

**S1 Table. Origins, alternative names, and drug resistance haplotypes of the strains used in this study.**

	<b>3D7</b>	<b>V1/S</b>	<b>W2</b>	<b>Dd2</b>	<b>Dd2 <i>exo1A</i></b>	<b>PH0167-C<sup>a</sup></b>	<b>PH0306-C<sup>a,b</sup></b>	<b>PH0164-C<sup>a,b</sup></b>	<b>PH0482-C<sup>a</sup></b>	<b>PH0212-C<sup>a</sup></b>
<b>Original ID</b>	-	-	-	-	-	RF 918	RF 967	RF 915	RF 976	RF 986
<b>Sanger ID</b>	-	-	-	-	-	PH0167-C	PH0306-C	PH0164-C	PH0482-C	PH0212-C
<b>Other ID<sup>b</sup></b>	-	-	-	-	-	-	Cam3.II	CamWT	-	-
<b><i>k13</i></b>	WT	WT	WT	WT	WT	WT	WT	R539T	C580Y	C580Y
<b><i>pfert</i></b>	WT	Dd2 <sup>c</sup>	Dd2	Dd2	Dd2	Dd2	Dd2	Dd2	Dd2	Dd2
<b><i>KH group</i><sup>d</sup></b>	-	-	-	-	-	1	1	3	2	2
<b>Origin</b>	Netherlands	Vietnam	Vietnam	W2-mef	Dd2	Cambodia	Cambodia	Cambodia	Cambodia	Cambodia
<b>Year</b>	1970s	1980	1980	1980s	2014	2010	2010	2010	2010	2010

<sup>a</sup> Published in [20]. <sup>b</sup> Published in [21]. <sup>c</sup> Dd2 *pfert* allele: 74I/75D/76T/220S/271E/326S/356T/371I. <sup>d</sup> KH groups are Cambodian subpopulation cluster groups determined in [9].