

TABLE S4. Number of repeats of sequenced private alleles

Isolate number	Population	Locus	Amplicon size (bp) ^b	# repeats	Repeat motif
CMW45653	South Sumatra, Indonesia	A	198	8	TGG
CMW48638	Malaysia	A	192	6	TGG
CMW44962 ^a	South Africa	A	195	7	TGG
CMW45653	South Sumatra, Indonesia	C	233	11	TG
CMW48654	Malaysia	C	237	13	TG
CMW44962 ^a	South Africa	C	235	12	TG
CMW45653	South Sumatra, Indonesia	G	415	11	TG
CMW44962 ^a	South Africa	G	413	10	TG
CMW48652	South Sumatra, Indonesia	H	311	5	CCA
CMW44962 ^a	South Africa	H	320	8	CCA
CMW48640	Malaysia	J	382	22	AG
CMW44962 ^a	South Africa	J	372	17	AG
CMW45686	North Sumatra, Indonesia	K	444	10	TCA
CMW44962 ^a	South Africa	K	441	9	TCA
CMW48638	Malaysia	L	393	7	ATC
CMW44962 ^a	South Africa	L	402	10	ATC

^a CMW44962 genome isolate used as positive control

^b Determined with GeneScan