyl group at C-16 with propionic acid. The main difference consisted in the presence of an aldehyde group at C-19 in compound 3 instead of the angular methyl group at C-19 of dyscinobufotalin [10]. HMBC correlations between aldehyde proton at $\delta_{\rm H}$ 10.00 and the quaternary carbon at C-10 allowed for the assignment of the aldehyde group at C-19 (Figure 1). On the basis of spectral data, the compound 3 was identified as 19-formyl-dyscinobufotalin.

Table 1. NMR (¹H 400 MHz, ¹³C 100 MHz) data for compound 3 in methanol-d4 (δ in ppm, *J*, in Hz).

No.	¹³ C, Type	¹ H, Mult. (<i>J</i> in Hz)
1a	18.6, CH ₂	2.26, m
1a 1b	16.6, C112	1.77, m
2a	27.5, CH ₂	1.77, m 1.77, m
2a 2b	27.3, C112	
3	69.0 CH	1.77, m
3 4a	68.0, CH	4.15, m 1.99, m
4a 4b	36.7, CH ₂	
5	75.8, C	1.61, m
6a	38.3, CH ₂	2.23, m
6b	38.3, CH2	1.49, m
7a	24.1, CH ₂	1.49, m 1.77, m
7a 7b	24.1, C112	1.77, m 1.11, m
8	43.1, CH	1.77, m
9	34.4, CH	2.47, td (3.8, 12.2)
10	56.0, C	2.47, ta (3.0, 12.2)
11a	22.9, CH ₂	1.66, m
11b	22.5, C112	1.39, m
12a	40.5, CH ₂	1.77, m
12b	10.0, C112	1.49, m
13	46.0, C	1115) 111
14	73.1, C	
15	60.7, CH	3.75, m
16	76.3, CH	5.47, m
17	51.2, CH	2.93, d (9.2)
18	17.3, CH₃	0.81, s
19	210.1, CH	10.00 s
20	118.2, C	
21	153.6, CH	7.36, m
22	150.9, CH	8.02, m
23	114.2, CH	6.24, dd (0.8, 9.7)
24	164.0, C=O	
1'	174.9, C=O	
2'a	28.1, CH ₂	2.23, m
2'b		2.11, m
3'	9.3, CH ₃	0.94, t (7.8)

4

Furthermore, six steroids of the bufadienolide family were identified as 16β -hydroxylhellebrigenin (1) [11], desacetyl-bufotalin (2) [5], bufotalin (4) [5], cinobufotalin (5) [12], dyscinobufotalin (6) [10], and bufalin (7) [5].

2.2. Anti-trypanosomal activity and mammalian cytotoxicity

The trypanosomaticidal activity of the bufadienolides 1-7 was evaluated in vitro against $T.\ cruzi$. The ability of these compounds to inhibit growth of the trypomastigote form was evaluated at four different concentrations. The results showed that bufadienolides 1, 3, 4, 5, 6 and 7 reduce the growth of $T.\ cruzi$ while compound 2 was inactive at the screening concentration of 10 µg/mL. The bufadienolides 3 and 7 presented EC50 values (0.19 and 3.93 µM, respectively) being compound 3 even more active than the positive control benznidazole (Table 2). The cytotoxicity of bufadienolides against mammalian cells was also tested $in\ vitro$ using healthy epithelial kidney Vero cells. The IC50 values of bufadienolides 1-7 ranged from 0.02 to > 135 µM. As shown below, compounds 1, 2 and 4 presented the lowest cytotoxic effects. Both the antiparasitic and cytotoxic activities exhibited by each bufadienolide assessed are shown in Table 2. Selectivity was analyzed by the cytotoxicity on Vero cells/anti-trypanosomal activity ratio.

Table 2. Anti-trypanosomal activity and mammalian cytotoxic of bufadienolides **1-7** isolated from the oocytes of *Rhinella alata*.

Commound	T. cruzi	Vero cell	Selectivity (CC50/EC50)	
Compound	EC ₅₀ (μM)	CC50 (µM)		
1	22.80 ± 4.11	> 135	> 5	
2	Inactive	> 28		
3	0.19 ± 0.07	0.22 ± 0.09	1.16	
4	19.60 ± 2	> 46	> 2	
5	27.80 ± 3.77	1.88 ± 0.20	0.07	
6	12.30 ± 1.87	0.02 ± 0.002	< 0.01	
7	3.93 ± 0.57	10.61 ± 1.54	2.69	
benznidazole	2.73 ± 0.10	Not tested		
doxorubicin	Not tested	0.23		

2.3. Molecular docking of bufadienolides 1, 3 and 7

A docking protocol was validated and its reproducibility confirmed by the root mean square deviation (RMSD) values obtained. The RMSD for each cruzipain-co-crystallized inhibitor complex were as follows: 2.3586 for 1EWO-VSC, 1.9060 for 2AIM-ZRA, 1.6817 for 1U9Q-186 and 2.9319 for 4XUI-2VC. In addition, relative docking scores for each complex resulted in -7.0484, -6.9388, -8.2353 and -8.1351, respectively. After validation of the protocol accuracy, interactions of anti-trypanosomal bufadienolides 1, 3 and 7 with the active site of proteins were studied via docking calculations. Table 3 shows the calculated parameters for the best pose obtained for each bufadienolide tested. The energy variation for the predictions obtained by simulation ranged from -2.2 to +1.3 kcal/mol. The isolated bufadienolides 1, 3 and 7 displayed interactions with cruzipain. Bufadienolide 1 showed H-bonds with ASP-158 and GLN-19 at a distance of 2.10 Å and 2.32 Å, respectively. On the other hand, bufadienolide 3 presents three H-bonds with amino acid residues GLN-19, HIS-159 and TRP-177. Bufadienolide 7 formed one H-bond with ASP-158 at a distance of 1.95 Å (Figure 2).

Table 3. Calculations for molecular docking of bufadienolides with cruzipain.

Bufadienolide	RMSD (Å)	Docking Score	Ligand group	Aminoacid residue	Energy (kcal/mol)	Distance (Å)
1	2.5812	-4.2777	CH=O at 19	GLN-19	-1.5	2.32
			OH at 14	ASP-158	-1.7	2.10
3	2.0099	-5.3984	CH=O at 19	GLN-19	-2.2	2.09
			CH=O at 19	TRP-177	-0.5	2.59
			OH at 5	HIS-159	-1.5	2.35
7	3.0381	-4.1206	OH at 14	ASP-158	+1.3	1.95

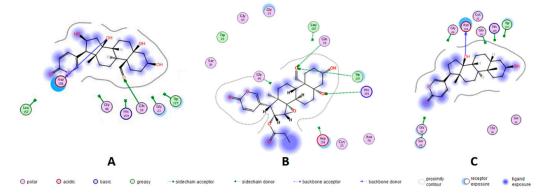


Figure 2. Representation in 2D format for molecular docking of bufadienolides **1**, **3** and **7** with active site amino acid residues of cruzipains. X-ray crystallographic structures of the proteins were extracted from the Protein Data Bank (PDB). Bufadienolide/cruzipain PDB ID= A: 16β -hydroxyl-hellebrigenin (**1**)/1EWO; B: 19-formyl-dyscinobufotalin (**3**)/1EWO; and C: bufalin (**7**)/2AIM.

Currently, there is no effective chemotherapeutic treatment available for American trypanosomiasis [4]. Poisonous terrestrial vertebrates are rarely considered for natural product research programs; however, it has been reported that bioactive materials derived from animals, have resulted in effective medicines, highlighting the potential of vertebrate animals' toxins in pharmacology. For instance, teprotide a nonapeptide isolated from the venom of the snake *Bothrops jararaca*, has served as template to produce captopril and other angiotensin-converted enzyme inhibitors that currently are routinely medicated for hypertension patients [13]. The alkaloid epibatidine obtained from the skin of the dendrobatid frog *Epipedobates tricolor* presents analgesic properties 200 times more potent than morphine. Epibatidine has served as template for producing anti-nociceptive derivatives such as tebanicline (ABT-594) that reached clinical phase II [14,15].

In this study, through the analysis of the spectroscopic data, we elucidated the chemical structure of the new bufadienolide 3 from the oocytes of *R. alata*. Moreover, five bufadienolides (2, 4-7) were previously reported in amphibians and one in plants (1). Bufadienolide 2 was isolated from the poison of the toad *Duttaphrynus melanostictus* as a minor constituent [5]. Compounds 4 and 5 have been obtained from skin and parotoid gland secretions of bufonids, but not in toads of the genus *Rhinella* [16]. The bufadienolide 6 was isolated from Chan Su, a traditional chinese remedy prepared with the poison of Asian toads [10]. The bufadienolide 7 has been identified in bufonid extracts including oocytes [17].

Due to the marked inhibitory activity against the pump (Na $^+$ -K $^+$)ATPase, the antiparasitic potential of bufadienolides against human pathogens has currently received some attention [18]. In this study, bufadienolides **1**-7 were evaluated in vitro for inhibition of the growth of the trypomastigote form of *T. cruzi*. According to our data, substitution of methyl at C-19 by formyl group increases the anti-trypanosomal activity as seen in bufadienolide **3** in comparison with bufadienolide **6**. Similarly, in a previous study hellebrigenin showed trypanocidal activity at 91.7 μ g/mL while telocinobufagin, which contains a C-19 methyl group instead, was inactive [19]. Likewise, comparing the activities of bufadienolides **2** and **7**, it can be inferred that the lack of a hydroxyl group at C-16 is

required for the anti-trypanosomal activity, however if this hydroxyl group is esterified, a moderate bioactivity is observed (compounds **4-6**, Table 2).

Also, previous evaluations on cancer cell lines with bufadienolides revealed a decrease in growth inhibition by introduction of a hydroxyl group at C-16 position [20]. Bufadienolide 7 presents higher anti-trypanosomal activity than its 19-OH analogue (IC50 = 19.4 μ M) [21]. On the basis of these results, we can suggest that the presence of a formyl group at C-19 and the esterification of the hydroxyl group at C-16 are essential elements required for the effective biomolecular interactions with components present in the trypomastigote form of *T. cruzi*, which led to the anti-protozoal activity observed.

In mammals, bufadienolides block the transport activity of the (Na*-K*)ATPase pump by binding to the α -subunit. This inhibition provokes that intracellular sodium concentration increases until cells get depolarized [22]. In T. cruzi, there is no report on the identification of this protein; maintenance of the ionic steady-state is carried out by K+ channels, Na+ efflux pumps and (H⁺)ATPases [23]. Kyoichi, I. et al. [24] cloned and characterized a (Na⁺)ATPase from a gene encoding of T. cruzi. After biochemical evaluations it was evidenced that the enzymatic activity of this pump was not inhibited by the steroid (cardenolide) ouabain. In the search for anti-trypanosomal drugs, the most considered targets for Chagas disease are cruzipain, sterol- 14α -demethylase, transsyalidase, trypanothione reductase and tubulin [4,24]. Currently, the mechanism by which bufadienolides cause inhibition of T. cruzi is unknown, despite of previous findings suggest that bufadienolides inhibit the growth of T. cruzi parasites by a biochemical pathway other than the blockage of the (Na*-K*)ATPase pump. In this study, in silico analysis by molecular docking was carried out, using the *T. cruzi* protease cruzipain. Redocking with previously reported co-crystallized inhibitors was employed for protocol validation and active site prediction (Figure S9). Molecular docking with the anti-trypanosomal bufadienolides 1, 3 and 7 revealed strong interactions with cruzipains. These interactions correspond to H-bonds between amino acid residues GLN-19, ASP-158, HIS-159 and TRP-177 with hydroxyl groups at C-5 and C-14 position; and formyl at C-19 position of bufadienolides. The most potent anti-trypanosomal bufadienolide (3), presented three interactions with cruzipain while bufadienolides 1 and 7 showed two and one H-bonds, respectively. Bufadienolide 7 interacts with cruzipain by the hydroxyl at C-14, a functional group essential for bioactivity [25]. No interaction with polar groups at C-16 such as hydroxyl and propionyloxy were observed for bufadienolides 1 and 3. Cruzipain contains, at the active site, a catalytic triad formed by GLY-23, CYS-25 and GLY-65 [26]. Furthermore, some amino acid residues have been revealed to be involved in enzymatic catalysis of cruzipain such as residues GLN-19, GLY-66, ASP-158, HIS-159 and TRP-177 that keep peptidyl inhibitors anchored to the active site [27]. A molecular docking analysis with 173 compounds reported to be inhibitors of cruzipain showed that in addition to the catalytic triad, the amino acid residues GLN-19 and ASP-158 are involved in ligand interactions with a frequency of 47 % and 44 %, respectively [26].

Selectivity is a well-accepted parameter of the safety and pharmacological activity of a compound [28]. Bufadienolides from oocytes of R. alata were evaluated against kidney monkey Vero cells. The bufadienolides 1 (EC50 = 22.80 μ M, S.I. > 5) and 7 (EC50 = 3.93 μ M, S.I. = 2.7) presented significant selectivity (Table 2). Although bufadienolide 7 shows a promising S.I., less cytotoxicity is required. On the contrary, the low cytotoxicity showed by the bufadienolide 1 highlights its potential for further studies. Recently, 19-hydroxy-bufalin was evaluated against Vero cells and its low cytotoxicity, in comparison with bufadienolide 7, suggest that hydroxyl at position 19 decrease cytotoxic effects against mammals [21]. Despite of the potent anti-trypanosomal effect shown by compounds 3, 5 and 6, these bufadienolides were toxic to Vero cells, resulting in a reduced selectivity for the assayed parasite. In a previous study, hellebrigenin and telocinobufagin, two bufadienolides with antiprotozoal activity were tested for cytotoxicity on mouse macrophages and erythrocytes, showing no toxic effects [19]. However, it has been found that rodent cells are over 1000-fold more resistant than human cells to cytotoxic effects of cardiotonic sterols [29]. Differential cytotoxicity and water solubility are great attributes for potential anti-trypanosomals. In this context, microbial transformations carried out with bufadienolides 4, 5 and 7 have produced nontoxic derivatives to

human cells [30–32]. Nanoparticles containing bufadienolide 7 were injected in mice xenografted with human colon cancer cells, and their therapeutic potential was tested. The particles were able to inhibit the growth of tumor cells with efficient tumor targeting and cellular uptake [33].

Some bufadienolides have been found to be active against amphibian pathogens. Arenobufagin, γ -bufotalin and telocinobufagin isolated from the skin secretion of the boreal toad *Anaxyrus boreas* inhibit the harmful fungus Batrachochytrium dendrobatidis in vitro [34]. Amphibians are also affected by parasitic diseases caused by protozoa [35]. In fact, different families of anurans, including Bufonidae, are hosts of trypanosomes and are frequently parasitized by more than one species [36]. Blood samples from R. alata have not been examined for the presence of trypanosomes; however, congeneric species, as well as syntopic species such as the frog Engystomops pustulosus, are known to be parasitized by trypanosomes [37,38]. Trypanosomiasis in amphibians is characterized by anemia, food refusal, listlessness and localized hemorrhages with swollen lymph glands. Species of leeches from the family Hirudinea are considered to be the aquatic vectors of amphibian trypanosomes, while mosquitos, sand flies and midges, their main terrestrial vectors [37-39]. The presence of antitrypanosomal bufadienolides in oocytes of toads is intriguing. Bufonids biosynthesize bufadienolides, as shown by studies using cholesterol marked isotopically [40]. These compounds have been isolated from parotoid glands and skin of toads, as well as from their plasma, bile, ovaries and oocytes [17,41–43]. Because they inhibit (Na⁺-K⁺)ATPase pump, bufadienolides are considered as possible regulators of ionic equilibrium in anurans, and the high levels of bufadienolides present in some bufonid species presumably serve as defense against predation [44]. However, a possible role as endogenous antiprotozoal substances should not be discarded.

3. Materials and Methods

3.1. General Experimental Procedures

Optical activity was determined on a digital P-2000 polarimeter (Jasco, Easton, MD, USA) at room temperature. Ultraviolet spectra were registered using a UV-2401 PC recording spectrophotometer (Shimadzu, Columbia, MD, USA). Infrared spectra were measured using a Platinum ATR Alpha instrument (Bruker, Billerica, MA, USA). Reverse-phase high-performance liquid chromatography was carried out on an Agilent 1100 HPLC equipped with a Diode Array detector 1200 series (Agilent, Santa Clara, CA, USA). Nuclear magnetic resonance (NMR) experiments were carried out at 25 °C on an Eclipse+ 400 Fourier transform spectrometer with a field strength of 9.38 T and equipped with a 5 mm TH tunable probe (JEOL, Peabody, MA, USA). Chemical shifts (δ) were expressed in parts per million (ppm) and coupling constants (J) in Hz. NMR spectra were processed using the MestReNova software version 12.0.3-21384 (© Mestrelab Research S.L., Spain) and referenced based on the residual signal of chloroform-d (δ_H 7.26 and δ_C 77.00), dimethyl sulfoxide-d (δ_H 2.50 and δ_C 39.51) or methanol-d (δ_H 3.31 and δ_C 49.00). For High Resolution Mass Spectrometry analysis, compounds were diluted in methanol and mass spectra were acquired on either a micrOTOF-QIIITM spectrometer (Bruker Daltonics, Billerica, MA) or a MALDI-TOF-TOF Ultraflex (Bruker Daltonics, Billerica, MA, USA). For HR-ESI-Q-TOF analysis, compounds were directly infused into the spectrometer and MS spectra were obtained in positive electrospray ionization mode in the range of 50 to 2500 mass to charge ratio (m/z) and capillary voltage set at 4500 V. Nitrogen was used as the nebulizer gas at 2.0 bar, 200 °C and 9.0 L/min. External calibration was performed with Agilent ESI-L Low Concentration Tuning Mix (Agilent Technologies, Santa Clara, CA) prior and throughout data collection. Hexakis (1H, 1H, 2H-di-fluoro-ethoxy-phosphazene; m/z 622.0290 [M+H]+) (Synquest Laboratories, Alachua, FL) was used as internal standard. For HR-MALDI-TOF analysis, the samples were transferred to a MTP plate (Bruker Daltonics, Billerica, MA, USA), and mixed with the α -cyano-4-hydroxy-cinammic acid matrix (Sigma Aldrich). All experiments were carried out in positive mode. Pepcalibstandard was used as external calibrant (Bruker, Daltonics, Billerica, MA, USA).

3.2. Sample collection

Ninety grams of oocytes were obtained from six gravid females of *R. alata* collected at Parque Nacional Soberanía, Republic of Panama (9.134994° N 79.722597° W). These gravid females spontaneously laid their oocytes when held in isolation inside plastic containers. The toads were collected under the authorization of the Ministry of Environment of Panama (Permit: SE/AQ-2-14) and released at the same collection site. The Ministry of Environment of Panama reviewed and approved the experimental protocols of the study before granting collection permits. All methods were carried out in accordance with relevant guidelines and regulations of the INDICASAT. All methods are reported in accordance with ARRIVE guidelines.

3.3. Extraction and Isolation of bufadienolides

Freshly collected oocytes were macerated with 100 mL of chloroform-methanol (1:1) three times overnight in a shaker at room temperature. Extractions were pooled and dried under reduced pressure. The pool was suspended with deionized water and extracted with equal volume of chloroform-methanol (2:1). The organic phase was dried to provide 641.2 mg of residue. This extract was loaded onto Supelclean™ LC-18 SPE cartridges (Supelco, Bellefonte, PA, USA) and fractionated by employing a stepwise gradient of methanol in water at 40, 80 and finally to 100 % of methanol to produce fractions 1-3, respectively. Solvent was removed by rotary evaporation under vacuum and the remaining water was removed by lyophilization. Fraction 3 was diluted in methanol and the compounds were separated by HPLC. We used a semipreparative Synergy Hydro C18 column (250 x 10 mm, 4 μm, 80 Å) (Phenomenex, Torrance, CA, USA). The mobile phase consisted of water with 0.1 % of trifluoroacetic acid (TFA) (Solvent A), and a mixture of methanol-acetonitrile (1:1) acidified with TFA at 0.1 % (Solvent B). Elution detection was performed at λ 300 nm at a flow rate of 1 mL/min. The fractions were collected individually and as result the isolation of compounds 1 (5.7 mg, Rt 42.1 min), 2 (8.3 mg, Rt 63.0 min), 3 (1.7 mg, Rt 65.8 min), 4 (3.4 mg, Rt 67.5 min), 5 (4.4 mg, Rt 70.7 min), 6 (2.1 mg, Rt 75.4 min) and 7 (2.4 mg, Rt 77.1 min) was achieved. The purity of all compounds was found to be higher than 95 % as analyzed by ¹H NMR spectroscopy.

3.4. Compounds characterization data

 16β -hydroxyl-hellebrigenin (1): yellow solid; for 1 H and 13 C NMR data see supplementary information file; HR-ESI-MS m/z 433.2222 [M + H] $^{+}$ (calculated for C₂₄H₃₃O₇, 433.2221).

Desacetyl-bufotalin (2): yellow solid; For ^{1}H and ^{13}C -NMR data see supplementary information file. HR-ESI-MS m/z 403.2472 [M + H]+ (calculated for $C_{24}H_{35}O_{5}$, 403.2479).

19-formyl-dyscinobufotalin (3): white solid; $[\alpha]_D^{16}$ = + 11° (0.001 c, methanol); UV (CH₃OH) λ_{max} (log ϵ) 201 nm (4.45) 300 nm (3.90); IR ν_{max} 3589, 2864, 1715 cm⁻¹; for ¹H and ¹³C NMR data see Table 1 and spectra in supporting information (Figures S2–S7); HR-MALDI-MS m/z 487.2283 [M + H]⁺ (calculated for C₂₇H₃₅O₈, 487.2326) (Figure S8).

Bufotalin (4): white solid; For 1 H and 13 C-NMR data see supplementary information file. HR-ESI-MS m/z 445.2616 [M + H] $^{+}$ (calculated for C_{26} H₃₇O₆, 445.2585).

Cinobufotalin (5): yellow solid; For 1H and ^{13}C -NMR data see supplementary information file. HR-ESI-MS m/z 459.2385 [M + H] $^+$ (calculated for C₂₆H₃₅O₇, 459.2377).

Dyscinobufotalin (6): white solid; For ¹H and ¹³C-NMR data see supplementary information file. HR-ESI-MS m/z 473.2518 [M + H]⁺ (calculated for C₂₇H₃₇O₇, 473.2534).

Bufalin (7): white solid; For 1H and ^{13}C -NMR data see supplementary information file. HR-ESI-MS m/z 387.2512 [M + H] $^+$ (calculated for $C_{24}H_{35}O_4$, 387.2530).

3.5. Anti-trypanosomal activity

The evaluation of activity against trypanosomes was carried out using the recombinant strain Tulahuen, clone C4, of *T. cruzi* trypomastigotes (ATCC, Manassas, VA, USA). This strain expresses the β -galactosidase enzyme [45]. The parasites were maintained at 37 °C under a 5 % CO₂ atmosphere in RPMI-1640 culture medium with L-glutamine,4-(2-hydroxyethyl)-piperazine-1-ethanesulfonic

acid (HEPES) buffer, NaHCO3, 10 % FBS and 0.05 % gentamicin (50 mg/mL) as supplements. Epithelial kidney monkey Vero cells from the American Type Culture Collection (ATCC, Manassas, VA, USA) were cultured for 24 hours, and one day prior to the addition of the compound, cells were infected with trypomastigotes. After additional 24 hours of infection, bufadienolides (1-7) were dissolved in di-methyl-sulfoxide and tested at 10, 2, 0,4 and 0.08 μ g/mL for five days. Benznidazole was used as positive control. To determine the anti-trypanosomal activity, chlorophenol-red- β -D-galactopyranoside (Roche Applied Science) was added to each well and allowed to react with the β -galactosidase of the remaining living parasites for 4.5 hours. Absorbance was calculated at 570 nm using a microtiter plate reader (Sinergy HT, BioTek Instruments Inc., Winooski, VT, USA).

3.6. Mammalian cytotoxicity assay

Epithelial kidney monkey Vero cells were cultivated in 96-well plates at 37 °C under a 5% CO2 atmosphere, using RPMI-1640 medium (Sigma-Aldrich, USA) sterilized with 0.05 % gentamicin (50 mg/mL) and supplemented with 10 % FBS (fetal bovine serum; Gibco, Invitrogen, Carlsbad, CA, USA). Vero cells were allowed to adhere for 24 hours prior to incubation for five days with the samples. During the treatment, dimethyl-sulfoxide was used as negative control and doxorubicin as positive control. After incubation, 3-(4, 5-di-methyl-thiazol-2-yl)-2, 5-di-phenyl-tetra-zolium bromide was added to each well and absorbance was determined 4 hours later at 570 nm, using a color plate reader. Cytotoxicity was evaluated colorimetrically by calculating the ability of the remanent living Vero cells to reduce the pale yellow MTT into the black purple formazan product, as described [46].

3.7. Statistical analysis of bioassays

All anti-trypanosomal and cytotoxicity assays represent independent analysis and were performed in duplicate. The statistical analysis for the 50 % inhibiting (IC50) and cytotoxic (CC50) concentration values were carried out employing the Data Analysis complement Wizard of Excel 2000 by adjusting the dose-response curve to a sigmoidal model (Microsoft, Seattle, WA, USA).

3.8. Molecular docking

Molecular docking analysis was performed with the Molecular Operating Environment (MOE), software version 2018.01 (Chemical Computing Group, Montreal, QC, Canada). Molecular docking was carried out by allowing ligands to move keeping proteins rigid. Cruzipains and ligands were prepared with the standard protocol of MOE. The X-ray crystallographic data of twenty-four proteins were extracted from the Protein Data Bank. The structures of the proteins were corrected on its misallocated amino acid residues, protonated in 3D and the energies were minimized using the MMFF94x force field. Downloaded proteins were aligned before docking simulation. All cruzipains showed protein similarity from 99.1 to 100 %.

Validation of the protocol was carried out by redocking of the cruzipains 1EWO [47], 2AIM [48], 1U9Q [49] and 4XUI [50] (resolution between 2.10 - 2.51 Å); with their respective co-crystallized ligands VSC, ZRA, 186 and 2VC. Co-crystallized ligands were subjected to a simulated auto mode coupling induced fit. The binding pocket was defined as the set of amino acid residues within 4.5 Å from the co-crystallized ligand. Bufadienolides (1, 3, 7) were selected for molecular docking with cruzipain since they present the highest bioactivity and a significant selectivity in vitro. The docking conformations that showed lowest docking scores were selected for analysis.

4. Conclusions

Seven steroids from the bufadienolide class were isolated from the oocytes of the toad R. alata. Among them, 19-formyl-dyscinobufotalin (3) is reported as new natural product. While compound 2 was the only one reported as inactive, new bufadienolide 3 showed the most potent anti-trypanosomal activity, even more active than the control drug benznidazole. The *in vitro* assays and structural-activity relationship analysis suggest the formyl group at C-19 and esterification of the hydroxy group at C-16 increases anti-trypanosomal activity. 16β -hydroxyl-hellebrigenin showed

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significant selectivity and a promising therapeutic window. Furthermore, molecular docking analysis revealed bufadienolides **1**, **3** and **7** interacts with the main protease from *T. cruzi*, cruzipain, at the amino acid residues GLN-19, ASP-158, HIS-159 and TRP-177 through H-bonds. Chemical interactions involve the groups hydroxyl at positions C-5, and C-14 and formyl at C-19 at the cruzipain active site. Given the lack of therapeutic options to currently treat the American trypanosomiasis, this work can serve as the basis for further studies that aim the development of bufadienolides or their derivatives as drugs against Chagas disease.

Supplementary Materials: The following supporting information can be downloaded at the website of this paper posted on Preprints.org, Figure S1: RP-HPLC chromatogram profile of fraction-3 from oocytes of *Rhinella alata*; Figure S2: ¹H-NMR spetrum of 19-formyl-dyscinobufotalin (3); Figure S3: ¹³C-NMR spetrum of 19-formyl-dyscinobufotalin (3); Figure S5: HMBC spetrum of 19-formyl-dyscinobufotalin (3); Figure S6: HSQC spetrum of 19-formyl-dyscinobufotalin (3); Figure S7: COSY spetrum of 19-formyl-dyscinobufotalin (3); Figure S8: HR-MALDI-MS spetrum of 19-formyl-dyscinobufotalin (3); Figure S9: Alignment and superposition of 24 cruzipains and cocrystallized ligands; NMR data for compounds 1, 2, 4-7 isolated from the oocytes of *Rhinella alata*; Figure S10: ¹H NMR spectra for compound 16β-Hydroxyl-hellebrigenin (1); Figure S11: ¹H NMR spectra for compound Desacetyl-bufotalin (2); Figure S12: ¹H NMR spectra for compound Cinobufotalin (5); Figure S14: ¹H NMR spectra for compound Dyscinobufotalin (6); Figure S15: ¹H NMR spectra for compound Bufotalin (7); Supplementary references.

Author Contributions: Conceptualization, M.G., and C.R.; methodology, M.G., C.R., C.S. A.A.D.-A.; software, D.O.; validation, M.G., C.R., D.O. and C.S.; formal analysis, M.G., C.R., M.N., D.O. and C.S.; investigation, C.R., M.G., C.S. M.N., R.I.; resources, M.G., C.S., R.I.; data curation, C.R., M.G. C.S., D.O.; writing—original draft preparation, C.R., M.G., A.A.D.-A., R.I., D.O.; writing—review and editing, M.G., C.S. A.A.D.-A., R.I., D.O.; visualization, C.R., M.G.,; supervision, M.G., C.S.; project administration, M.G.; funding acquisition, M.G., C.S., R.I. All authors have read and agreed to the published version of the manuscript.

Funding: This research was funded by the INDICASAT-BID Program (grant number IND-JAL-05). C.R., R.I., A.A.D.-A., C.S. and M.G. were supported by the Sistema Nacional de Investigación (SNI), Panama.

Institutional Review Board Statement: The toads were collected under the authorization of the Ministry of Environment of Panama (Permit: SE/AQ-2-14) and released at the same collection site. The Ministry of Environment of Panama reviewed and approved the experimental protocols of the study before granting collection permits. All methods were carried out in accordance with relevant guidelines and regulations of the INDICASAT. All methods are reported in accordance with ARRIVE guidelines.

Informed Consent Statement: Not applicable.

Data Availability Statement: All data are available in the main text or the Supplementary Materials.

Acknowledgments: CR was supported by a scholarship of the Instituto para la Formación y Aprovechamiento de Recursos Humanos (IFARHU), Panama and The Secretaría Nacional de Ciencia, Tecnología e Innovación (SENACYT), Panama (No. **270-2014-007**). We gratefully acknowledge the Government of Panama (MiAmbiente) for granting permission to collect the frogs used in this study.

Conflicts of Interest: The authors declare no conflict of interest.

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