

9M6 (Liverpool)	1:	MVLLDLLVLIPIVSYLLYRWAVATYDFFEKRKIPYVKPYPFVGGGLWPFVSGKLHPTDAA	60
9M6v1 (SP)	1:	-----	60
9M6v2 (SP)	1:	-----	60
9M6v3 (SMK)	1:	-----	60
9M6v4 (SMK)	1:	-----	60
9M6v5 (SMK)	1:	-----	60
9M6 (Liverpool)	61:	VLGYNLFPENRFSGFFAFRRPGYL I HDPALAKQIMIKDFDHFTHMNTISVDVDP I FGRA	120
9M6v1 (SP)	61:	-----	120
9M6v2 (SP)	61:	-----	120
9M6v3 (SMK)	61:	-----T-----	120
9M6v4 (SMK)	61:	-----	120
9M6v5 (SMK)	61:	-----	120
9M6 (Liverpool)	121:	LFFMDGQRWRHGRSGLSPAFTGSKMRNMFTLLSKYVEGAMQRLAQDAGQGKMELEIRDLF	180
9M6v1 (SP)	121:	-----R-----V-----	180
9M6v2 (SP)	121:	-----	180
9M6v3 (SMK)	121:	-----	180
9M6v4 (SMK)	121:	-----	180
9M6v5 (SMK)	121:	-----	180
9M6 (Liverpool)	181:	QKLGNDIITSISFGVEIDSVHNPNEFFKRGKQLAATGGFQGLKFFFSLVVPDSVFKLFG	240
9M6v1 (SP)	181:	-----	240
9M6v2 (SP)	181:	-----	240
9M6v3 (SMK)	181:	-----	240
9M6v4 (SMK)	181:	-----	240
9M6v5 (SMK)	181:	-----	240
9M6 (Liverpool)	241:	IRFLPKEADFYVDVVSKT I KHREEYK I VRPDF I HLFVQARKNELKEETADDELKSAGFT	300
9M6v1 (SP)	241:	-----	300
9M6v2 (SP)	241:	-----	300
9M6v3 (SMK)	241:	-----	300
9M6v4 (SMK)	241:	-----	300
9M6v5 (SMK)	241:	-----	300
9M6 (Liverpool)	301:	TVEEHIEASTENSQYTDLDITAVAASFFFGGIETTTMLCFALYELAGNKEVQQKLQAEI	360
9M6v1 (SP)	301:	-----	360
9M6v2 (SP)	301:	-----E-----	360
9M6v3 (SMK)	301:	-----	360
9M6v4 (SMK)	301:	-----	360
9M6v5 (SMK)	301:	-----	360
9M6 (Liverpool)	361:	DSVRKELGGGSLTYEVLQKMKYLDMVVTETLRRWPPLGI TNRVCVKPYTFEDHEGTVKVI	420
9M6v1 (SP)	361:	-----E-----	420
9M6v2 (SP)	361:	-----ED-----M-----	420
9M6v3 (SMK)	361:	-----E-----	420
9M6v4 (SMK)	361:	-----	420
9M6v5 (SMK)	361:	-----E-----	420

9M6 (Liverpool)	421: EKGQLIQIPVQSFHRDPSFFPDYRFDPERFSEENKHKINQDAFLPFGSGPRNCIGSRLA	480
9M6v1 (SP)	421: -----N-----	480
9M6v2 (SP)	421: -----N-----	480
9M6v3 (SMK)	421: -----N-----	480
9M6v4 (SMK)	421: -----	480
9M6v5 (SMK)	421: -----N-----*	473
9M6 (Liverpool)	481: LMQAKCLLYLFSAFSLEYSKMDVPIKLNKMSLTYTAKNGFWFNLLPKKVAV*	533
9M6v1 (SP)	481: -----*	533
9M6v2 (SP)	481: -----V-R-----*	533
9M6v3 (SMK)	481: -----V-R-----*	533
9M6v4 (SMK)	481: -----*	533
9M6v5 (SMK)	473: -----	473

Figure S4 Multiple alignments of CYP9M6 amino acid sequences identified from Liverpool, SP, and SMK strains of *Aedes aegypti*. Hyphens indicate identical amino acids to the sequence of Liverpool strain which was used for the genome project. Asterisks indicate stop codons.