SUPPLEMENTARY FIG. S2. Demographic history of the expansion of HIV-1 subtype BJ-B lineages in China. Bayesian skyline plot was estimated to reconstruct the demographic history of subtype BJ-B in China using BEAST 1.4. The *x*-axis is the time in units of years and the *y*-axis is equal to the effective population size. The *thick solid line* is the mean estimates and the 95% HPD credible region is shown by *blue* areas.