



Citation: Klug DM, Diaz-Gonzalez R, Pérez-Moreno G, Ceballos-Pérez G, García-Hernández R, Gomez-Pérez V, et al. (2019) Evaluation of a class of isatinoids identified from a high-throughput screen of human kinase inhibitors as anti-Sleeping Sickness agents. PLoS Negl Trop Dis 13(2): e0007129. https://doi.org/10.1371/journal. pntd.0007129

Editor: Igor C. Almeida, University of Texas at El Paso, UNITED STATES

Received: October 2, 2018

Accepted: January 4, 2019

Published: February 8, 2019

Copyright: © 2019 Klug et al. This is an open access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Data Availability Statement: All relevant data are within the manuscript and its Supporting Information files.

Funding: The authors wish to acknowledge funding from the National Institute of Allergy and Infectious Diseases (MPP & MN: R01AI114685; MPP R21AI126296, R21AI127594; CRC: R21AI126296, R21AI133393. https://www.niaid.nih.gov/), the Spanish Ministerio de Economía, Industria y RESEARCH ARTICLE

Evaluation of a class of isatinoids identified from a high-throughput screen of human kinase inhibitors as anti-Sleeping Sickness agents

Dana M. Klug¹, Rosario Diaz-Gonzalez², Guiomar Pérez-Moreno², Gloria Ceballos-Pérez², Raquel García-Hernández², Veronica Gomez-Pérez², Luis Miguel Ruiz-Pérez², Domingo I. Rojas-Barros², Francisco Gamarro², Dolores González-Pacanowska², María S. Martínez-Martínez³, Pilar Manzano³, Lori Ferrins¹, Conor R. Caffrey⁴, Miguel Navarro², Michael P. Pollastri¹*

1 Northeastern University Department of Chemistry & Chemical Biology, Boston, MA, United States of America, 2 Instituto de Parasitología y Biomedicina "López-Neyra" Consejo Superior de Investigaciones Científicas (CSIC), Granada, Spain, 3 Tres Cantos Medicines Development Campus, DDW and CIB, GlaxoSmithKline, Tres Cantos, Spain, 4 Center for Discovery and Innovation in Parasitic Diseases, Skaggs School of Pharmacy and Pharmaceutical Sciences, University of California, San Diego, La Jolla, CA, United States of America

* m.pollastri@northeastern.edu

Abstract

New treatments are needed for neglected tropical diseases (NTDs) such as Human African trypanosomiasis (HAT), Chagas disease, and schistosomiasis. Through a whole organism high-throughput screening campaign, we previously identified 797 human kinase inhibitors that grouped into 59 structural clusters and showed activity against *T. brucei*, the causative agent of HAT. We herein report the results of further investigation of one of these clusters consisting of substituted isatin derivatives, focusing on establishing structure-activity and -property relationship scope. We also describe their *in vitro* absorption, distribution, metabolism, and excretion (ADME) properties. For one isatin, **NEU-4391**, which offered the best activity-property profile, pharmacokinetic parameters were measured in mice.

Author summary

Human African trypanosomiasis (HAT) is a parasitic disease prevalent in sub-Saharan Africa. Current treatments cause severe toxicity, are difficult to administer, and are susceptible to resistance. In order to quickly discover new leads for HAT drug discovery, we screened human kinase inhibitors against *Trypanosoma brucei*, the parasite that causes HAT, and discovered several hundred compounds that demonstrated antiparasitic activity. In this paper, we present the results of medicinal chemistry follow-up work on a group of compounds known as isatins.

Competitividad, (MN: SAF2015-71444-P; DG-P: SAF2016-79957-R. http://www.mineco.gob.es), Junta de Andalucía (FG: CTS-7282. https://www. juntadeandalucía.es) and Subdirección General de Redes y Centros de Investigación Cooperativa (RICET) (MN: RD16/0027/0019; DG-P: RD16/ 0027/0014. https://bit.ly/2OegOnX). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing interests: We have read the journal's policy and the authors of this manuscript have the following competing interests: Two of the co-authors (MSM-M, PM) are employed by GlaxoSmithKline. Data was provided, free of charge, by AstraZeneca.

Introduction

Human African trypanosomiasis (HAT), also known as sleeping sickness, is a parasitic disease endemic in 36 African countries. Along with 19 other indications, HAT is designated as a neglected tropical disease (NTD) by the World Health Organization [1, 2]. Although there were fewer than 3,000 reported cases in 2015, the current burden of HAT amounts to 390,100 disability-adjusted life years [2, 3]. Caused by two subspecies of the parasite *Trypanosoma brucei* (*T. b. gambiense* and *T. b. rhodesiense*), HAT proceeds in two stages following initial infection via the bite of a tsetse fly. In the first stage, the parasite is present in the blood and lymph systems of the patient and causes mild, flu-like symptoms. In the second stage, the parasite crosses the blood-brain barrier (BBB) into the central nervous system (CNS) and causes a variety of neurological and behavioral changes, including disrupted sleeping patterns [2]. HAT is 100% fatal if left untreated, and the available drugs are associated with safety concerns, susceptibility to resistance, and lack of efficacy against all *T. brucei* subspecies and stages of the disease [4]. Currently, there are two new compounds for HAT in clinical trials: fexinidazole and acoziborole [5, 6]. However, given the high failure rate of compounds in clinical trials [7], it is prudent to continue to search for compounds to fill the drug discovery pipeline for HAT.

It has been shown by others that *T. brucei* expresses essential kinases [8], and furthermore, by our group, that human kinase inhibitors can be successfully re-optimized against these parasites [9–11]. As part of a lead repurposing strategy [12], we tested over 40,000 human kinase inhibitors in a high-throughput screen (HTS) against *T. b. brucei* [13]. This initial screening set was narrowed to 797 compounds with *T. b. brucei* pEC₅₀ >6 and >100× selectivity over HepG2 cells. These final hits were then clustered based on structural similarity. We herein report the development of structure-activity and structure-property relationships (SAR and SPR) for one of these clusters.

The compounds **NEU-1183**, **NEU-1184**, and **NEU-1185** (Fig 1) are representatives of a cluster of isatinoids that were identified in our kinase-targeted HTS as inhibitors of *T. brucei* growth. Various measured and computed properties of this cluster are shown in Table 1, along with our targeted values for each property. In addition to physicochemical properties such as clogP (calculated partition coefficient) and topological polar surface area (TPSA), we also considered metrics such as lipophilic ligand efficiency (LLE) [14] and CNS multi-parameter optimization (CNS-MPO) scores [15] when evaluating compounds. Overall, the isatinoids had good to excellent physicochemical properties that made them an attractive starting point for further development. Their generally low clogP and high LLE values suggested that



NEU-1184





https://doi.org/10.1371/journal.pntd.0007129.g001

| | Targeted Value | Cluster Average | NEU-1183 | NEU-1184 | NEU-1185 |
|--------------------------------------|--|-----------------|----------|----------|----------|
| <i>T.b.b.</i> pEC ₅₀ | ≥7.0 | 7.0 | 7.6 | 7.1 | 6.4 |
| HepG2 pTC ₅₀ ^a | <5.0 | 4.3 | 5.0 | 4.0 | 4.0 |
| cLogP | <u>≤</u> 3 | 1.1 | 2.3 | 0.52 | 0.41 |
| LLE | ≥5 | 5.9 | 5.3 | 6.5 | 6.0 |
| TPSA (Å ²) | 40 <x≤90< th=""><th>127</th><th>114</th><th>127</th><th>140</th></x≤90<> | 127 | 114 | 127 | 140 |
| MWt (Da) | \leq 360 | 374 | 370 | 367 | 383 |
| CNS-MPO Score | <u>≥</u> 3 | 4.3 | 4.2 | 4.5 | 4.1 |
| Kinetic aq. solubility (µM) | >10 | nd | <5 | 7.7 | 18 |

Table 1. Targeted values, cluster average, and individual values for the physicochemical properties of interest of NEU-1183, NEU-1184 and NEU-1185. Data from original HTS [13]. *nd* = no data.

^aTC₅₀ = 50% toxic concentration.

https://doi.org/10.1371/journal.pntd.0007129.t001

expansion of the structure would be tolerated from a property standpoint if necessary, and their high CNS-MPO scores indicated a likelihood of brain penetration (necessary for treatment of stage 2 infection). We therefore looked at ways to improve the potency and aqueous solubility of these compounds while maintaining their desirable physicochemical profile.

Methods

Bioactivity assays

In order to determine the *T. b. brucei* EC_{50} values, 4 µL per well from compound master plates were dispensed into a new plate and 96 µL of HMI-9 per well were added to generate a 4% DMSO intermediate plate. Mid-log phase growth *T. b. brucei* was diluted to a working cell density of 2,750 cells/mL and 90 µL/well dispensed into 96-well flat-bottom transparent assay plates (Nunc). Ten µL/well from intermediate plates were added. The final top concentration of compounds was 40 µM in 0.4% DMSO per well.

Assay plates were incubated for 72 h at 37°C and 5% CO₂. Four hours prior to the end of the incubation, 20 μ L of a 440 μ M resazurin solution in prewarmed HMI-9 was added to each well and incubated for another 4 h. Fluorescence was then measured in an Infinite F200 plate reader (Tecan) at 550 nm (excitation filter) and 590 nm (emission filter). A 4-parameter equation was employed to fit the dose-response curves and determine of EC₅₀ using the SigmaPlot 13.0 software. Assays were performed in duplicate at least twice, to achieve a minimal n = 3 per dose response.

Detailed protocols for rate of action assays, *Trypanosoma cruzi* and *Leishmania donovani* EC₅₀ assays, MRC5 and THP-1 cytotoxicity assays, and *Schistosoma mansoni* assays are provided in **S2 Text**.

Pharmacokinetics protocols

NEU-4391 was administered intraperitoneally (IP) to two groups of female NMRI mice (Group 1 n = 3; Group 2 n = 6). The compound was prepared in 1% (v/v) DMSO:99% (v/v) 20% (w/v) sulfobutyl ether-beta-cyclodextrin (SBE- β -CD) (Captisol) in water and the dosing volume was 10 mL/kg for a total dose of 10 mg/kg. Food and tap water were available *ad libi-tum*. Following IP dosing, Group 1 blood samples were collected from the tail vein into capillary tubes containing K2EDTA at the following time-points: 0.0833, 0.25, 0.5, 1, 2, 4, 6, 8 and 24 h.

In order to obtain simultaneous blood and brain samples, Group 2 mice were placed under terminal anaesthetic (isoflurane) and blood samples (0.3 mL) collected from the retro-orbital sinus into K2EDTA tubes at 0.5 h (n = 3) and 4 h (n = 3) after compound administration. Immediately following blood sample collection, death was confirmed by cervical dislocation and the brain removed. Aliquots of each blood sample were diluted in an equal volume of water. Mouse brain samples were weighed, water was added at a 1/2 (w/v) ratio (brain/water), and then homogenized. Both blood and brain samples were stored -80°C until analysis.

Diluted blood and brain homogenates were processed under standard liquid-liquid extraction procedures using acetonitrile containing an internal standard (Nifedipine) and analyzed by LC-MS/MS. Non-compartmental analysis was performed using the Phoenix pharmacokinetic software version 1.4 (Certara) and C_{max} , t_{max} , AUC_{las}t, AUC, and $t_{1/2}$ were estimated.

Ethics statement

All animal studies were ethically reviewed and carried out in accordance with Animals (Scientific Procedures) Act 1986 and the GSK Policy on the Care, Welfare and Treatment of Animals. This work was performed at Charles River Laboratories, Edinburgh Ltd. under the UK Home Office Project Procedure Project License No. PPL 70/8781: Drug Metabolism and Pharmacokinetics, Protocol Reference Number 1 and 6.

Chemistry

All sulfonamide-replacement analogs were synthesized by first preparing the requisite hydrazine from an aryl chloride or bromide, and subsequently coupling with isatin by stirring in methanol at room temperature. A mixture of isomers was isolated by vacuum filtration and further resolved by purification or recrystallization. All compounds tested had a purity of >95% as measured by LCMS. Details regarding compound syntheses and characterization are provided in <u>S1 Text</u>.

Results

In addition to the three compounds shown in Table 1, analogs with substituents at the C4, C5, and C6 (\mathbb{R}^3 , \mathbb{R}^2 , and \mathbb{R}^1 , respectively) were obtained from the GSK compound collection and tested against T. brucei; the biological activity of these compounds is presented in Table 2 (the general structure of analogs presented is shown in Fig 2). The most potent of these compounds was **NEU-5469**, with methyl groups at R^1 and R^3 and a hydroxy group at R^2 . Changing R^3 to a chloro, as in NEU-5485, resulted in a log unit drop in potency; further changing R¹ to an isopropyl group (NEU-5455) resulted in another drop in potency of over half a log unit. Further exploration of the R³ substituent showed that in general, small alkyl substituents were best in terms of both potency and LLE. The ethyl (NEU-5489), isopropyl (NEU-5491), isobutyl (NEU-5487), and isobutylene (NEU-1183) were approximately equipotent, although NEU-1183 has a slightly improved LLE due to its low clogP (2.2 versus NEU-5487's 3.4). Amides (NEU-5460) and larger substituents (NEU-5492 and NEU-5493) at this position generally resulted in less potent compounds as compared to the alkyl substituents. Analogs substituted at both R^2 and R^3 (NEU-5464) were significantly more potent than analogs substituted only at R^2 (NEU-2319). Comparing NEU-5464 to NEU-5479, the methyl substituent at R^2 results in a loss of potency of almost half a log unit as compared to the chlorine, showing that in this case, chlorine does not act as a bioisostere for a methyl group. Aromatic fused ring systems, such as those of NEU-1184 and NEU-5456, were more potent than the aliphatic ring of NEU-5459. As a general measure of mammalian cell toxicity, compounds were tested against either

| | R ¹ | R ² | R ³ | T.b.b. pEC ₅₀ | T.b.b. LLE | pTC ₅₀ |
|-----------|------------------|------------------|--|--------------------------|-------------------|-------------------|
| NEU-1183* | -H | -H | -CHC(CH ₃) ₂ | 7.5 | 5.3 | 5.0 ^a |
| NEU-1185* | -H | 5-oxazole | -H | 6.4 | 6.0 | <4.0 ^a |
| NEU-2319 | -H | -Cl | -H | 6.0 | 3.7 | <5.0 ^b |
| NEU-5455 | -iPr | -OH | -Cl | 6.4 | 3.2 | <5.0 ^b |
| NEU-5460 | -H | -H | -CONH ₂ | 6.2 | 5.7 | <5.0 ^b |
| NEU-5464 | -H | -Cl | -CH ₃ | 7.3 | 5.2 | <5.0 ^b |
| NEU-5469 | -CH ₃ | -OH | -CH3 | 8.2 | 5.8 | <5.0 ^b |
| NEU-5470 | -Et | -H | -H | 6.1 | 3.9 | <5.0 ^b |
| NEU-5479 | -H | -CH ₃ | -CH ₃ | 6.9 | 4.2 | <5.0 ^b |
| NEU-5485 | -CH ₃ | -OH | -Cl | 7.1 | 4.6 | <5.0 ^b |
| NEU-5487 | -H | -H | -CH ₂ CH(CH ₃) ₂ | 7.4 | 4.0 | <5.0 ^b |
| NEU-5489 | -H | -H | -Et | 7.5 | 4.9 | <5.0 ^b |
| NEU-5491 | -H | -H | -iPr | 7.6 | 4.7 | <5.0 ^b |
| NEU-5492 | -H | -H | -CH ₂ cyBu | 6.9 | 3.6 | <5.0 ^b |
| NEU-5493 | -H | -H | -CH2CH2-4-hydroxybenzene | 6.0 | 2.1 | <5.1 ^b |
| NEU-1184* | See Fig 2 | | | 7.1 | 6.4 | <4.0 ^a |
| NEU-5456 | See Fig 2 | | 7.4 | 5.5 | <5.0 ^b | |
| NEU-5459 | See Fig 2 | | 6.1 | 4.9 | <5.0 ^b | |

Table 2. Potency, LLE, and toxicity data for substituted-core analogs.

^aHepG2 toxicity

^bMRC5 toxicity.

*Data from original HTS [13]. All experimental error was within ±0.20 log units.

https://doi.org/10.1371/journal.pntd.0007129.t002

HepG2 (original hits) or MRC5 (follow-up compounds) cell lines; no analogs showed significant toxicity against either cell type.

Given the established SAR around the isatin core, we decided to focus further efforts on replacing the sulfonamide moiety. Because of easier synthetic accessibility, initial analogs were synthesized without any substituents on the core by first making the requisite hydrazine **2** from the aryl halide **1** (X = Br or Cl) and coupling with isatin as shown in Fig **3**. The double bond geometry of **3** was confirmed by X-ray crystallography (S1 Fig; CCDC ID 1865900) to be the (Z) isomer as shown in Fig **3**, where an intramolecular hydrogen bond is formed between







NH₂

NEU-5456



Fig 2. General structure of analogs presented in Table 2 as well as structures of NEU-1184, -5456, and -5459.

https://doi.org/10.1371/journal.pntd.0007129.g002





Fig 3. General scheme for synthesis of sulfonamide replacements and construction of substituted isatin core. Reagents and reaction conditions: a) Aryl halide, hydrazine monohydrate; 120°C, 12 h (52–86%). b) Isatin, methanol; RT, 12 h (10–100%). c) 4-Bromoisatin, boronic acid pinacol ester, K_2CO_3 , Pd(PPh₃)₄; 100°C, 12 h (66%).

https://doi.org/10.1371/journal.pntd.0007129.g003

the amide carbonyl and the hydrazone -NH. Later, analogs with an isobutylene substituent at R^3 were also synthesized; this core was constructed via a Suzuki coupling with 4-bromoisatin 4 and boronic ester 5 as shown in Fig 3.

Additionally, we noted the structural similarity of this chemotype to hesperadin, an investigational human Aurora kinase inhibitor that we previously demonstrated as active against *T*. *brucei* (Fig 4) [16]. With this in mind, NEU-4893 was designed as a crossover analog. Synthesis, shown in Fig 4, began with a reductive amination between 4-chlorobenzaldehyde 7 and piperidine, followed by displacement of the chloride under Buchwald conditions with bochydrazine to yield 9. This compound was then deprotected using HCl to yield the HCl salt 10, which was coupled with isatin to yield the final compound NEU-4893.

The biological activities of the sulfonamide replacement analogs are shown in Table 3 (the general structure of analogs presented is shown in Fig 5). The removal of the olefin at R³ resulted in the loss of nearly two log units of potency against *T. brucei*, illustrated by NEU-2115. Replacement of the sulfonamide invariably resulted in a loss of potency as compared to the unsubstituted primary sulfonamide NEU-2115, demonstrating that this moiety is critical for activity. Tuning the electronics of the aryl ring at R⁴ had a modest effect on potency; where the electron-rich NEU-2117 lost 0.5 log units of activity compared to these two analogs. Alkylation of the primary sulfonamide resulted in a similar loss of activity, although secondary (NEU-2124) and tertiary (NEU-2118) sulfonamides were better tolerated than the excision of the sulfonamide moiety altogether. Re-introduction of the olefin substituent at R³ (as in NEU-4391 and NEU-4405) predictably resulted in improved potency in the case of NEU-4391, although this came at the cost of increased clogP. Incorporation of the benzyl piperidine moiety of NEU-4893 led to decreased activity.



Fig 4. *T. b. brucei* activity of hesperadin and synthetic scheme for synthesis of NEU-4893. Reagents and reaction conditions: a) Piperidine, TEA, AcOH, NaHB(OAc)₃, DCM; RT, 12 h (85%). b) Boc-NH-NH₂, NaOtBu, Pd₂(dba)₃, XPhos, dioxane; μw, 150°C, 2 h (74%). c) 4M HCl in dioxane; RT, 3 h (74%). d) NaOtBu, isatin, MeOH; RT-50°C, 48 h (28%).

https://doi.org/10.1371/journal.pntd.0007129.g004

The absorption, distribution, metabolism, and excretion (ADME) properties of the sulfonamide replacement analogs were also assessed (<u>S1 Table</u>). In general, the aqueous solubility of these compounds is low, with only two (**NEU-2124** and **NEU-4893**) achieving a solubility

Table 3. Potency, LLE, and toxicity data for sulfonamide replacement analogs.

| | R ³ | R ⁴ | <i>T.b.b.</i> pEC ₅₀ | T.b.b. LLE | MRC5 pTC ₅₀ |
|-----------|-------------------------------------|------------------------------------|---------------------------------|------------|------------------------|
| NEU-1183* | -CHC(CH ₃) ₂ | 4-SO ₂ NH ₂ | 7.6 | 5.3 | <4.3 |
| NEU-2114 | Н | 4-OCH ₃ | 5.0 | 2.1 | <4.3 |
| NEU-2115 | Н | 4-SO ₂ NH ₂ | 5.9 | 4.2 | <4.3 |
| NEU-2116 | Н | n/a | 4.9 | 1.8 | <4.3 |
| NEU-2117 | Н | 3,5-dichloro | <4.4 | 0.12 | <4.3 |
| NEU-2118 | Н | 4-SO ₂ NMe ₂ | 5.8 | 3.6 | <4.3 |
| NEU-2124 | Н | 4-SO ₂ NHMe | 5.3 | 3.4 | <4.3 |
| NEU-4391 | -CHC(CH ₃) ₂ | 4-SO ₂ NHMe | 6.8 | 3.5 | <4.3 |
| NEU-4405 | -CHC(CH ₃) ₂ | 4-SO ₂ NMe ₂ | 5.3 | 1.8 | <4.3 |
| NEU-4893 | Н | 4-CH ₂ -piperidine | 5.0 | 1.5 | <4.3 |

*Data from original HTS [13]. All experimental error within ±0.08 log units.

https://doi.org/10.1371/journal.pntd.0007129.t003



Fig 5. General structure of analogs presented in Table 3.

https://doi.org/10.1371/journal.pntd.0007129.g005

above 10 μ M. Apart from **NEU-2116**, human liver microsome (HLM) clearance is low for this cluster. However, rat hepatocyte clearance shows significantly higher variability across the series, suggesting that mechanisms other than metabolism by CYPs may be involved in clearance. Table 4 shows the ADME profile of two compounds of interest, **NEU-4893** and **NEU-4391**. **NEU-4893** shows the best overall ADME profile of the sulfonamide replacement analogs, but was inactive against *T. brucei* and for this reason was not pursued further. **NEU-4391** showed sub-micromolar activity against *T. brucei* and had a reasonable ADME profile, aside from high rat hepatocyte clearance.

Of the active compounds for which ADME data was available, **NEU-4391** displayed the best combination of potency and favorable ADME properties. We therefore progressed this compound to mouse pharmacokinetic (PK) studies using an IP dose of 10 mg/kg (Fig 6). The concentration of **NEU-4391** in blood is represented in Fig 6A, with a C_{max} of 50.1 ng/ml

| | Targeted Value | NEU-4893 | NEU-4391 |
|---|----------------|----------|----------|
| clogP | ≤3 | 3.5 | 3.3 |
| LogD (7.4) | ≤2 | 2.2 | nd |
| Aq sol (μM) | >10 | 26 | 4 |
| HLM Cl _{int} (μL/min/mg) | <9 | <3 | 8.6 |
| Rat Hepatocyte Cl _{int} (μL/min/10 ⁶ cells) | <5 | 148 | 103 |
| DDB (%) | < 95 | 90 | >96 |

Table 4. ADME profile of NEU-4893.

Values that meet the target are shaded green, those in an intermediate range are shaded yellow, and those that are well outside the target value are shaded red.

https://doi.org/10.1371/journal.pntd.0007129.t004



Fig 6. a) Peripheral blood levels of NEU-4391 after IP administration at 10 mg/kg. b) Blood and brain levels at 0. 0.5, and 4 h. Female NMRI mice (n = 3). 1% DMSO:99%, 20% Captisol in water. Individual values for each time point are represented in the plot. Concentrations after 1 h in **Fig 6A** were below the lower limit of quantitation.

https://doi.org/10.1371/journal.pntd.0007129.g006

 $(0.130 \ \mu\text{M})$ (S2 Table). This concentration is within two-fold of the EC₅₀ of NEU-4391 (0.084 μ M) and is unlikely to produce a therapeutic effect *in vivo*. Additionally, the blood and brain concentrations of NEU-4391 were measured in a separate experiment (Fig 6B). These results show that the concentration of NEU-4391 in the brain was below the limit of detection for all time points (S3 Table). The combined PK results clearly indicate that NEU-4391 is unlikely to be an effective therapeutic for either stage 1 or stage 2 HAT.

Discussion

The modest potency and suboptimal ADME-PK properties suggest that this cluster of isatins is unsuitable for further development as anti-HAT agents. Although we discovered a compound, **NEU-4893**, with an improved ADME profile over the original HTS hits, this compound lost significant activity against *T. brucei* (approximately 2.5 log units as compared to **NEU-1183**). The compound with the best combination of potency and ADME parameters, **NEU-4391**, was progressed to *in vivo* PK studies, but both blood and brain concentrations were too low to warrant further progression.

Given our success with parasite cross-screening campaigns in the past [17], a selection of the compounds synthesized were also tested against the related parasites *T. cruzi*, the causative agent of Chagas disease and *L. donovani*, one of the causative agents of visceral leishmaniasis (S3 Table). We have also previously observed activity against *S. mansoni*, one of the causative agents of schistosomiasis, in compounds derived from other HTS clusters; we therefore tested a selection of isatin derivatives against *S. mansoni* (S5 Table). These compounds did not show activity against either *T. cruzi* or *L. donovani*, apart from the weak activity of NEU-4391 and NEU-1183 against *T. cruzi* ($pEC_{50} = 5.65$ and 6.14, respectively). NEU-1183, -2118, and -2124 showed moderate and variable activity against *S. mansoni* adults and/or post-infective larvae (somules), but based on the overall results this cluster did not warrant further investigation as anti-schistosomal agents. We report these results in the interest of informing others who may be working on optimization programs based on our initially-reported HTS results.

Supporting information

S1 Table. ADME properties of selected analogs. (DOCX)

S2 Table. Individual blood pharmacokinetic parameters of NEU-4391. (DOCX)

S3 Table. Blood and brain levels of NEU-4391 after intraperitoneal administration of 10 mg/kg single dose.

(DOCX)

S4 Table. *T. cruzi* and *L. donovani* activity for all analogs in the isatinoid cluster. (DOCX)

S5 Table. *S. mansoni* activity of selected analogs. (DOCX)

S1 Fig. Crystal structure of NEU-2114. (TIF)

S2 Fig. Rate of action curve for NEU-4391. (TIF)

S1 Text. Detailed chemical synthesis and characterization. (DOCX)

S2 Text. Cell assay protocols. (DOCX)

S3 Text. ADME experiment protocols. (DOCX)

Acknowledgments

We thank AstraZeneca and GlaxoSmithKline for the provision of some of the *in vitro* ADME and physicochemical properties data, and Dr. Bo Li for the X-ray crystallography.

Author Contributions

Conceptualization: Dana M. Klug, Rosario Diaz-Gonzalez, Pilar Manzano, Miguel Navarro, Michael P. Pollastri.

Formal analysis: Dana M. Klug.

- Funding acquisition: Francisco Gamarro, Dolores González-Pacanowska, Miguel Navarro, Michael P. Pollastri.
- Investigation: Dana M. Klug, Rosario Diaz-Gonzalez, Guiomar Pérez-Moreno, Gloria Ceballos-Pérez, Raquel García-Hernández, Veronica Gomez-Pérez, Luis Miguel Ruiz-Pérez, Domingo I. Rojas-Barros, Francisco Gamarro, Dolores González-Pacanowska, María S. Martínez-Martínez, Pilar Manzano, Conor R. Caffrey.

Methodology: Rosario Diaz-Gonzalez.

- **Project administration:** Francisco Gamarro, Dolores González-Pacanowska, Lori Ferrins, Miguel Navarro, Michael P. Pollastri.
- Supervision: Rosario Diaz-Gonzalez, Pilar Manzano, Conor R. Caffrey, Miguel Navarro, Michael P. Pollastri.

Validation: Rosario Diaz-Gonzalez, Guiomar Pérez-Moreno, Gloria Ceballos-Pérez, Raquel García-Hernández, Veronica Gomez-Pérez, Luis Miguel Ruiz-Pérez, Domingo I. Rojas-Barros, Francisco Gamarro, Dolores González-Pacanowska, María S. Martínez-Martínez, Pilar Manzano, Conor R. Caffrey.

Writing - original draft: Dana M. Klug.

Writing - review & editing: Dana M. Klug, Lori Ferrins, Michael P. Pollastri.

References

- 1. World Health Organization. Neglected Tropical Diseases [November 24, 2015]. Available from: http://www.who.int/neglected_diseases/diseases/en/.
- 2. Drugs for Neglected Diseases Initiative. About Sleeping Sickness [September 1, 2016]. Available from: http://www.dndi.org/diseases-projects/hat/.
- Murray CJL, Barber RM, Foreman KJ, Ozgoren AA, Abd-Allah F, Abera SF, et al. Global, regional, and national disability-adjusted life years (DALYs) for 306 diseases and injuries and healthy life expectancy (HALE) for 188 countries, 1990–2013: quantifying the epidemiological transition. The Lancet. 2015; 386 (10009): 2145–91. http://dx.doi.org/10.1016/S0140-6736(15)61340-X.
- World Health Organization. Human African Trypanosomiasis Fact Sheet [November 24, 2015]. Available from: http://www.who.int/mediacentre/factsheets/fs259/en/.
- Drugs for Neglected Diseases Initiative. Fexinidazole (HAT) [February 13, 2018]. Available from: <u>http://www.dndi.org/diseases-projects/portfolio/fexinidazole/</u>.
- 6. Drugs for Neglected Diseases Initiative. Acoziborole [February 13, 2017]. Available from: http://www. dndi.org/diseases-projects/portfolio/scyx-7158/.
- Hay M, Thomas DW, L. CJ, Economides C, Rosenthal J. Clinical development success rates for investigational drugs. Nat Biotechnol. 2014; 32(1): 40–51. https://doi.org/10.1038/nbt.2786 PMID: 24406927
- Merritt C, Silva LE, Tanner AL, Stuart K, Pollastri MP. Kinases as druggable targets in trypanosomatid protozoan parasites. Chem Rev. 2014; 114: 11280–304. <u>https://doi.org/10.1021/cr500197d</u> PMID: 26443079
- Patel G, Karver CE, Behera R, Guyett PJ, Sullenberger C, Edwards P, et al. Kinase scaffold repurposing for neglected disease drug discovery: Discovery of an efficacious, lapatanib-derived lead compound for trypanosomiasis. J Med Chem. 2013; 56(10): 3820–32. <u>https://doi.org/10.1021/jm400349k</u> PMID: 23597080
- Ochiana SO, Pandarinath V, Wang Z, Kapoor R, Ondrechen MJ, Ruben L, et al. The human Aurora kinase inhibitor danusertib is a lead compound for anti-trypanosomal drug discovery via target repurposing. Eur J Med Chem. 2013; 62: 777–84. https://doi.org/10.1016/j.ejmech.2012.07.038 PMID: 22889561
- Seixas JD, Luengo-Arratta SA, Diaz R, Saldivia M, Rojas-Barros DI, Manzano P, et al. Establishment of a structure-activity relationship of 1*H*-imidazo[4,5-*c*]quinoline-based kinase inhibitor NVP-BEZ235 as a lead for African sleeping sickness. J Med Chem. 2014; 57(11): 4834–48. https://doi.org/10.1021/ jm500361r PMID: 24805946
- Klug DM, Gelb MH, Pollastri MP. Repurposing strategies for tropical disease drug discovery. Bioorg Med Chem Lett. 2016. https://doi.org/10.1016/j.bmcl.2016.03.103 PMID: 27080183
- Diaz R, Luengo-Arratta S, Seixas JoD, Amata E, Devine W, Cordon-Obras C, et al. Identification and characterization of hundreds of potent and selective inhibitors of *Trypanosoma brucei* growth from a kinase-targeted library screening campaign. PLoS Negl Trop Dis. 2014; 8(10): e3253. https://doi.org/ 10.1371/journal.pntd.0003253 PMID: 25340575
- Shultz MD. The thermodynamic basis for the use of lipophilic efficiency (LipE) in enthalpic optimizations. Bioorg Med Chem Lett. 2013; 23: 5992–6000. https://doi.org/10.1016/j.bmcl.2013.08.030 PMID: 24054120
- Wager T, Hou X, Verhoest PR, Villalobos A. Moving beyond rules: The development of a central nervous system multiparameter optimization (CNS MPO) approach to enable alignment of druglike properties. ACS Chem Neurosci. 2010; 1: 435–49. https://doi.org/10.1021/cn100008c PMID: 22778837
- Patel G, Roncal NE, Lee PJ, Leed SE, Erath J, Rodriguez A, et al. Repurposing human Aurora kinase inhibitors as leads for anti-protozoan drug discovery. Med Chem Commun. 2014; 5(5): 655–8. https:// doi.org/10.1039/c4md00045e PMID: 24910766

 Devine W, Woodring JL, Swaminathan U, Amata E, Patel G, Erath J, et al. Protozoan parasite growth inhibitors discovered by cross-screening yield potent scaffolds for lead discovery. J Med Chem. 2015; 58: 5522–37. https://doi.org/10.1021/acs.jmedchem.5b00515 PMID: 26087257