Alkoxyamines designed as potential drugs against *Plasmodium* and *Schistosoma* parasites

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Supporting Information

Figure S1. Structures of alkoxyamines whose biological evaluation is reported in Table S1. (A) Structures of nitroxide radicals R_1, R_2 –NO•; (B) Structures of alkyl radicals R_3, R_4, R_5 –C•.

Table S1. Antiplasmodial activities of alkoxyamines against chloroquine resistant FcB1-Columbia strain, and antischistosomal activities on adult *S. mansoni*. The numbering of alkoxyamines is made of the number **1-14**, corresponding to the nitroxide moiety, and the letter **a-z** or **A-O**, corresponding to the alkyl moiety (for the structures, see Figure S1, panels A and B, respectively). The IC₅₀ values of artemisinin and chloroquine are provided for comparison. ND = not determined.

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Figure S1. Structures of alkoxyamines whose biological evaluation is reported in Table S1.



(A) Structures of nitroxide radicals R_1, R_2 -NO•.









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EtO

EtO-

P=0

10

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(B) Structures of alkyl radicals R₃,R₄,R₅–C.



Table S1. Antiplasmodial activities of alkoxyamines against the *P. falciparum* chloroquine-resistant strain, FcB1-Columbia, and antischistosomal activities on adults and worms of *S. mansoni*. The numbering of alkoxyamines is made of the number **1-14**, corresponding to the nitroxide moiety, and the letter **a-z** or **A-O**, corresponding to the alkyl moiety (for the structures, see Figure S1, panels A and B, respectively). The IC₅₀ values of artemisinin and chloroquine are provided for antiplasmodial comparison; The value of praziquantel is provided for antischistosomal comparison. ND = not determined.

Compounds	IC ₅₀ (μM) on <i>P. falciparum</i>	Mean (±SD) survival time – Adult S. mansoni (hours)	Time for the death of 100% worms (hours)	Compounds	IC ₅₀ (μM) on <i>P. falciparum</i>	Mean (±SD) survival time – Adult S. mansoni (hours)	Time for the death of 100% worms
1	>> 10	> 8	>8	(<i>RR/SS</i>)-1v	>> 10	3.4 ± 0.2	4
(<i>RS/SR</i>)-1a	3.8	1.2 ± 0.1	2	(<i>RS/SR</i>)-1v	>> 10	> 8	>8
(<i>RR/SS</i>)-1a	>> 10	> 8	>8	(<i>RR/SS</i>)-1w	>> 10	2.1 ± 0.1	3
(<i>RS/SR</i>)-1b	>> 10	ND	ND	(<i>RS/SR</i>)-1w	>> 10	> 8	>8
(<i>RR/SS</i>)-1b	22.5	ND	ND	(<i>RR/SS</i>)-1x	>> 10	> 8	>8
(<i>RR/SS</i>)-1c	>> 10	ND	ND	(<i>RS/SR</i>)-1x	>> 10	2.4 ± 0.3	4
(<i>RR/SS</i>)-1d	>> 10	> 8	>8	(<i>RR/SS</i>)-1y	>> 10	> 8	>8
(<i>RS/SR</i>)-1e	>> 10	> 8	>8	(<i>RS/SR</i>)-1y	>> 10	> 8	>8
(<i>RS/SR</i>)-1f	>> 10	ND	ND	(<i>RS/SR</i>)-1A	>> 10	1.0 ± 0.05	1
(<i>RS/SR</i>)-1g	>> 10	2.6 ± 0.3	4	(<i>RR/SS</i>)-1A	>> 10	1.0 ± 0.05	1
(<i>RR/SS</i>)-1g	>> 10	1.9 ± 0.1	2	(<i>RS/SR</i>)-1B	>> 10	ND	ND
(<i>RS/SR</i>)-1h	>> 10	> 8	>8	(<i>RS/SR</i>)-1C	>> 10	1.4 ± 0.2	2
(<i>RS/SR</i>)-1i	>> 10	> 8	>8	(<i>RR/SS</i>)-1C	>> 10	ND	ND
(<i>RR/SS</i>)-1i	>> 10	> 8	>8	(<i>RS/SR</i>)-1D	>> 10	3.8 ± 0.5	6
(<i>RS/SR</i>)-1j	>> 10	> 8	>8	(<i>RS/SR</i>)-1E	>> 10	> 8	>8
(<i>RR/SS</i>)-1j	>> 10	> 8	>8	(<i>RR/SS</i>)-1E	>> 10	5.0 ± 0.5	6
(<i>RR/SS</i>)-1k	>> 10	1.0 ± 0.05	1	1F	>> 10	ND	ND
(<i>RS/SR</i>)-1k	>> 10	> 8	>8	1H	>> 10	> 8	>8
(<i>RR/SS</i>)-11	>> 10	4.5 ± 0.5	6	(<i>RS/SR</i>)-2c	>> 10	ND	ND
(<i>RS/SR</i>)-11	>> 10	> 8	>8	(<i>RR/SS</i>)-2c	>> 10	ND	ND
(<i>RR/SS</i>)- 1m	>> 10	> 8	>8	(<i>RS/SR</i>)-2z	>> 10	> 8	>8

(<i>RS/SR</i>)- 1m	>> 10	5.1 ± 0.4	6	(<i>RS/SR</i>)-2F	0.3	2.1 ± 0.1	3
(<i>RR/SS</i>)-10	25	2.3 ± 0.2	3	(<i>RR/SS</i>)-2F	0.2	2.5 ± 0.2	5
(<i>RS/SR</i>)-10	25	1.1 ± 0.1	2	(<i>RS/SR</i>)-3z	>> 10	ND	ND
(<i>RR/SS</i>)-1n	>> 10	> 8	>8	(<i>RS/SR</i>)-4z	>> 10	> 8	>8
(<i>RS/SR</i>)-1n	>> 10	> 8	>8	(<i>RR/SS</i>)-4z	>> 10	> 8	>8
(<i>RR/SS</i>)-1p	20.5	1.0 ± 0.05	1	(<i>RS/SR</i>)-4F	2.9	5.2 ± 0.2	7
(<i>RS/SR</i>)-1p	20.5	1.4 ± 0.3	3	(RR/SS)- 4 F	1.3 ± 0.7	> 8	>8
(<i>RR/SS</i>)-1q	>> 10	3.3 ± 0.9	6	(<i>RS/SR</i>)-5z	>> 10	> 8	>8
(<i>RS/SR</i>)-1q	>> 10	1.3 ± 0.2	2	51	>> 10	ND	ND
(<i>RR/SS</i>)-1r	>> 10	3.0 ± 0.8	4	6c	ND	ND	ND
(<i>RS/SR</i>)-1r	>> 10	3.1 ± 0.1	6	7z	ND	ND	ND
(<i>RR/SS</i>)-1s	>> 10	> 8	>8	10z	>> 10	ND	ND
(<i>RS/SR</i>)-1s	>> 10	3.3 ± 0.1	4	11z	>> 10	ND	ND
(<i>RR/SS</i>)-1t	>> 10	ND	ND	12z	>> 10	ND	ND
1u	>> 10	ND	ND	13z	>> 10	ND	ND
(<i>RR/SS</i>)- 2G	>> 10	6.4 ± 0.1	7	8 F	0.14 ± 0.03	5.0 ± 0.0	5
(<i>RS/SR</i>)-2G	>> 10	> 8	>8	2	>> 10	> 8	>8
5N	>> 10	ND	ND	8	>> 10	> 8	>8
5a	>> 10	ND	ND	22	0.91 ± 0.06	> 8	>8
7z	>> 10	ND	ND	2b	>> 10	>8	>8
(<i>RS/SR</i>)-7F	2.1 ± 2.2	2.7 ± 0.1	3	9b	>> 10	3.0 ± 0.0	3
(<i>RR/SS</i>)-7F	1.5 ± 0.8	2.5 ± 0.1	3	8b	>> 10	> 8	>8
1J	3.1 ± 1.1	1.1 ± 0.05	2	9z	>> 10	> 8	>8
14a	>> 10	2.5±0.10	3	9a	8 ± 3	3.0 ± 0.0	3
2a	>> 10	> 8	>8	8a	>> 10	> 8	>8
1h	>> 10	ND	ND	8z	>> 10	1.9 ± 0.1	2
1g	>> 10	ND	ND	(<i>RS/SR</i>)-1a	>> 10	> 8	>8
1K	1.3 ± 0.3	1.1 ± 0.04	2	(<i>RS/SR</i>)-1a	>> 10	> 8	>8
2J	2.5 ± 1.5	1.0 ± 0.0	1	(RR/SS)-1M	4.8	ND	ND
(RS/SR)-10	>> 10	ND	ND	Artemisinin	0.04 ± 0.007	ND	ND
(RS/SR)-1L	12	ND	ND	Chloroquine	0.16 ± 0.036	ND	ND
(RR/SS)-1L	28	ND	ND	Praziquantel	ND	1.0 ± 0.05	1
(KS/SK)- 1M	2.7	ND	ND				



¹³C-NMR (CDCl₃)



Figure S3. NMR of 4'-(1-bromoethyl)-2,2':6',2"-terpyridine 18.



Figure S4. NMR of (*RS/SR*)-6F.







Figure S5. NMR of (*RR/SS*)-6F. ¹H-NMR (CDCl₃)



¹³C-NMR (CDCl₃)















¹³C-NMR (CDCl₃)



















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Figure S14. NMR of (*RR/SS*)-6G.





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¹³C-NMR (CDCl₃)





Table S2. XRD data for (*RR/SS*)-6F

Compound	(<i>RR/SS</i>)-6F
Empirical formula	C ₃₆ H ₅₇ N ₄ O ₅ PSi
Formula weight	684.91
Temperature/K	293
Crystal system	monoclinic
Space group	$P2_1/c$
a/Å	20.9011(7)
b/Å	14.8947(5)
c/Å	13.4391(4)
α/°	90
β/°	107.315(4)
γ/°	90
Volume/Å ³	3994.2(2)
Z	4
$\rho_{calc}g/cm^3$	1.139
μ/mm^{-1}	0.141
F(000)	1480.0
Crystal size/mm ³	$0.5\times0.28\times0.28$
Radiation	MoKa ($\lambda = 0.71073$)
2Θ range for data collection/°	5.838 to 52.744
Index ranges	$-26 \le h \le 25, -18 \le k \le 18, -16 \le l \le 16$
Reflections collected	52925
Independent reflections	$8087 [R_{int} = 0.1388, R_{sigma} = 0.0555]$
Data/restraints/parameters	8087/15/432
Goodness-of-fit on F ²	1.084
Final R indexes [I>= 2σ (I)]	$R_1 = 0.0910$, $wR_2 = 0.2628$
Final R indexes [all data]	$R_1 = 0.1166, wR_2 = 0.3005$
Largest diff. peak/hole / e Å ⁻³	0.81/-0.77

Table S3. XRD data for (*RR/SS*)-6G.

Compound	(<i>RR/SS</i>)-6G
Empirical formula	C33H46NO5P
Formula weight	567.68
Temperature/K	295
Crystal system	monoclinic
Space group	Pc
a/Å	11.7459(5)
b/Å	18.2946(7)
c/Å	16.2012(6)
$\alpha/^{\circ}$	90
β/°	110.150(4)
γ/°	90
Volume/Å ³	3268.3(2)
Z	4
$\rho_{calc}g/cm^3$	1.154
µ/mm ⁻¹	1.049
F(000)	1224.0
Crystal size/mm ³	$0.26 \times 0.2 \times 0.12$
Radiation	$CuK\alpha \ (\lambda = 1.54184)$
2Θ range for data collection/°	7.558 to 142.748
Index ranges	$-14 \le h \le 14, -22 \le k \le 22, -19 \le l \le 19$
Reflections collected	27391
Independent reflections	9765 [$R_{int} = 0.0417$, $R_{sigma} = 0.0321$]
Data/restraints/parameters	9765/45/884
Goodness-of-fit on F ²	1.023
Final R indexes [I>= 2σ (I)]	$R_1 = 0.0773, wR_2 = 0.2160$
Final R indexes [all data]	$R_1 = 0.0829, wR_2 = 0.2293$
Largest diff. peak/hole / e Å ⁻³	0.91/-0.32
Flack parameter	0.01(3)

Figure S17. Proposed mechanism for the formation of compounds 24 and 25 (see Oxygen Radicals, J. K. Kochi, in *Free Radicals*, J. K. Kochi, ed., John Wiley & Sons (1973), Vol. II, p. 686ss.



Figure S18. ¹H NMR spectrum of (RS/SR)-2F and (RS/SR)-2FH⁺ in *t*-BuPh.



Figure S19. ¹H NMR spectrum of (RR/SS)-2F and (SS/RR)-2FH⁺ in *t*-BuPh.



12.6 12.2 11.8 11.4 11.0 10.6 10.2 58 5.6 5.4 5.2 5.0 8.8 8.6 8.4 8.2 8.0 7.8 7.6 7.4 7.2 7.0 rf.formi