

1 **S2 Table. Percentage similarity based on L segment of HTNV nucleotide and amino acid sequences**
 2 **between HTNV from Jeju Island and representative rodent-borne orthohantaviruses.**

Sample (Host)	Homology (%)							
	HTNV		HTNV		SOOV		SEOV	
	76-118		HV004		SC1		80-39	
	<i>(A. agrarius)</i>		<i>(A. agrarius)</i>		<i>(A. peninsulae)</i>		<i>(R. norvegicus)</i>	
	nt (%)	aa (%)	nt (%)	aa (%)	nt (%)	aa (%)	nt (%)	aa (%)
Ac19-6	82.6	96.1	81.4	95.4	81.0	95.2	74.5	83.8
Ac20-5	82.4	96.8	81.6	96.1	81.3	95.9	74.6	84.6
Ac20-6	82.2	97.1	81.6	96.2	81.3	96.0	74.7	84.8
Ac20-30	82.9	97.2	82.0	96.4	81.6	96.2	74.8	84.8
Ac20-31	83.1	97.0	82.1	96.2	81.4	95.8	74.7	84.6
Ac20-32	82.9	97.0	81.8	96.0	81.1	95.8	74.8	84.8

3 HTNV, Hantaan virus; SOOV, Soochong virus; SEOV, Seoul virus; nt, nucleotide; aa, amino acid.