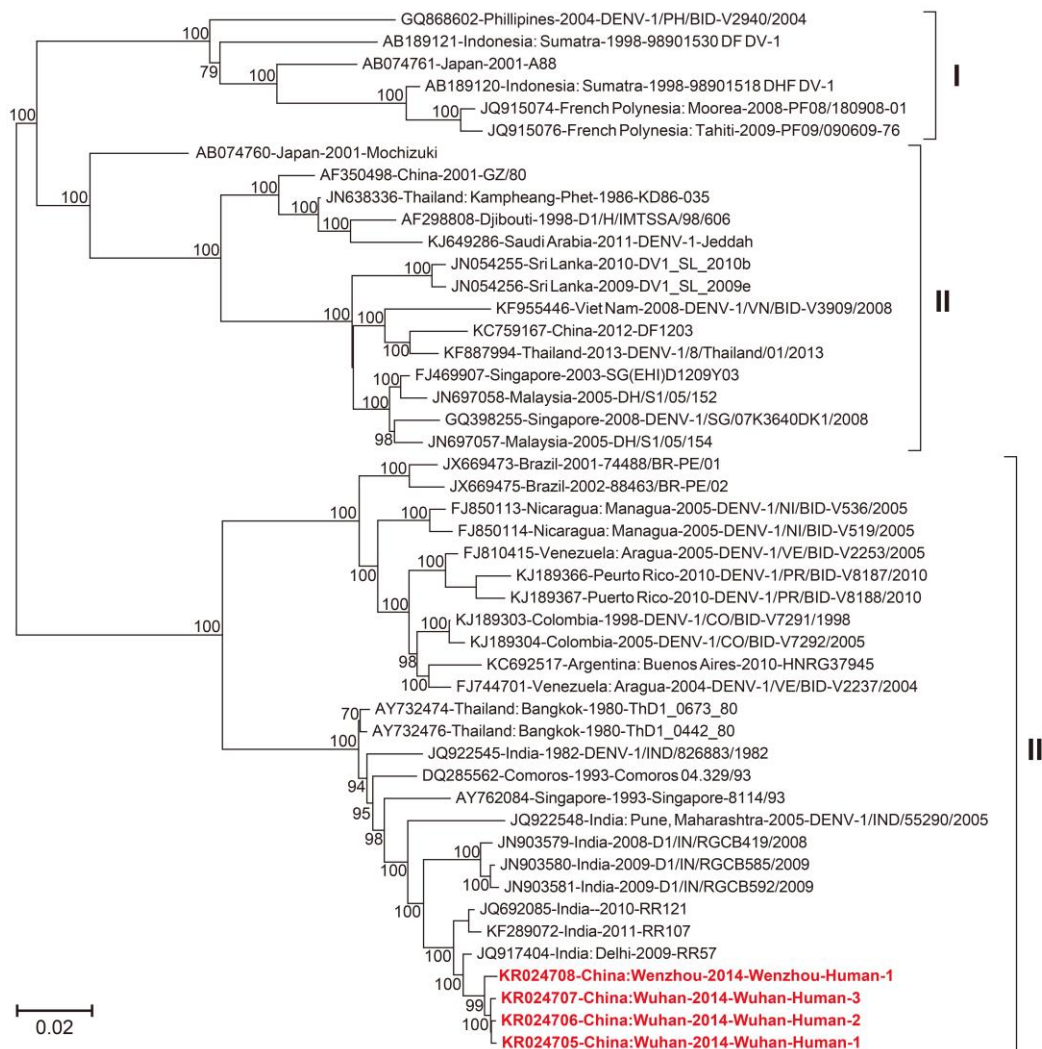


Reemergence and Autochthonous Transmission of Dengue Virus, Eastern China, 2014

Technical Appendix



Technical Appendix Figure. Phylogenetic tree of whole genome sequences of DENV-1 showing the position of the Wenzhou and Wuhan viruses within viral genotype III. Bootstrap values (>70%) are shown at relevant nodes. The tree is mid-point rooted for clarity. Scale bar indicates nucleotide substitutions per site.