

Structural basis of parasitic HSP90 ATP-ase inhibition by small molecules

Giusy Tassone¹, Marco Mazzorana^{2,*} and Cecilia Pozzi^{1,*}

¹ Department of Biotechnology, Chemistry and Pharmacy, Department of Excellence 2018–2022, University of Siena, Via Aldo Moro 2, 53100 Siena, Italy

² Diamond Light Source Ltd., Diamond House, Harwell Science & Innovation Campus, Didcot, Oxfordshire, OX11 0DE, UK.

* Correspondence: M.M. marco.mazzorana@diamond.ac.uk Tel.: +44 01235 778643; C.P. cecilia.pozzi@unisi.it; Tel.: +39 0577 232132

Table of contents

Figure S1.	S2
Table S1.	S3
References	S9

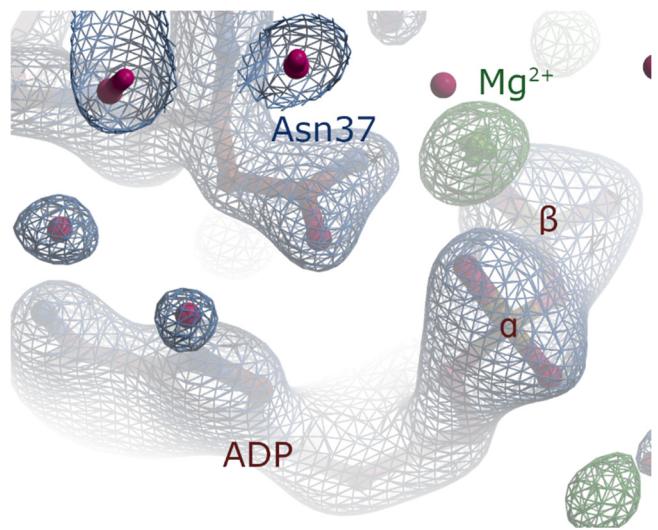
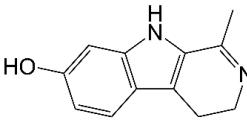
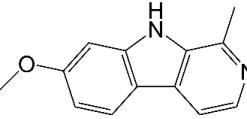
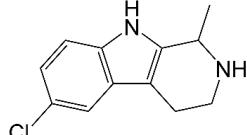
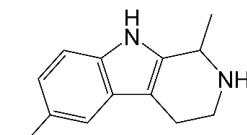
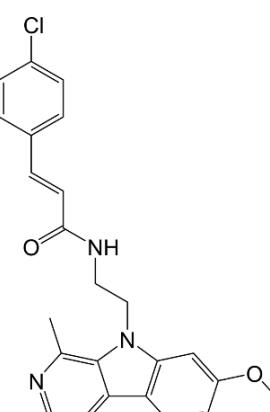
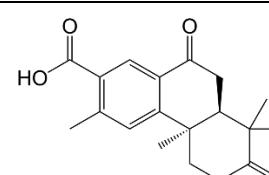
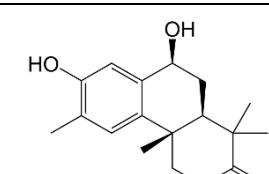
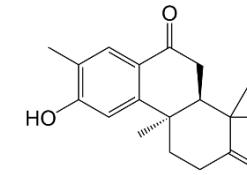


Figure S1. Electron density map detail from *Pf*-NTD structure (PDB id 3K60 [1]) reprocessed with PDB-redo (<https://pdb-redo.eu/db/3k60>) [2]. The $f_0 - f_c$ map (3σ level) shows a clear positive density not accounted for between the side chain of Asn37 and the α and β phosphates of ADP. The green blob, unaccounted for in the original structure, is compatible with the location of Mg(II) ions in other HSP90 structures.

Table S1. HSP90 binding and/or inhibition data and cytotoxicity on parasite and human cells by other natural products and synthetic compounds.

Compound	Structure	Parasite HSP90 binding and/or inhibition	Parasite cytotoxicity	Cytotoxicity on human cells
Harmine		<i>Pf</i> -NTD $K_d = 0.04$ mM [3]	N.A. [3]	N.A. [3]
Harmanol		<i>Pf</i> -NTD $K_d = 7.0$ mM [3]	N.A. [3]	N.A. [3]
17A		<i>Pf</i> $IC_{50} = 12.2 \pm 2.3$ μM [4]	<i>Pf</i> W2 $IC_{50} = 4.2 \pm 1.3$ μM [4]	HeLa $IC_{50} = 1.1 \pm 0.3$ mM [4] HepG2 $IC_{50} = 0.16 \pm 0.01$ mM [4]
21A		<i>Pf</i> $IC_{50} = 23.1 \pm 8.8$ μM [4]	<i>Pf</i> W2 $IC_{50} = 5.7 \pm 1.7$ μM [4] <i>Pf</i> MRA-1236 $IC_{50} = 9.2 \pm 0.4$ μM [4] <i>Pf</i> MRA-1240 $IC_{50} = 9.6 \pm 2.0$ μM [4]	HeLa $IC_{50} = 0.48 \pm 0.14$ mM [4] HepG2 $IC_{50} = 0.140 \pm 0.001$ mM [4]

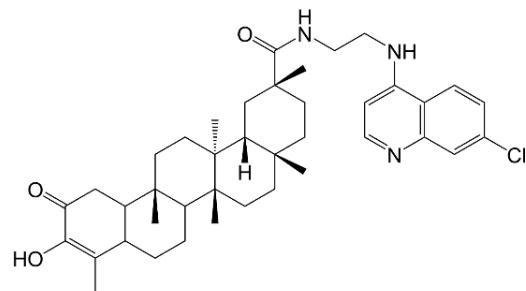
NATURAL COMPOUNDS

 <p>5e</p>	<p>N.A. [5]</p>	<p><i>Pf</i>3D7 $IC_{50} = 0.04 \pm 0.02 \mu\text{M}$ [5]</p> <p><i>Pf</i>Dd2 $IC_{50} = 0.17 \pm 0.01 \mu\text{M}$ [5]</p>	<p>HepG2 $IC_{50} = 2.91 \pm 1.75 \mu\text{M}$ [5]</p>
<p>Margolonone</p> 	<p>N.A. [6]</p>	<p>N.A. [6]</p>	<p>N.A. [6]</p>
<p>Nimbione</p> 	<p>N.A. [6]</p>	<p>N.A. [6]</p>	<p>N.A. [6]</p>
<p>Nimbinone</p> 	<p>N.A. [6]</p>	<p>N.A. [6]</p>	<p>N.A. [6]</p>

CE		<i>Lb</i> $IC_{50} = 210 \pm 10 \mu\text{M}$ [7]	<i>Li</i> Promatigotes $IC_{50} = 0.21 \pm 0.07 \mu\text{M}$ [7] <i>Lt</i> Promatigotes $IC_{50} = 0.49 \pm 0.17 \mu\text{M}$ [7]	HMEC-1 $IC_{50} = 0.38 \pm 0.08 \mu\text{M}$ [7] THP-1 $IC_{50} = 0.90 \pm 0.30 \mu\text{M}$ [7]
1		<i>Lb</i> $IC_{50} = 83 \pm 2 \mu\text{M}$ [7]	<i>Li</i> Promatigotes $IC_{50} = 0.06 \pm 0.02 \mu\text{M}$ [7] <i>Lt</i> Promatigotes $IC_{50} = 0.09 \pm 0.03 \mu\text{M}$ [7] <i>Lt</i> Amastogotes $IC_{50} = 0.19 \pm 0.11 \mu\text{M}$ [7]	HMEC-1 $IC_{50} = 0.51 \pm 0.08 \mu\text{M}$ [7] THP-1 $IC_{50} = 2.08 \pm 0.24 \mu\text{M}$ [7]
3		<i>Lb</i> $IC_{50} = 81 \pm 1 \mu\text{M}$ [7]	<i>Li</i> Promatigotes $IC_{50} = 0.06 \pm 0.01 \mu\text{M}$ [7] <i>Lt</i> Promatigotes $IC_{50} = 0.08 \pm 0.04 \mu\text{M}$ [7] <i>Lt</i> Amastogotes $IC_{50} = 0.13 \pm 0.09 \mu\text{M}$ [7]	HMEC-1 $IC_{50} = 0.53 \pm 0.17 \mu\text{M}$ [7] THP-1 $IC_{50} = 2.79 \pm 0.44 \mu\text{M}$ [7]
5		<i>Lb</i> $IC_{50} = 20 \pm 2 \mu\text{M}$ [7]	<i>Li</i> Promatigotes $IC_{50} = 0.12 \pm 0.02 \mu\text{M}$ [7] <i>Lt</i> Promatigotes $IC_{50} = 0.12 \pm 0.05 \mu\text{M}$ [7] <i>Lt</i> Amastogotes $IC_{50} = 0.52 \pm 0.28 \mu\text{M}$ [7]	HMEC-1 $IC_{50} = 0.12 \pm 0.03 \mu\text{M}$ [7] THP-1 $IC_{50} = 1.88 \pm 0.34 \mu\text{M}$ [7]

SYNTHETIC COMPOUNDS

6



Lb $IC_{50} = 65 \pm 4 \mu\text{M}$ [7]

Li Promatigotes $IC_{50} = 0.09$

$\pm 0.03 \mu\text{M}$ [7]

Lt Promatigotes $IC_{50} = 0.14$

$\pm 0.03 \mu\text{M}$ [7]

Lt Amastigotes $IC_{50} = 0.66$

$\pm 0.06 \mu\text{M}$ [7]

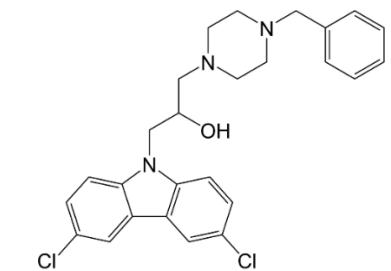
HMEC-1 $IC_{50} = 0.40 \pm 0.03$

μM [7]

THP-1 $IC_{50} = 0.85 \pm 0.24$

μM [7]

5B

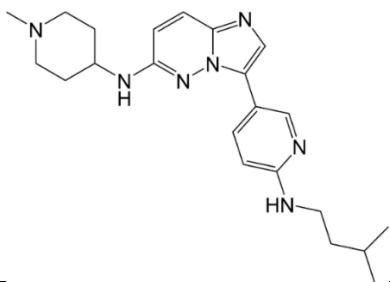


Pf-HSP90 $K_d = 28.1 \mu\text{M}$
[8]

Pf $IC_{50} = 82 \text{ nM}$ [8]

N.A. [8]

Compound D



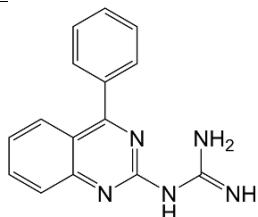
Pf-HSP90 $K_d = 6.17 \mu\text{M}$
[9]

Pf-HSP90-NTD $K_d = 10.73$
 μM [9]

N.A. [9]

N.A. [9]

Glb08



Lb $IC_{50} = 30 \pm 1 \mu\text{M}$ [10]

Lb-HSP90 $K_{Dapp} = 13 \pm 4$

μM [10]

Lb-NTD $K_{Dapp} = 17 \pm 6 \mu\text{M}$
[10]

Lb $IC_{50} = 43 \pm 2 \mu\text{M}$ [10]

N.A. [10]

Glb15		$Lb\ IC_{50} = 36 \pm 1\ \mu M$ [10] $Lb\text{-HSP90}\ K_{Dapp} = 14 \pm 4\ \mu M$ [10] $Lb\text{-NTD}\ K_{Dapp} = 13 \pm 3\ \mu M$ [10]	$Lb\ IC_{50} = 71 \pm 2\ \mu M$ [10]	N.A. [10]
Glb23		$Lb\ IC_{50} = 35 \pm 1\ \mu M$ [10] $Lb\text{-HSP90}\ K_{Dapp} = 26 \pm 7\ \mu M$ [10] $Lb\text{-NTD}\ K_{Dapp} = 20 \pm 10\ \mu M$ [10]	$Lb\ IC_{50} > 100\ \mu M$ [10]	N.A. [10]
PU-H71		$Pf\text{-NTD}\ K_d = 70.8 \pm 0.006\ \mu M$ [11]	$Pf\ IC_{50} = 111\ nM$ [11]	N.A. [11]
Compound 1		$Tc\text{-NTD}\ K_d = 9 \pm 2\ nM$ [12]	$Tc\ IC_{50} = 0.31\ \mu M$ [12]	N.A. [12]
Compound 3		$Tc\text{-NTD}\ K_d = 26 \pm 9\ nM$ [12]	$Tc\ IC_{50} = 0.21\ \mu M$ [12]	N.A. [12]

Compound 4		$Tc\text{-NTD } K_d = 3 \pm 1 \text{ nM}$ [12]	$Tc\text{ } IC_{50} = 0.10 \mu\text{M}$ [12]	N.A. [12]
D1U		N.A.	N.A.	N.A.

N.A. = Not Available

References

1. Corbett, K.D.; Berger, J.M. Structure of the ATP-Binding Domain of Plasmodium Falciparum Hsp90. *Proteins* **2010**, *78*, 2738–2744, doi:10.1002/prot.22799.
2. Joosten, R.P.; Long, F.; Murshudov, G.N.; Perrakis, A. The PDB_RED0 Server for Macromolecular Structure Model Optimization. *IUCrJ* **2014**, *1*, 213–220, doi:10.1107/S2052252514009324.
3. Shahinas, D.; Macmullin, G.; Benedict, C.; Crandall, I.; Pillai, D.R. Harmine Is a Potent Antimalarial Targeting Hsp90 and Synergizes with Chloroquine and Artemisinin. *Antimicrob Agents Chemother* **2012**, *56*, 4207–4213, doi:10.1128/AAC.00328-12.
4. Bayih, A.G.; Folefoc, A.; Mohon, A.N.; Eagon, S.; Anderson, M.; Pillai, D.R. In Vitro and in Vivo Anti-Malarial Activity of Novel Harmine-Analog Heat Shock Protein 90 Inhibitors: A Possible Partner for Artemisinin. *Malar J* **2016**, *15*, 579, doi:10.1186/s12936-016-1625-7.
5. Marinović, M.; Perković, I.; Fontinha, D.; Prudêncio, M.; Held, J.; Pessanha de Carvalho, L.; Tandarić, T.; Vianello, R.; Zorc, B.; Rajić, Z. Novel Harmicines with Improved Potency against Plasmodium. *Molecules* **2020**, *25*, doi:10.3390/molecules25194376.
6. Daniyan, M.O.; Ojo, O.T. In Silico Identification and Evaluation of Potential Interaction of Azadirachta Indica Phytochemicals with Plasmodium Falciparum Heat Shock Protein 90. *Journal of Molecular Graphics and Modelling* **2019**, *87*, 144–164, doi:10.1016/j.jmgm.2018.11.017.
7. Bassanini, I.; Parapini, S.; Ferrandi, E.E.; Gabriele, E.; Basilico, N.; Taramelli, D.; Sparatore, A. Design, Synthesis and In Vitro Investigation of Novel Basic Celastrol Carboxamides as Bio-Inspired Leishmanicidal Agents Endowed with Inhibitory Activity against Leishmania Hsp90. *Biomolecules* **2021**, *11*, doi:10.3390/biom11010056.
8. Wang, T.; Mäser, P.; Picard, D. Inhibition of Plasmodium Falciparum Hsp90 Contributes to the Antimalarial Activities of Aminoalcohol-Carbazoles. *J Med Chem* **2016**, *59*, 6344–6352, doi:10.1021/acs.jmedchem.6b00591.
9. Green, J.L.; Moon, R.W.; Whalley, D.; Bowyer, P.W.; Wallace, C.; Rochani, A.; Nageshan, R.K.; Howell, S.A.; Grainger, M.; Jones, H.M.; et al. Imidazopyridazine Inhibitors of Plasmodium Falciparum Calcium-Dependent Protein Kinase 1 Also Target Cyclic GMP-Dependent Protein Kinase and Heat Shock Protein 90 To Kill the Parasite at Different Stages of Intracellular Development. *Antimicrob Agents Chemother* **2015**, *60*, 1464–1475, doi:10.1128/AAC.01748-15.
10. Batista, F.A.H.; Ramos, S.L.; Tassone, G.; Leitão, A.; Montanari, C.A.; Botta, M.; Mori, M.; Borges, J.C. Discovery of Small Molecule Inhibitors of Leishmania braziliensis Hsp90 Chaperone. *J Enzyme Inhib Med Chem* **2020**, *35*, 639–649, doi:10.1080/14756366.2020.1726342.
11. Shahinas, D.; Folefoc, A.; Taldone, T.; Chiosis, G.; Crandall, I.; Pillai, D.R. A Purine Analog Synergizes with Chloroquine (CQ) by Targeting Plasmodium Falciparum Hsp90 (PfHsp90). *PLoS ONE* **2013**, *8*, e75446, doi:10.1371/journal.pone.0075446.
12. Pizarro, J.C.; Hills, T.; Senisterra, G.; Wernimont, A.K.; Mackenzie, C.; Norcross, N.R.; Ferguson, M.A.J.; Wyatt, P.G.; Gilbert, I.H.; Hui, R. Exploring the Trypanosoma Brucei Hsp83 Potential as a Target for Structure Guided Drug Design. *PLOS Neglected Tropical Diseases* **2013**, *7*, e2492, doi:10.1371/journal.pntd.0002492.