**Table S2. Intra-genotype pairwise nucleotide distances of CRF58\_01B and its putative parental reference strains (CRF01\_AE and B/B') (supplementary of Figure 1)**

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| HIV-1 genotypes/ Sub-region trees | I  (*gag*) | II  (*pol*) | III  (*pol*) | IV  (*pol*) | V  (*pol*-*env*) | VI  (*env*) | VII  (*env-*3' LTR) |
| 1. CRF01\_AE | 0.030±0.002 |  | 0.037±0.003 |  | 0.064±0.002 |  | 0.071±0.005 |
| 2. B/B' |  | 0.057±0.004 |  | 0.069±0.008 |  | 0.077±0.006 |  |
| 3. CRF58\_01B | 0.029±0.003 | 0.035±0.005 | 0.019±0.004 | 0.038±0.009 | 0.053±0.003 | 0.027±0.005 | 0.070±0.006 |