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MODEL-BASED SCREENING OF COMPOUNDS FOR THE TREATMENT OF CHAGAS DISEASE, A NEGLECTED TROPICAL DISEASE

S. D'Agate¹, I. Cotillo Torrejon², P. Healy¹, O. Della Pasqua^{1,3}

¹Clinical Pharmacology & Therapeutics Group, University College London, London, UK

²Kinetoplastid Discovery Performance Unit, GlaxoSmithKline, Tres Cantos, Spain

³Clinical Pharmacology Modelling & Simulation, GlaxoSmithKline, Uxbridge, UK

Chagas Disease

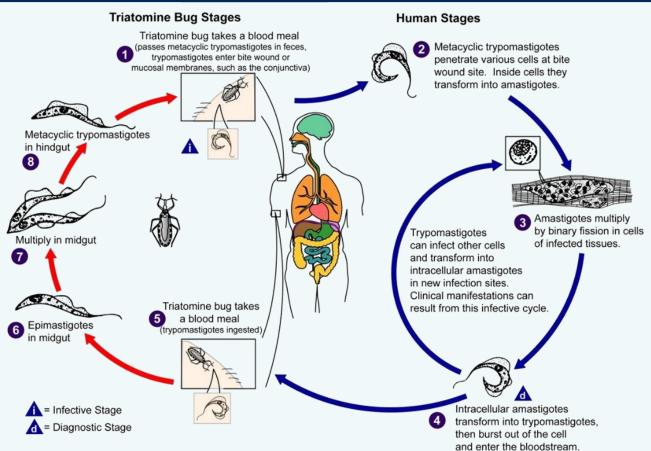


(American trypanosomiasis) people affected worldwide 0000 people die of complications linked to Chagas disease **Estimated number of cases** Status of vector transmission <1 000 Countries without vector transmission available 1 000-99 999 Countries with accidental vector transmission 100 000-999 999 drugs to Countries with ongoing vector transmission Data Source: World Health Organization World Health Organization >=1 000 000 Map Production: Control of Neglected treat Tropical Diseases (NTD) No officially estimated cases World Health Organization



Trypanosoma cruzi

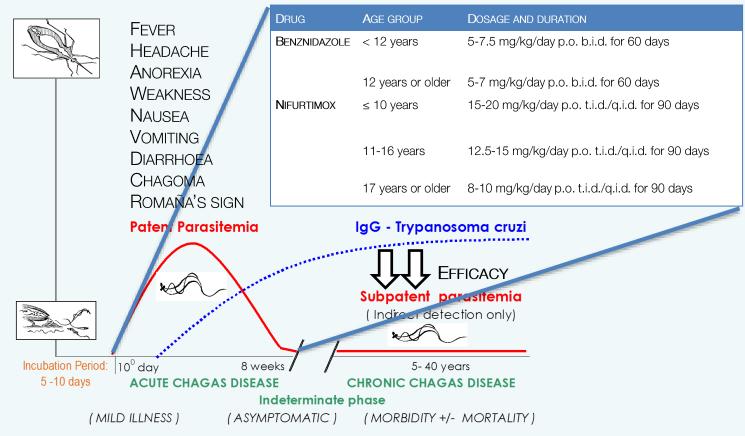






Chagas Disease



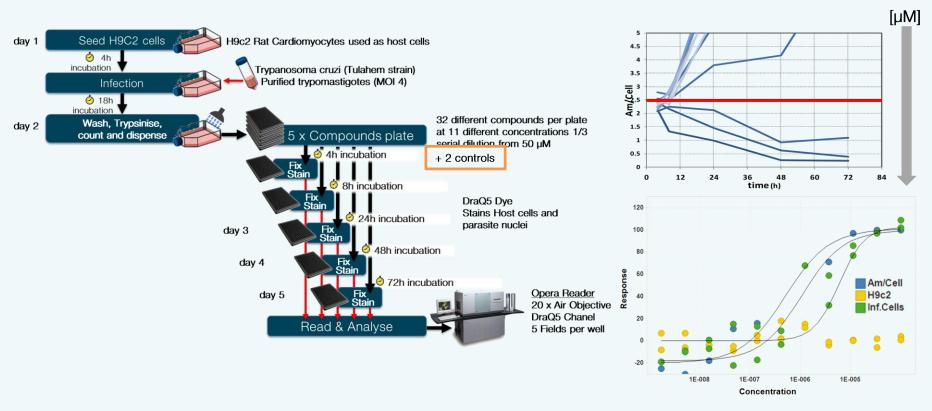




Courtesy of Dr Patricia Paredes, Guadalajara, Jal., Mexico.

Materials - experimental protocol[1]



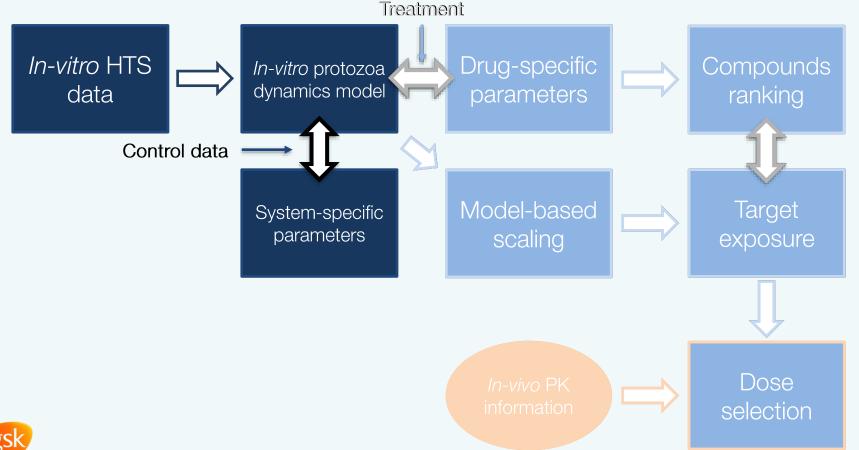




1. Alonso-Padilla et al. PLoS neglected tropical diseases 2015; 9: e0003493.

Aims of the current investigation







Methods - model development

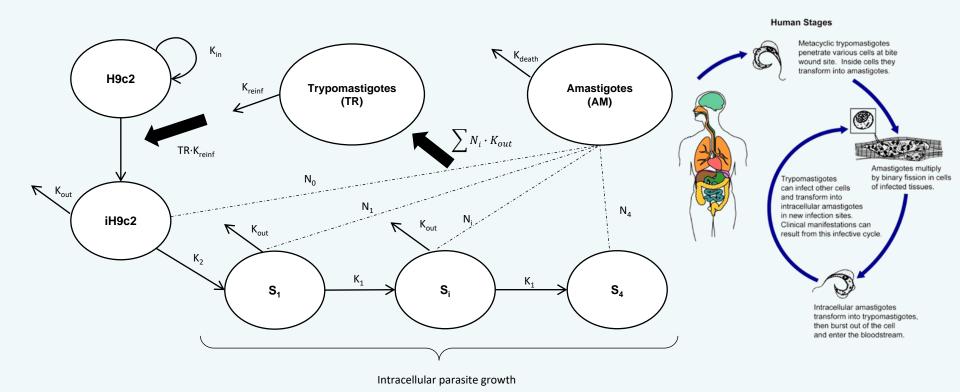


- Semi-mechanistic model parameterisation, aimed at disentangling drug-specific from system-specific parameters.
- Compartmental model describing protozoa growth and infection processes.
- Sequential PK/PD modelling approach:
 - 1. Control data used for the estimation of system-related parameters,
 - 2. Compound data used for the estimation of drug-related parameters.
- All analyses were carried out in NONMEM V7.3 using first-order conditional estimation with interaction (FOCE-I);
- Goodness of fit assessed by graphical methods, visual predictive check and bootstrap techniques.



Methods – Protozoa dynamics



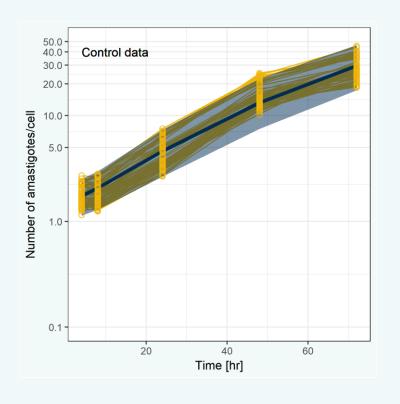




Visual predictive checks/Protozoa dynamics



Parameters of interest	Estimate
Reinfection rate (h^{-1})	0.0012
Premature host cell lyse (h^{-1})	0.0046
Initial number of amastigotes/cell	3.303
Parasite replication rate (h^{-1})	0.0716

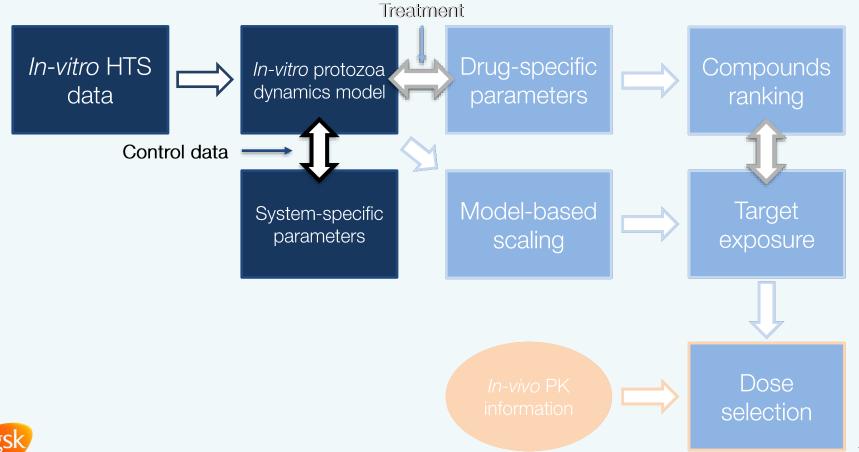




Data from control wells (N_{samples}=310) were used in the model building process.

Aims of the current investigation

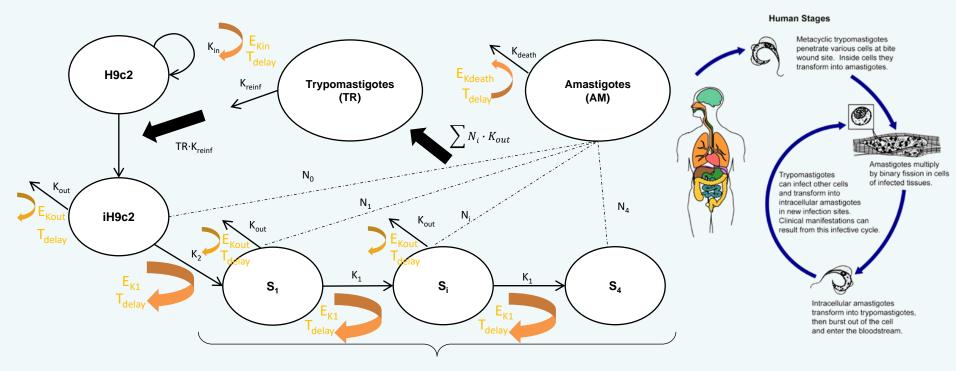






Methods - final model





Intracellular parasite growth



$$E = 1 - \frac{I_{max} \cdot C(t)}{IC_{50} + C(t)} \cdot D$$
 or $E = \frac{E_{max} \cdot C(t)}{EC_{50} + C(t)} \cdot D$ and $D = \frac{t}{T_{delay} + t}$

Drug-specific parameters



$$E = 1 - \frac{I_{max} \cdot C(t)}{IC_{50} + C(t)} \cdot D$$
 or $E = \frac{E_{max} \cdot C(t)}{EC_{50} + C(t)} \cdot D$ and $D = \frac{t}{T_{delay} + t}$

- IC_{50K1} inhibitory effect on the parasite replication rate
- EC_{50KD} \Longrightarrow direct effect on the death rate of the parasite
- IC_{50KIN} inhibitory effect on the host cells growth rate
- IC_{50KOUT} inhibitory effect on the premature lysis of host cells



Compound analysis



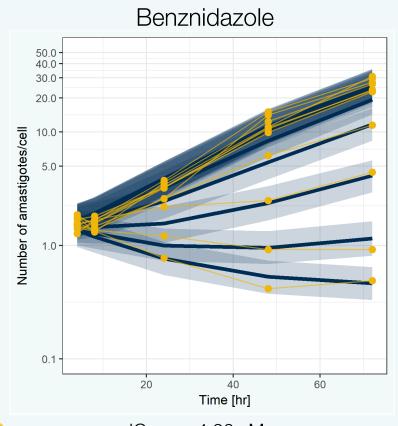
- Screening data:
 - 2 reference drugs (benznidazole, nifurtimox);
 - 44 new compounds

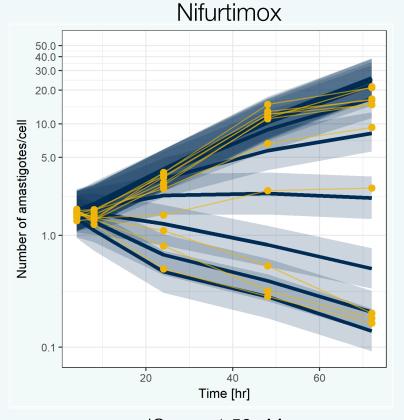
- Experimental protocol:
 - Exposure-response curves for each compound include 11 levels, ranging from 0.85 nM to 50 μM.



Visual predictive checks/Drug effect







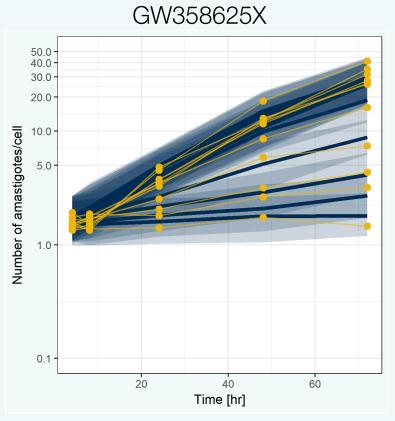


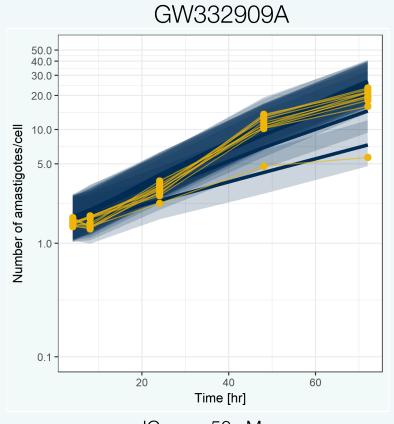
 $IC_{50K1} = 4.26 \,\mu\text{M}$ $EC_{50KD} = 14.73 \,\mu\text{M}$

 $IC_{50K1} = 1.59 \mu M$ $EC_{50KD} = 13.01 \mu M$

Visual predictive checks/Drug effect





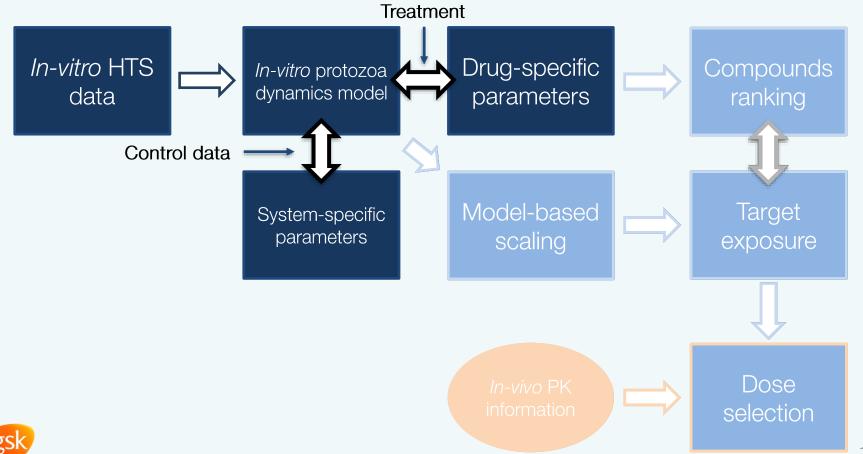




 $IC_{50K1} = 0.93 \, \mu M$

Aims of the current investigation





Compound ranking



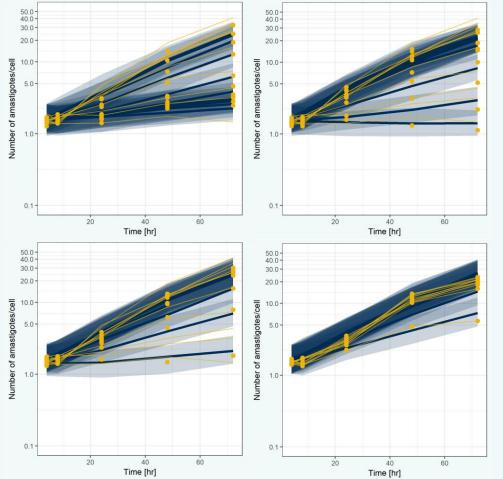
- Sensitivity analysis was performed to identify the most sensitive parameter describing parasiticidal and parasitostatic effects;
- IC_{50K1} represents the drug effect on parasite replication rate (parasitostatic);
- EC_{50KD} represents the drug induced increase in parasite clearance (parasiticidal).



Compound ranking



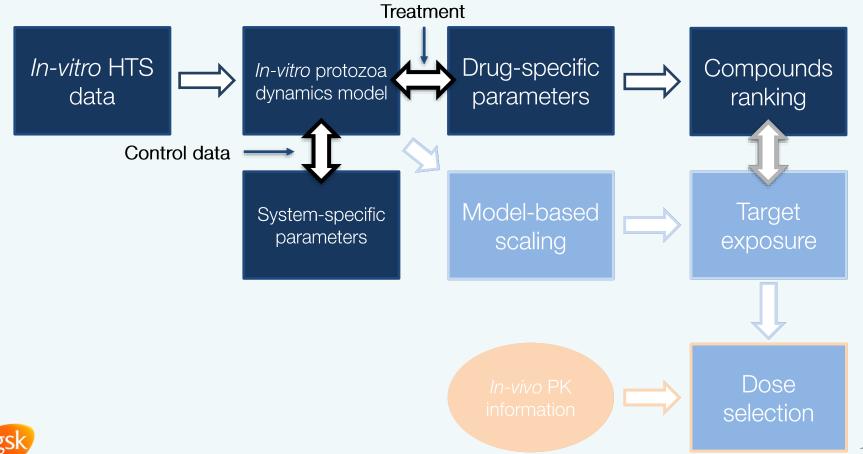
	Compound	IC _{50K1} (μΜ)
\Rightarrow	GW380944A	0.037
	GSK2249069A	0.066
	GSK503900A	0.071
	Posaconazole	1.26
	Nifurtimox	1.59
	GW368548X	1.61
	GW866668A	1.67
	GSK2960477A	1.71
	GSK1172530A	2.16
\Rightarrow	GW368763X	2.63
	Benznidazole	4.26
\Rightarrow	GSK121884A	16.96
\Rightarrow	GW332909A	>50





Aims of the current investigation





Dose rationale in humans

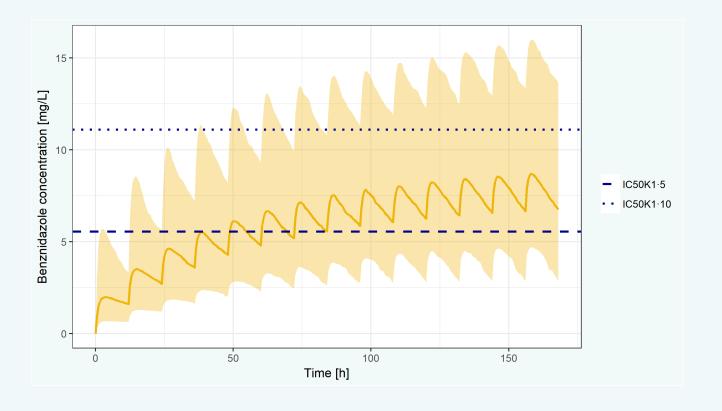


- Benznidazole and posaconazole were used to exemplify how model derived parameters can support the dose rationale for first-time-in humans and proof-of-concept studies.
- Target exposure corresponding to >IC80 and >IC90 values were assumed to be required for the therapeutic effect.
- Predicted exposure (plasma concentrations) were derived using allometric scaling principles and/or available pharmacokinetic data[2]. Results were then compared to currently used/tested clinical doses.



Target exposure - benznidazole

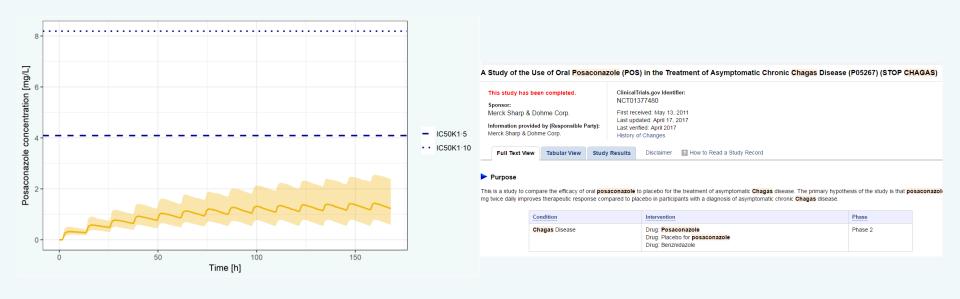






Target exposure - posaconazole







Conclusions



- We have shown that model-based approach can be used to describe parasite growth dynamics *in-vitro*;
- The model was parameterised using system and drug-specific parameters, enabling more precise estimates of the parameters of interest (IC_{50});
- Most compounds appear to have a predominant parasitostatic, rather than parasiticidal effect, which potentially explains the failure of some compounds in clinical trials;
- Given that target exposure can be derived from model parameters, our approach also allows a more robust dose rationale for proof of concept in Chagas disease.



Acknowledgements



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Nadia Terranova



References



[1] Benzekry S, Lamont C, Beheshti A, Tracz A, Ebos JM, Hlatky L, Hahnfeldt P. Classical mathematical models for description and prediction of experimental tumor growth. PLoS computational biology 2014; 10: e1003800.

[2] Nielsen EI, Friberg LE. Pharmacokinetic-pharmacodynamic modeling of antibacterial drugs. Pharmacol Rev 2013; 65: 1053-90.

[3] Alonso-Padilla, J., et al. (2015). "Automated high-content assay for compounds selectively toxic to Trypanosoma cruzi in a myoblastic cell line." PLoS Negl Trop Dis 9(1): e0003493.

[4] Soy, D., et al. (2015). "Population pharmacokinetics of benznidazole in adult patients with Chagas disease." Antimicrob Agents Chemother 59(6): 3342-3349.

[5] N. Terranova, M.B Jiménez-Díaz, I. Angulo-Barturen, P. Magni, O. Della Pasqua. Modelling of protozoa dynamics and drug effects in a murine model of malaria infection. PAGE. Abstracts of the Annual Meeting of the Population Approach Group in Europe. ISSN 1871-6032. PAGE 21 (2012) Abstr 2574.





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