

SUPPLEMENTAL TABLE S2. LIST OF REFERENCE FOR CHINESE SEQUENCES

<i>Accession No.</i>	<i>Title</i>	<i>Journal</i>
JX112847, 49, 54, 55, 59 JX112796, 98, 99 JX112800-08	The rapidly expanding CRF01_AE epidemic in China is driven by multiple lineages of HIV-1 viruses introduced in the 1990s	AIDS 27 (11), 1793–1802 (2013)
KM217833–KM217835 KM217837–KM217855	Comprehensive characterization of the transmitted/founder <i>env</i> genes from a single MSM cohort in China	J Acquir Immune Defic Syndr 69 (4), 403–412 (2015)
JX960603–JX960626 JX960628–JX960634	Reconstituting the epidemic history of CRF01_AE among MSM in Liaoning, northeastern China: implication in expanding MSM epidemic in China	J Virol 86 (22), 12402–12406 (2012)
KC183777	New emerging recombinant HIV-1 strains and close transmission linkage of HIV-1 strains in the Chinese MSM population indicate a new epidemic risk	PLoS One 8 (1), E54322 (2013)
KJ778895 KJ778896	Near full-length genomic characterization of a novel HIV type 1 CRF01_AE/07_BC recombinant in men who have sex with men from Shannxi, China	Unpublished