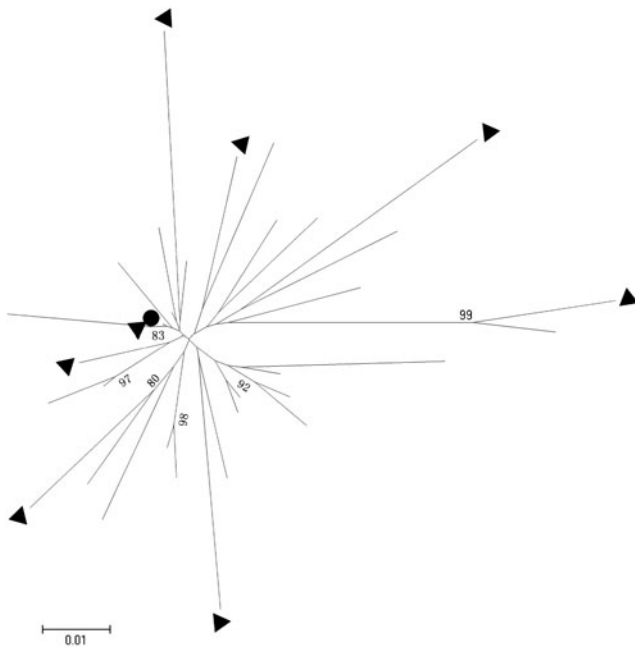# Supplementary Data

SUPPLEMENTARY FIG. S1. The neighbor-joining tree was constructed using an *env* c2v3 set. A dot represents predicted consensus sequences; a triangle represents chosen strains with full-length genomic sequences.