

**A semi-automated luminescence based standard membrane feeding assay  
identifies novel small molecules that inhibit transmission of malaria parasites by  
mosquitoes**

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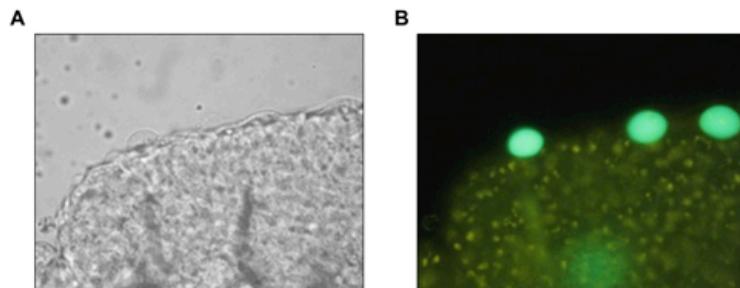
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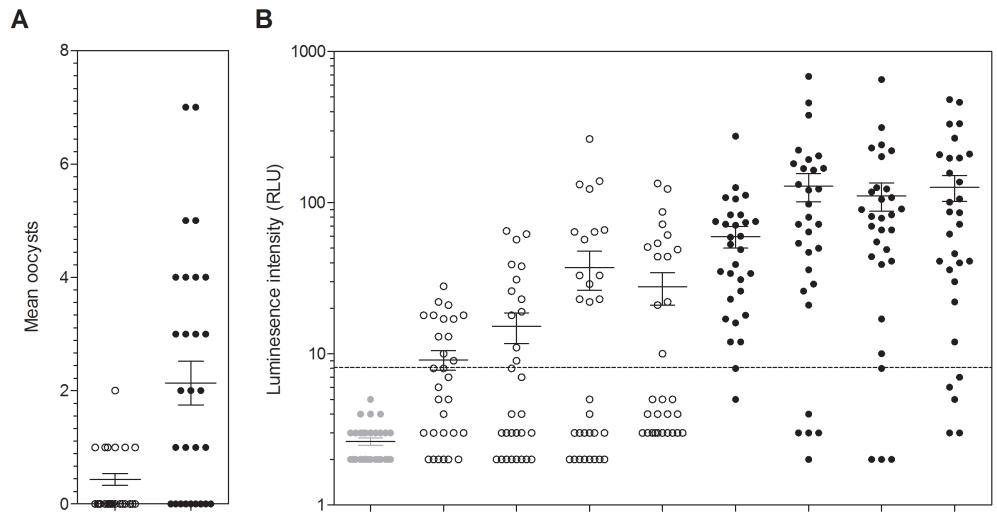
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**SUPPLEMENTARY INFORMATION**



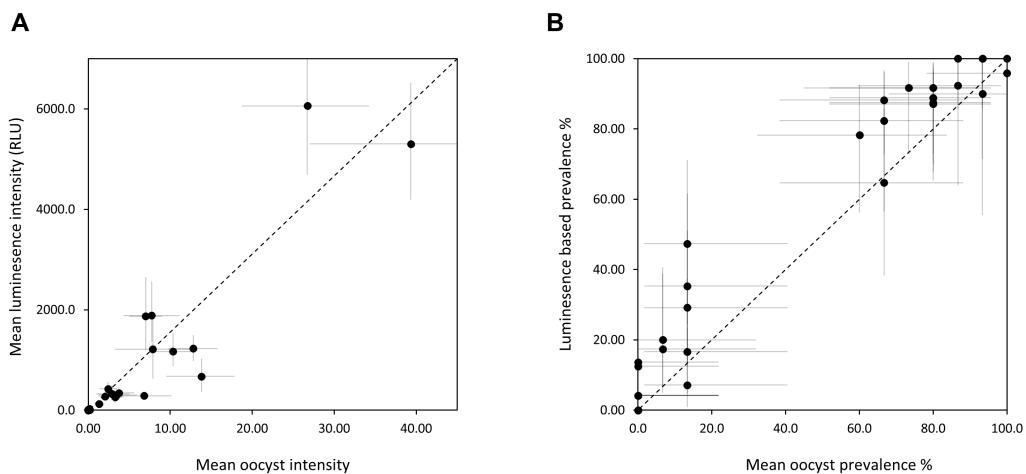
**Figure S1.** **A.** Brightfield micrograph of a section of mosquito midgut with 3 oocysts taken after mosquito dissection 6 days PI with NF54-HGL. **B.** Fluorescence micrograph of the same section of mosquito midgut with 3 oocysts taken after mosquito dissection 6 days PI with NF54-HGL.



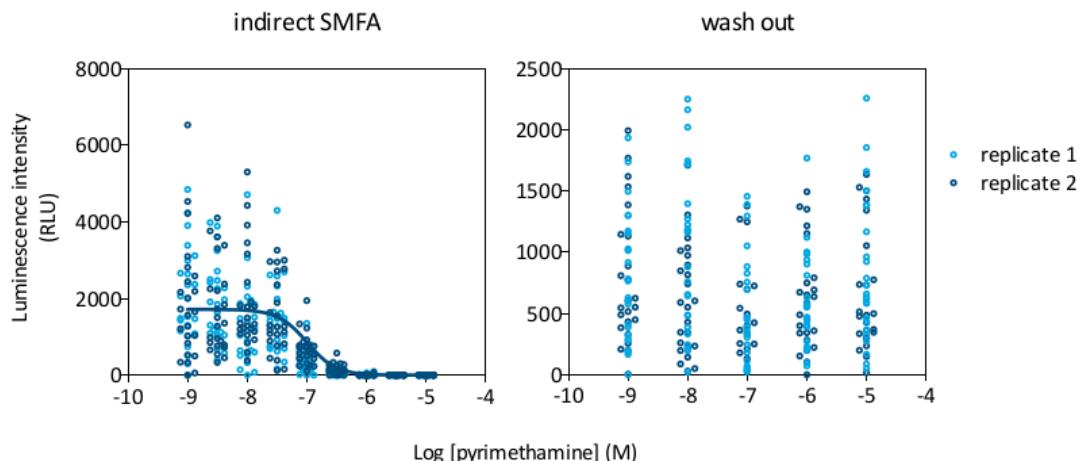
Feed	A	B
Day PI	8	8
Mean oocysts	0.43	2.13
Mean RLU	-	-
Prevalence	40%	70%

Neg	A	A	A	A	B	B	B	B
-	7	8	9	12	7	8	9	12
-	-	-	-	-	-	-	-	-
2.63	9.1	15.2	37.2	27.8	59.7	128.7	111.2	126.2
-	40%	40%	46.7%	43.3%	93.3%	83.3%	86.7%	83.3%

**Figure S2.** **A.** Oocyst intensity in individual mosquitoes from two groups of mosquitoes (*A* and *B*) infected with NF54-HGL, determined after dissection and standard midgut staining and microscopy at day 8 PI. **B.** Luminescence intensity in individual mosquitoes from the same two mosquito groups (*A* and *B*), assessed in separate mosquito samples removed from the primary mosquito storage cage at day 7, 8, 9 and 12 PI. The dashed line indicates the cut-off calculated statistically from the RLU values of all mosquitoes assayed in the current study (8.1 RLU). Prevalence estimates were based on this cut-off.



**Figure S3.** **A.** Relative oocyst and luminescence intensity among mosquitoes sampled from the same cages fed blood-meals without the addition of a test compound (16 feeds, 480 mosquitoes) or with the addition of DHA (16 feeds, 711 mosquitoes). **B.** Relative infection prevalence based on the observation of oocysts or calculated from luminescence intensity, among mosquitoes sampled from the same cages fed blood-meals without the addition of a test compound (16 feeds, 480 mosquitoes) or with the addition of DHA (16 feeds, 711 mosquitoes).



**Figure S4.** Efficacy of the washout protocol. The left panel shows the activity of pyrimethamine in the indirect SMFA as reported in Figure 2 of the main manuscript. The right panel shows the activity in the wash-out mode of the SMFA, where the compound was washed out prior to mosquito feeding.

primer	sequence 5'-3'
MWV 257	GTTAACGAAAGTCCCTACTATTTGAAGAATAGGAACCTACCGCGTGGCGCATGGTACCGTACCTAACTGTTTGAATCTC
MWV 259	CCGGGGTTCAATTCTATGCTAACATACATG
MWV 262	GGCGCCGATTTGGTAGTTGTACTCATCTTG
MWV 267	CGAATTGAAAGTCCCTATTCTAGAAAGTATAGGAACCTCCCGGCATAGTATCTACATACTGAAGCAAG
MWV 252	GGTACCACTAGTCCCGGGATCAACCGGTATGAGTAAGGAGAAGAACCTTCACTGG
MWV 253	TTGAAGTTAACCTTGATTCCATTCTTTGTTTGTCTG
MWV 198	TATCCGGGGCTAGCCGATAAAATCTGGTGAATAC
MWV 200	TATCGGATCCCCTTTGCACTAGCCAATTTTC
MWV 268	AATGGGGCCGAATTGCGGGCTACCTTATTGATGAACTAACAC
MWV 271	TCTAGATITCGACAGGATTATAATATATTATGTACTCACATGGG
MWV 300 (P1)	TACATTCAAATAACTCAGAGGGTAAC
MWV 301 (P2)	GTTTGTATATTTACCTTACATTATCTCC

**Table S1.** List of DNA primer sequences as described in methods.