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Supporting Information

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Synthesis of new analogs of morpholine and their antiplasmodial evaluation against human malaria parasite *Plasmodium falciparum*

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Figure S1. ¹H NMR spectrum of 3.



Figure S2. ¹³C NMR spectrum of 3.



90 80 f1 (ppm)



0.000



Figure S6. ¹³C NMR spectrum of 6b.



Figure S8. ¹³C NMR spectrum of 6c.



Figure S9. ¹H NMR spectrum of 6d.



Figure S10. ¹³C NMR spectrum of 6d.



Figure S11. ¹H NMR spectrum of 6e.



Figure S12. ¹³C NMR spectrum of 6e.



Figure S13. ¹H NMR spectrum of 6f.



Figure S14. ¹³C NMR spectrum of 6f.



Figure S15. ¹H NMR spectrum of 6g.



Figure S16. ¹³C NMR spectrum of 6g.







re S18. ¹³C NMR spectrum of 6h.

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Figure S20. ¹³C NMR spectrum of 6i.



Figure S22. ¹³C NMR spectrum of 6j.



Figure S23. ¹H NMR spectrum of 6k.



Figure S24. ¹³C NMR spectrum of 6k.







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Figure S27. ¹³C NMR spectrum of 6m.



Figure S28. ¹³C NMR spectrum of 6m.

		Qua	litative Com	pound Report		
Data File Sample Type	N-12.d Sample Instrument 1 Damo JK.m Success		Sample Name Position	N-12 P1-B9		
nstrument Name org Method RM Callbration Status			User Name Acquired Time DA Method	29-08-2018 11:40:23 Default.m		
ample Group cquisition SW 6 ersion Q	200 series TOF TOF B.05.01 (I /6500 series (85125.1)	nfo.			
ompound Table					MFG Diff	
Compound Label	RT 0.102	Mass 356 1946	Formula C19 U24 NA C2	MFG Formula	(ppm)	DB Formula
Mpound Label 1 5: C19 H24 N4 O3 MS Spectrum 10 7 Cpd 5: C19 H24	<i>m/z</i> 357.1916 N4 O3: +ESI	RT A 0.102 F	Algorithm Find by Molecular Featu n (0.050-0.851 min) Frag	Mass re 356.1846 =175.0V N-12.d		
mpound Label d 5: C19 H24 N4 O3 MS Spectrum 10 7 Cpd 5: C19 H24 2.5 2 1.5 1 0.5	m/z 357.1916 N4 O3: +ESI '357 ([C19 H24	RT 4 0.102 F MFE Spectrum 7,1916 N4 O3J+H)+	Algorithm Find by Molecular Featu n (0.050-0.851 min) Frag	Mass re 356.1846 =175.0V N-12.d		
Impound Label d 5: C19 H24 N4 O3 MS Spectrum 10 7 Cpd 5: C19 H24 2.5 2 1.5 1 0.5 0 150 200 2 MS Zoomed Spectrum	m/z 357.1916 N4 O3: +ESI '357 ([C19 H24 50 300 35	RT 4 0.102 F MFE Spectrur 7,1916 N4 03J+H)+	Algorithm Find by Molecular Featurn n (0.050-0.851 min) Frag 500 550 600 650 vs. Mass-to-Charge (m/z	Mass re 356.1846 a=175.0V N-12.d	950	
Impound Label d 5: C19 H24 N4 O3 MS Spectrum 10 7 Cpd 5: C19 H24 2.5 2 1.5 1 0.5 0 150 150 2 150 2 150 2 150 2 150 2 150 2 150 2 150 2 150 2 150 2 150 2 2 2 2 2 2 2 2 2 2 3 4 5	<i>m/z</i> 357.1916 `357 ([C19 H24] 50 300 35 \$4 O3 +ESI ([C	RT / 0.102 F 0.102 F MFE Spectrum 7,1916 N4 O3J+H)+ 0 400 450 Counts 0 400 450 Counts 0 357[1916 119 H24 H4 O3	Algorithm Find by Molecular Featu n (0.050-0.851 min) Frag 500 550 600 650 vs. Mass-to-Charge (m/2 n (0.050-0.851 min) Frag 3]+H)+	Mass re 356.1846 g=175.0V N-12.d	950	

Figure S29. ESI (HR-MS) spectrum of 6a.



Figure S30. ESI (HR-MS) spectrum of 6h.



Figure S31. ESI (HR-MS) spectrum of 6i.



Figure S32. HPLC purity analysis of hit compound 6k.

Figure S33: Initial Growth inhibition assay of all fourteen compounds at 1 μ M and 10 μ M with ART as positive control.

Figure S34: A) 3D modelled structure of PlmX. B) Ramachandran Plot, 3D modelled structure. C) Ramachandran Plot, **6k**-PlmX complex after MD. D) A timeline representation of the interactions and total contacts (H-bonds, hydrophobic interactions, ionic interactions, and water bridges) obtained during the molecular dynamics simulations. The panels show the total number of specific contacts the PlmX protein made with the **6k** over the course of the simulation.

Figure S35. Histogram (stacked bar chart) showing compound 6k-PlmX forming H-bonds interactions (green color), hydrophobic interaction (grey violet color), and water bridges (blue color) during 30 ns simulation.

Molecule	#H-bond	#H-	TPSA	XLOGP3	ESOL	ESOL Class	Gl	BBB	Lipinski
	acceptor	donors			LOg S		absorptio	permeant	#violation
3	5	2	71.03	2	-2.86	Soluble	High	Yes	0
6a	6	2	87.58	0.46	-2.15	Soluble	High	No	0
6b	4	2	61.8	3.19	-4.36	Moderatel y soluble	High	Yes	0
6c	7	2	61.8	3.09	-4.12	Moderatel y soluble	High	Yes	0
6d	5	2	74.27	2.02	-3.52	Soluble	High	No	0
6e	4	2	61.8	2.78	-3.69	Soluble	High	Yes	0
6f	4	2	61.8	2.2	-3.25	Soluble	High	Yes	0
6g	5	2	71.03	2.67	-3.64	Soluble	High	No	0
6h	6	2	87.58	0.86	-2.48	Soluble	High	No	0
<u>6i</u>	4	2	61.8	3.13	-3.92	Soluble	High	Yes	0
6j	5	2	68.28	2.2	-3.71	Soluble	High	No	0
6k	4	2	61.8	3.19	-4.36	Moderatel y soluble	High	Yes	0
61	5	2	61.8	2.37	-3.44	Soluble	High	Yes	0
6m	7	2	61.8	3.15	-4.15	Moderatel v soluble	High	Yes	0