

Table S2. Whole AIHV-1 genome sequencing data

Description	Total reads (no.)	Target reads (no.)	Target reads (%)	Coverage (reads/nt)
A7 ^{STOP-39} (BAC plasmid)	1,352,076	1,333,326	99	1,366
A7 ^{STOP-207} (BAC plasmid)	1,336,896	1,322,936	99	1,355
A8 ^{STOP-159} (BAC plasmid)	1,516,540	1,482,871	98	1,519
A7 ^{STOP-207} (BAC ⁻ virus in BT cells)	1,460,388	251,143	17	275
A8 ^{STOP-159} (BAC ⁻ virus in BT cells)	1,520,344	165,427	11	181