

Table S3. Sequence sampling to assess haplotype (strain) diversity of Villages

Cumulative density of each village level sample population given a value of diversity, θ , and number of sequences. Projections to capture 95% of the total probability are also provided.

Village	Peneng	Albulum2	Albulum1	Yautong1	Yautong2	Moilenge	Moihauk
ϑ	4.31	6.58	7.14	5.40	6.48	5.23	6.44
Number of sequences/village	69	134	147	23	42	25	47
Cumulative Density	0.94	0.95	0.95	0.79	0.86	0.81	0.87
Number of sequences/village to 95%	84	129	140	106	127	103	126