

## Electronic Supplementary Material

### Isolation, characterization, and phylogenetic analysis of two new Crimean-Congo hemorrhagic fever virus strains from the northern region of Xinjiang province, China

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**Table S1** Primers used in this study

Primer sets	Primers	Sequences (5'-3')	Positions (nt)	Length of Products
1	XHF-Fx-Q1/Q2	AGTGGAGGAARGACATAGG	450–468	246 bp
	XHF-Rx-Q2	CCATKATGTACTTGCCTTTG	670–689	
2	M13-F*	TGTAAAACGACGGCCAGT	2976–2993	348 bp
	XHF-Rx-Q2	CCATKATGTACTTGCCTTTG	670–689	
3	XHF-Fx-Q1/Q2	AGTGGAGGAARGACATAGG	450–468	176 bp
	XHF-Rx-Q1	TTTCATCRCCACCTCTGTT	599–617	

\* The nt position of primer M13-F is indicated in reference to the sequence of pGEM®-Teasy vector.