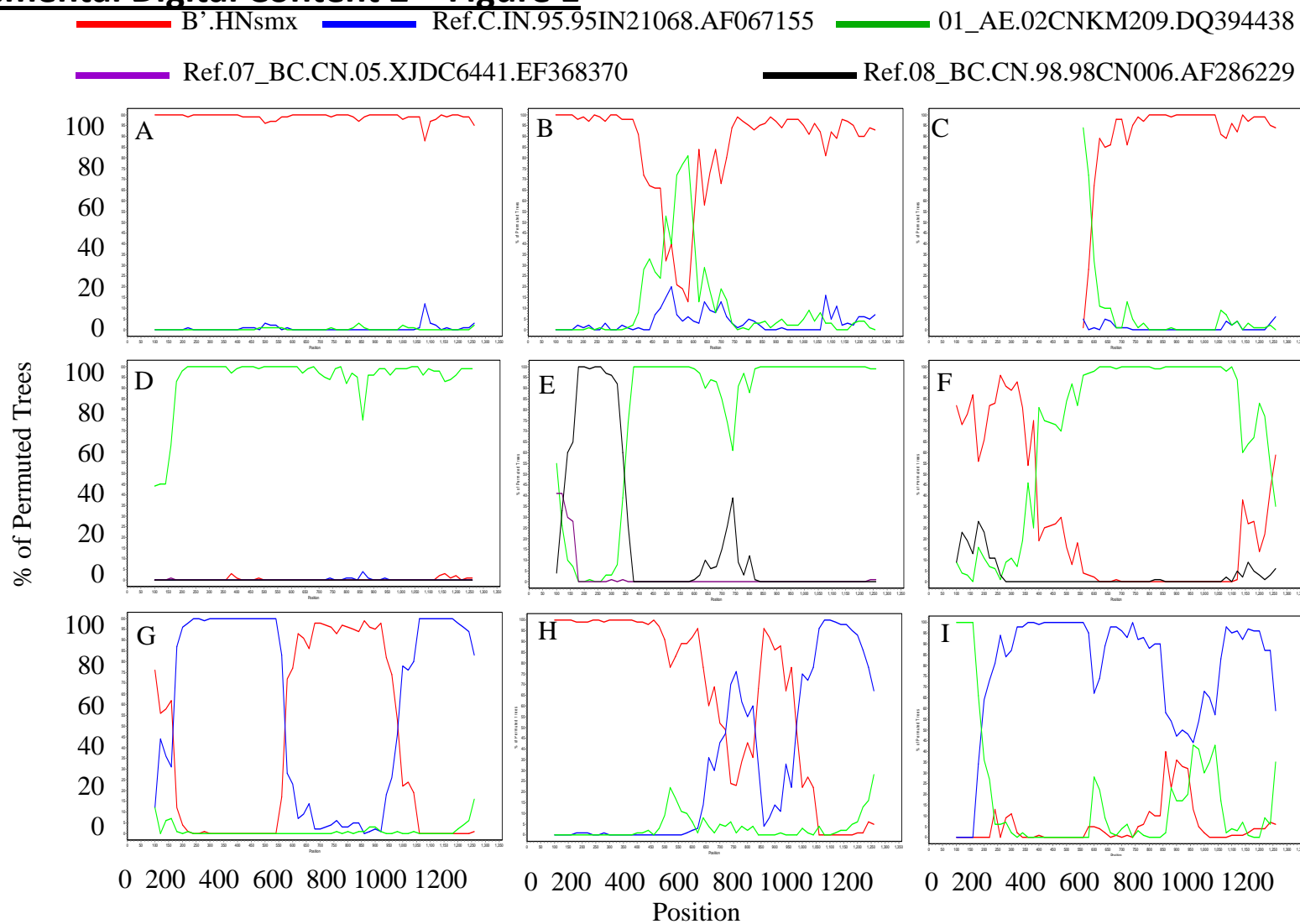


Supplemental Digital Content 2 – Figure 1



Window: 200bp, Step: 20bp, GapStrip: On, Reps: 100, Kimura (2-parameter), T/t: 2.0, Neighbor-Joining

Suppl Fig. 1. The Bootscan Analysis of HIV-1 RT Sequences of Nine Study Participants. These participants included one B' virus Ah.79H081554 (A), three B'/CRF01_AE recombinant viruses Cq.67H081498 (B), Hen.53BH081312 (C), Fj.Xm.69H081500 (F), one CRF01_AE virus Fj.Fz.12H080847 (D), one CRF01_AE/CRF08_BC recombinant virus Fj.Fz.46H081045 (E), one CRF08_BC virus Fj.Fz.23H080912 (G), one CRF07_BC virus Xj.87H082047 (H), and one C/CRF01_AE recombinant virus Fj.Fz.112BH081488 (I).