

	10	20	30	40	50	60
Sudan, Sud-60-93-11 isolate	CAAAGCC	TGCA	GATGCTG	TAA	GTACT	CAAAAGTGCT
Laos, RO33 type 1 (n=66)*	CAAAGCC	TGCA	GATGCTG	TAA	GTACT	CAAAAGTGCT
Savannakhet, RO33 type 2 (n=1)					TGT	
Savannakhet, RO33 type 3 (n=1)					T	G
Savannakhet, RO33 type 4 (n=1)					AG	
Savannakhet, RO33 type 5 (n=1)					A	G
Xekong, RO33 type 6 (n=1)						CAA
Xekong, RO33 type 7 (n=1)						
Thailand, clone 814C						
Thailand, isolate T838						
Cambodia, Cam84I-5 isolate						
Indonesia, isolate 4/M1						
India, RDTCF002						
Iran, isolate CH10						
Malawi, 2/040-1-8 isolate						
Malawi, 2/041-3-1 isolate						
Sudan, Sud-60-93-3						
Tanzania, isolate IFA17.29						
West Africa PFAMS1 isolate						
Ghana, isolate RO33						
Brazil, 7G8 isolate						
Peru, RO33-140.1 isolate						

	70	80	90
Sudan, Sud-60-93-11 isolate	CTTCAGG	TACTGCA	AGTACTAA
Laos, RO33 type 1 (n=66)*	CTTCAGG	TACTGCA	AGTACTAA
Savannakhet, RO33 type 2 (n=1)			
Savannakhet, RO33 type 3 (n=1)			
Savannakhet, RO33 type 4 (n=1)			
Savannakhet, RO33 type 5 (n=1)			
Xekong, RO33 type 6 (n=1)			
Xekong, RO33 type 7 (n=1)			TG
Thailand, clone 814C			
Thailand, isolate T838			
Cambodia, Cam84I-5 isolate			
Indonesia, isolate 4/M1			
India, RDTCF002			
Iran, isolate CH10			
Malawi, 2/040-1-8 isolate			
Malawi, 2/041-3-1 isolate			
Sudan, Sud-60-93-3			
Tanzania, isolate IFA17.29			
West Africa PFAMS1 isolate			
Ghana, isolate RO33			
Brazil, 7G8 isolate			
Peru, RO33-140.1 isolate			

* 10 sequences of Laos, RO33 type 1 derived from sequence analysis (2 from Oudomxay, 5 from Savannakhet and 3 from Xekong), 56 sequences were expected from SSCP screening.