

The antimalarial efficacy and mechanism of resistance of the novel chemotype DDD01034957

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Parasite	3D7 (parental)	Clone D5	Clone E2	Clone H9	Clone F3	Clone B2	Clone D2	Clone E11	Clone D3	Clone C11	Clone D3	Clone G8
IC ₅₀ (μM)	0.14	3.33	2.49	3.56	2.62	1.81	2.77	3.28	2.40	1.57	ND	ND
fold-change		24.47	18.30	26.16	19.25	13.30	20.35	24.10	17.63	11.54		
PfABC13 mutation		F2010L	F2010L	F2010L	F2010L	F2010L	F2010L	F2010L	F2010L	H2181D	L79F	H2181D
		Expt 1			Expt 2					Expt 3		

Supplementary Table S1 - *In vitro* selection of resistance and genotyping. *P. falciparum* asexual parasites were pulsed with DDD01034957 at 0.2μM or 2μM. When recovered, surviving parasites were cloned and tested for resistance to DDD01034957. From three independent selection experiments, twelve clones were obtained with a 12.1 to 26.2-fold increase in IC₅₀ to DDD01034957. Whole genomic sequencing revealed all clones had mutations in PF3D7_0319700 (*PfABC13*).