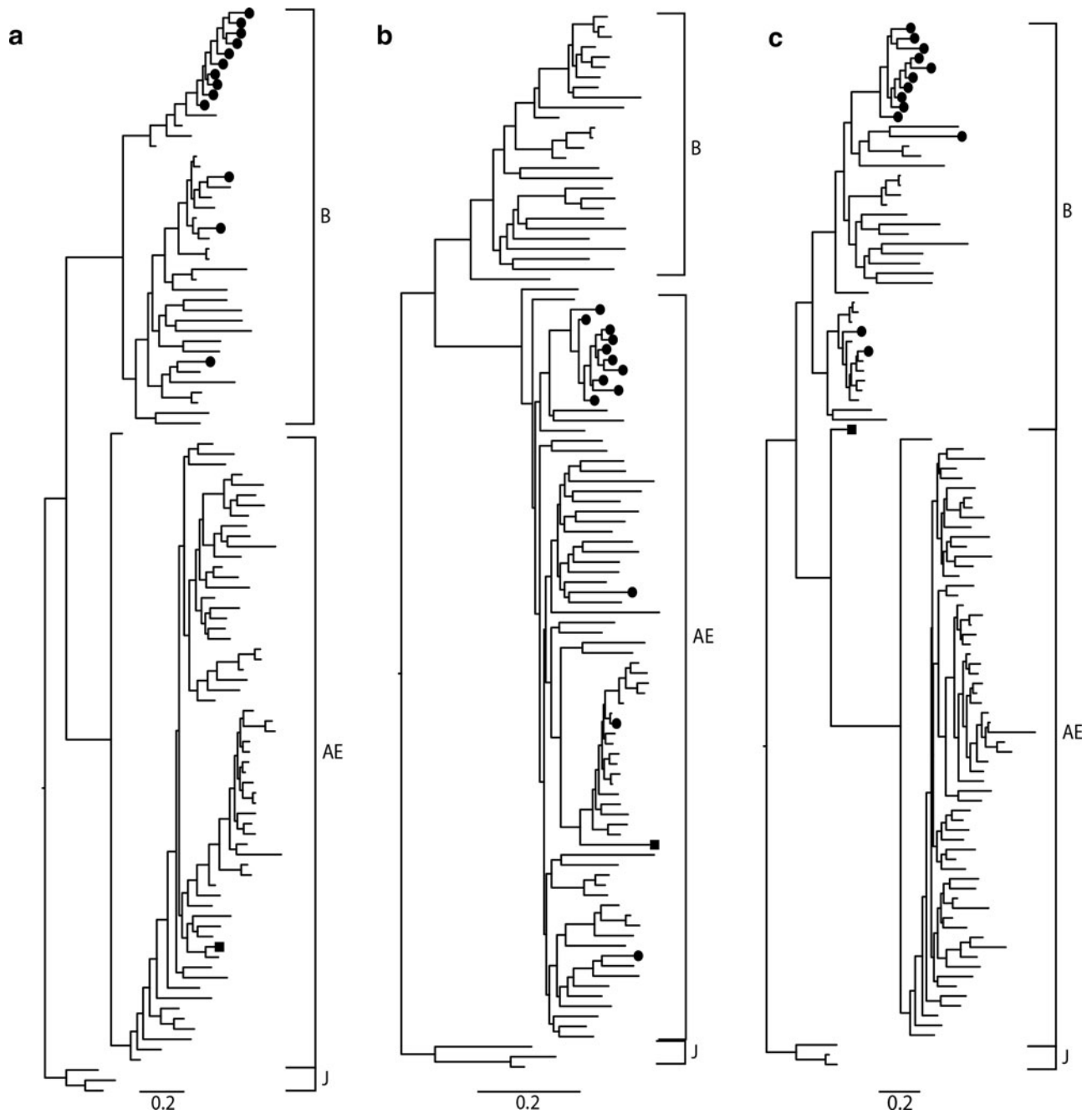


Supplementary Data



SUPPLEMENTARY FIG. S1. Phylogenies of 109 isolates successfully sequenced in all three regions. For clarity of presentation, only those clustering with subtype B and CRF01_AE are shown. Regions shown are (a) protease, (b) gp120, and (c) gp41. Filled circles = 13 sequences with an RF1 pattern; filled square = one sequence with an RF2 pattern. Subtype J was included as an outgroup.