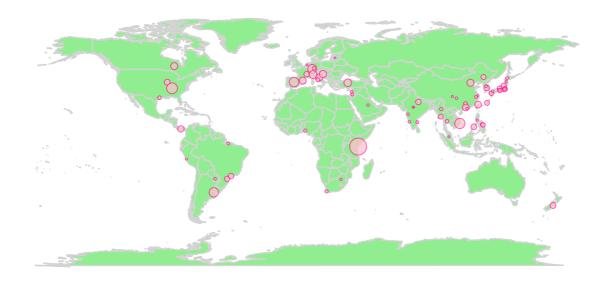
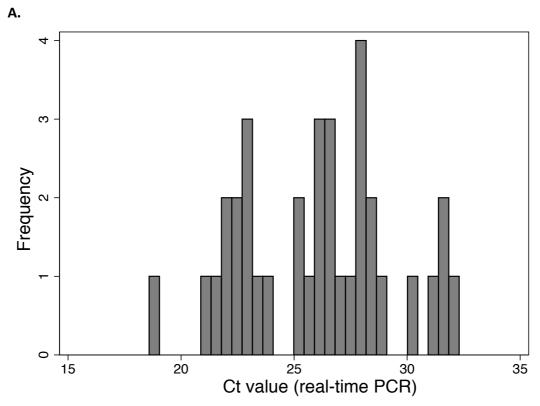
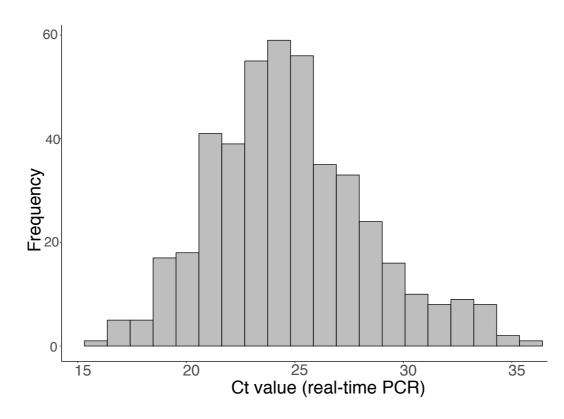
S1 Fig: Sampling locations of the global ON1 G-gene dataset with circles representative of relative proportion of contributing sequences by country



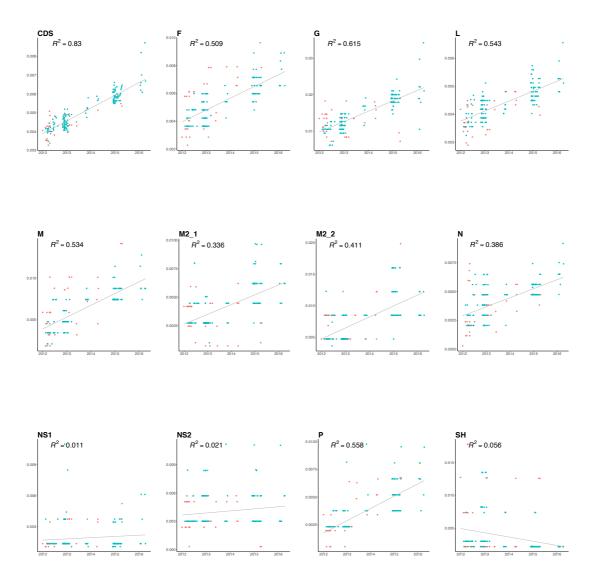
S2 Fig: Histograms showing distribution of Ct values for samples from the (A) KHDSS and (B) KCH



В.



S3 Fig: Root-to-tip regression analysis of Kilifi RSV-A ORFs



S4 Fig: BEAST MCC trees showing divergence between ON1(cyan) and GA2 (red) ORFs

