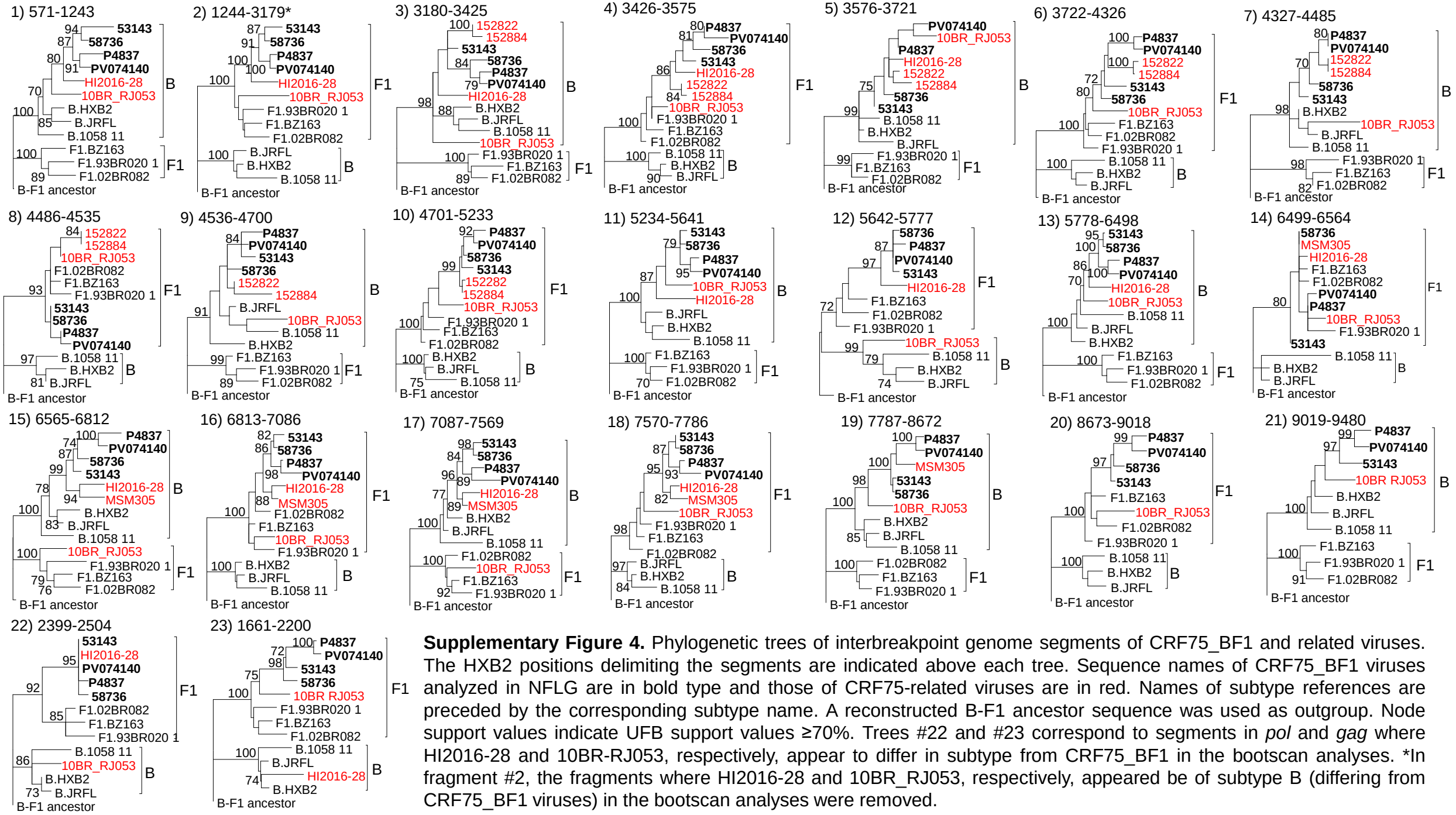

Supplementary Figure 4. Phylogenetic trees of interbreakpoint genome segments of CRF75_BF1 and related viruses. The HXB2 positions delimiting the segments are indicated above each tree. Sequence names of CRF75_BF1 viruses analyzed in NFLG are in bold type and those of CRF75-related viruses are in red. Names of subtype references are preceded by the corresponding subtype name. A reconstructed B-F1 ancestor sequence was used as outgroup. Node support values indicate UFB support values ≥70%. Trees #22 and #23 correspond to segments in *pol* and *gag* where HI2016-28 and 10BR-RJ053, respectively, appear to differ in subtype from CRF75_BF1 in the bootscan analyses. *In fragment #2, the fragments where HI2016-28 and 10BR_RJ053, respectively, appeared be of subtype B (differing from CRF75_BF1 viruses) in the bootscan analyses were removed.