Additional file 1 Additional compounds identified from virtual screening that were docked and tested *in vitro* against *Pf*SpdS.

Compound	Structure	Best Fit	Docking	%
		Value/number of PhFs	scores (kcal/mol)	Inhibition (at 100 μ M) ^a
DPM2				
3	NH ₃ ⁺ O NH ₃ ⁺ O NH ₂	0.91/4	-12.5	No Inhibition
4	*H ₃ N	0.15/4	-9.8	No Inhibition
DPM3				
5	HO NH ⁺ OH	0.10/5	-9.0	No Inhibition
6	*H ₃ N OH *HN	0.18/4	-9.7	No Inhibition
DPM4				
7	+H ₃ N O NH ₃ + NH NH NH NH ₃ +	0.14/5	-10.5	No Inhibition

^a Results represent inhibition of PfSpdS activity compared to untreated enzyme of three independent experiments performed in duplicate.