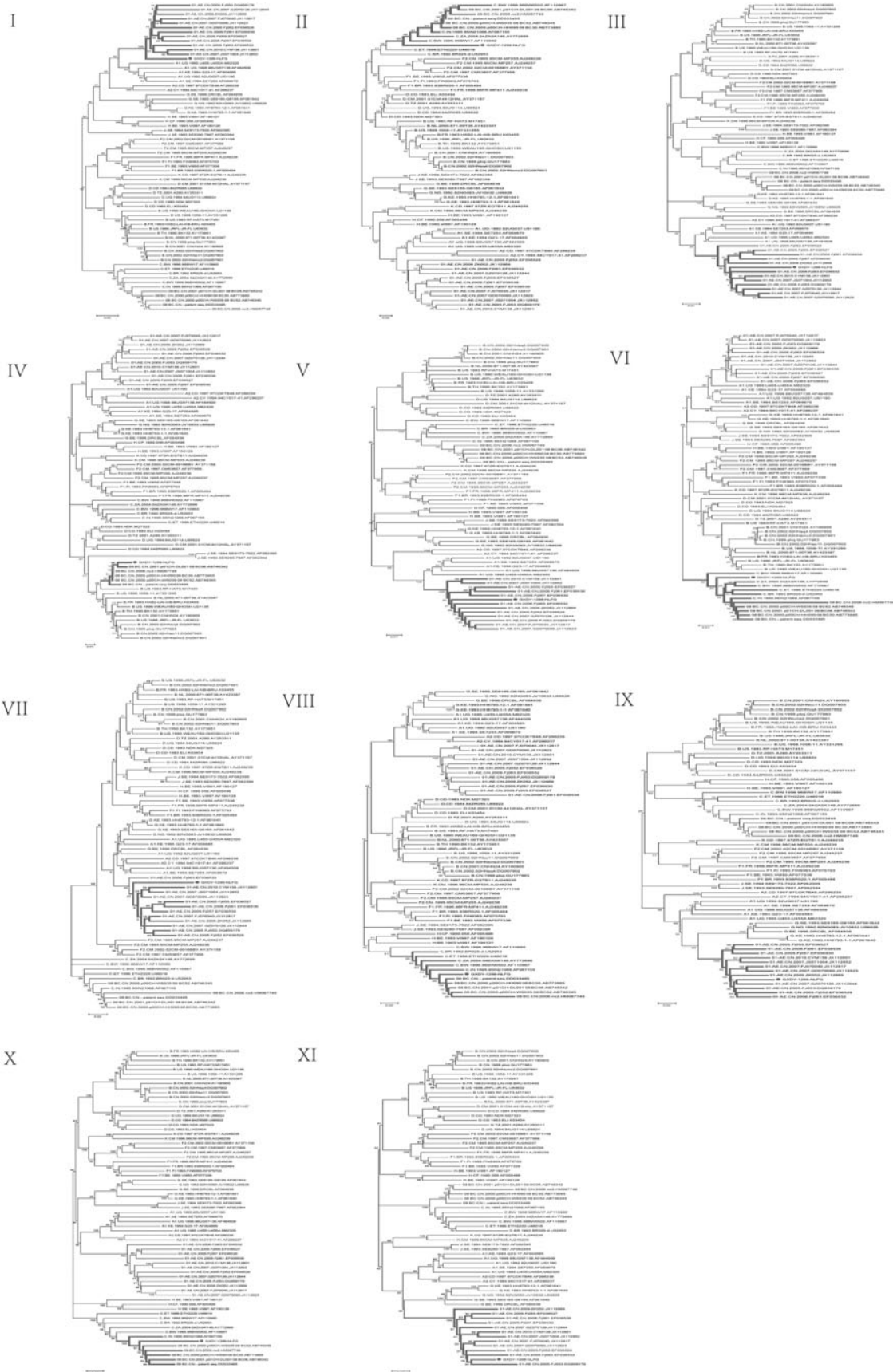# Supplementary Data

SUPPLEMENTARY FIG. S1. Phylogenetic analyses of 11 mosaic segments. The phylogenetic trees of the 11 mosaic segments were constructed with MEGA 5.0 using the neighbor-joining method with 1,000 replications. The subtype references are derived from the Los Alamos HIV Sequence Database. The clade including GXDY1299 is marked in red and is in bold.