Supporting Information

Peptidic boronic acid *Plasmodium falciparum* SUB1 inhibitors with improved selectivity over human proteasome

Chrislaine Withers-Martinez,^{§a} Elina Lidumniece,^{§b} Fiona Hackett,^a Christine R Collins,^a Zahie Taha,^a Michael J. Blackman,^{*a,c} and Aigars Jirgensons^{*b}

^aMalaria Biochemistry Laboratory, The Francis Crick Institute, London NW1 1AT, United Kingdom;

^bLatvian Institute of Organic Synthesis, Riga LV-1006, Latvia;

^cFaculty of Infectious and Tropical Diseases, London School of Hygiene & Tropical Medicine, London WC1E 7HT, United Kingdom;

[§]These authors contributed equally

Content

. Effect of human proteasome inhibitors on activity of recombinant PfSUB1	S2
2. 20S Human Proteasome dose response assay for peptidic boronic acid SUB1 inhibitors	S2
3. Docking of bortezomib and inhibitors 1a, 3b, 1c and 4c into PfSUB1	S3
4. Compound 2a–h , 3a–j , 4a–c characterization spectra	S4
5. Representative examples of HPLC purity	S46

1. Effect of human proteasome inhibitors on activity of recombinant PfSUB1



Figure S1. Human proteasome inhibitors do not inhibit recombinant PfSUB1. In vitro PfSUB1 activity assay showing lack of inhibition by bortezomib, ixazomib or carfilzomib (1 uM final concentration). EP_1200 (used as a positive control) completely inhibited recombinant PfSUB1 activity at 100 nM. The assay shown is typical of duplicate assays.

2. 20S Human Proteasome dose response assay for peptidic boronic acid SUB1 inhibitors



Figure S2. 20S Human proteasome dose response assay. The peptidic boronic acid PfSUB1 inhibitors **1a**, **3a**, **3b**, the boralactone PfSUB1 inhibitors **1c**, **4c**, **4a** and the bortezomib control were used at 500 nM final in a fluorescence-based enzymatic assay measuring the chymotrypsin-like activity (β 5) of the human 20S proteasome. All measurements were performed in duplicate, with points shown as mean values. Error bars, S.D.

3. Docking of bortezomib and inhibitors 1a, 3b, 1c and 4c into PfSUB1



Figure S3. Bortezomib, compound **1a**, **3b**, **1c** and **4c** docked into PfSUB1(PDB: 4lvn). SUB1 is shown as a cartoon with a semi-transparent molecular surface colored by elements (O: red, N: blue). Positions P1 to P5 of inhibitors are indicated, the boron atom is colored in pink. Stabilizing H-bonds are shown as black dashed lines. **A**.: Bortezomib docked into the active site of PfSUB1 (ICM score -7) is shown as green sticks colored by elements. The P1 Leu side chain did not fill the S1 polar pocket and faced outwards the pocket. **B**.: compound **1a** docked into the active site PfSUB1 (ICM score -30) is shown as purple sticks colored by elements. The P1 Ala side chain fitted the S1 pocket. **C**: compound **3b** docked into the active site PfSUB1(ICM score -41) is shown as magenta sticks colored by elements. The P1 Ala side chain fitted the S1 pocket. The P5 phenyl pyridine capping group was stabilized by Pi-stacking interactions involving Leu466 and Arg468 (in wheat colour). **D**.: compound **1c** docked into the S1 polar pocket. E: compound **4c** docked into the active site of PfSUB1 (ICM score -28) is shown as yellow sticks colored by elements. The boralactone group fitted the S1 polar pocket.

4. Compound 2a-h, 3a-j, 4a-c characterization spectra





















































































5. Representative examples of HPLC purity



Reported by User: Olita Report Method: Default Individual Report Report Method ID 12125 Page: 1 of 1 Project Name: Martins Date Printed: 2024.04.26. 16:25:36 Europe/Riga



Project Name: Martins Date Printed: 2024.04.26. 16:25:05 Europe/Riga



Project Name: Martins Date Printed: 2024.04.26. 13:37:22 Europe/Riga



Project Name: Martins Date Printed: 2024.04.26. 13:36:48 Europe/Riga



Project Name: Martins Date Printed: 2024.04.26. 13:36:11 Europe/Riga



Project Name: Martins Date Printed: 2024.04.26. 13:28:39 Europe/Riga