

Detection of Measles Virus Genotype B3, India



Technical Appendix Figure. Phylogenetic analysis of B3 genotype measles virus sequences detected in India with selected global B3 genotypes. A D8 genotype sequence from Thiruvananthapuram was used as the outgroup. We performed maximum-likelihood analysis with 1,000 bootstrap replications using Kimura2-parameter correction with gamma-distributed settings. Solid red triangles indicate strains from the current study; scale bar indicates nucleotide substitutions per site.