

1 **S4 Table. Percentage similarity based on S 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	85.4	96.3	85.3	97.2	84.3	96.3	69.6	82.3
Ac20-5	84.8	96.5	85.0	97.4	84.5	96.5	69.7	82.5
Ac20-6	84.9	96.5	85.1	97.4	84.6	96.5	69.7	82.5
Ac20-30	85.4	96.3	85.3	97.2	84.6	96.3	69.6	82.3
Ac20-31	85.1	96.5	85.1	97.0	84.8	96.0	69.8	82.5
Ac20-32	85.1	96.7	85.1	97.2	84.5	96.3	69.6	82.8

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