

S1 Table. Accession numbers and protein boundaries of the hepaciviruses used in this study.

Virus	Accession number	NS2 coding sequence	NS3_N coding sequence
BHV PDB-452	KC796090	nt 2504-3154	nt 3155-3796
GBV-B	AY243572	nt 2642-3265	nt 3266-3835
GHV-1 BWC08	KC551800	nt 2416-3051	nt 3052-3699
HCV H77 (1a)	NC_004102	nt 2769-3419	-
HCV Con1 (1b)	AJ238799	nt 2769-3419	-
HCV J6 (2a)	AF177036	nt 2780-3430	-
HCV JFH1 (2a)	AB047639	nt 2780-3430	nt 3431-4069
NPHV H3-011	JQ434008	nt 2758-3408	nt 3409-4047
RHV NLR07-oct70	KC411784	nt 2535-3131	nt 3132-3770

Boundaries of NS2 and NS3_N coding sequences used in constructs are indicated by their nucleotide (nt) positions within respective cDNAs.