Article DOI: https://doi.org/10.3201/eid2906.230056

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Novel Orthonairovirus Isolated from Ticks near China–North Korea Border

Appendix

	5' terminus	3' terminus
CCHFV-L	TCTCAAAGATATCAATCCCCCCGTT-	CTCTTTCTTAGCTATATCTTTGAGA
CCHFV-M	TCTCAAAGAAATACTTGCGGCACG-	GCGTGCCGCCACTATATCTTTGAGA
CCHFV-S	TCTCAAAGAAACACGTGCCGCTTA-	CTGTGCGGCAACGGTA TCTTTGAGA
HUGV-L	TCTCAAAGATATAAATCCCTACACC-	GGTTTAGGGAACACTT TCTTTGAGA
HUGV-M	TCTCAAAGAAAGACCTGCAGCAAC	GTTAGCTGCCACAATA TCTTTGAGA
HUGV-S	TCTCAAAGAAAGACGTGCCTATCC-	GTTAGCGGCAACAATATCTTTGAGA
ERVEV-L	TCTCAAAGAAAGCAATCCCCCCA	TAGGGGGAATACTATCTTTGAGA
ERVEV-M	TCTCAAAGAAAGACTAGCGGCAA-	GTATGCCGCTCCTATA TCTTTGAGA
ERVEV-S	TCTCAAAGAAAGTTGTGCTGTTACT-	GTATGCAGCAACACTATCTTTGAGA
TDYV-L	CACACCCA AATACATAGAACCAGG-	CCGGTGTTTATGGTAT <mark>TTGGGTGTG</mark>
TDYV-M	CACACCAAGCATTATAAACCAAGTT.	CAGGTGGATAAGATTTCTTGGTGTG
TDYV-S	CACACCCA AACTTTACACTTTAGG-	CCTTGAGCAATTTGCTTTGGGTGTG
TTV-L	TCTCAAAGATATATCCTGCACAC-	GCTGGCACAAATCGGTGAAATTGT
TTV-M	CTGCAGCACCAAAAGCCTTTCA	GCCGTGAAAAGAAAGAAATACA
TTV-S	AACGTGCTGCACACCAATAGCATTC-	ATCAGTTTACTACTGGTGTAAGTT
HTV-L	AAGATATATCCTGCACACCCAAA	GTGACCAACTCTTCTTGGTCTGATA
HTV-M	AAACAGTGTGAAGGCAATGATGAG-	AAACAAACAAAGAGAAAAA
HTV-S	CTGCACACCAAACCTAAAGCAAC	ATATAAAGAGATCGGAAGAGCGTCG
WTV-L	CCTAACACCACTTAACATCTGCCAA-	TAGGGTTARGGGTGTGCAGGAACA
WTV-M	CCCTACTAAAGGCTAAAGGTTAGCG-	TGGTGTTAGGTGTATTGGTGTGCTG
WTV-S	ATCACCTACATCGAATACCCATCCC-	GTAAGTTAAAGGGTGTGCAGCAACA
SGLV-L	TCTCAAAGATATATCCTGCACA-	GTGTGGGGAAGGTTGA <mark>TCCCATGT</mark>
SGLV-M	ACATGGGATAGTAACCTGTGCTAG-	CCTGTCTCGACCATGCCCCCCATGT
SGLV-S	TCTCAAAGAAAACGTGCTGCACAC	AGAAACAAAACAAATTCCCCATGT
ATV-L	GATATATCCTGCACACCCAAAC	AGCTGCTGTGGAGTCTGGTGACCT
ATV-M	TGTCAGCATCGAAGGAGGGAACA —	AAGGTCCCGCCTCTGCAACTGCCT
ATV-S	TCTCAAAGAAAACGTGCTGCACA	GTGCAGCAACAATATCTTTGAGA

Appendix Figure. Terminal sequences of ATV and other orthonairoviruses. Typical terminal sequences of orthonairoviruses are highlighted in yellow and putative terminal sequences in red. L, Large segment;

M, medium segment; S, small segment; CCHFV, Crimean-Congo hemorrhagic fever virus (L, NC_005301; M, NC_005300; S, NC_005302); HUGV, Hughes virus (L, NC_040512; M, NC_040513; S, NC_040514); ERVEV, Erve virus (L, JF911697; M, JF911698; S, JF911699); TDYV, Tamdy virus (L, MK757580; M, MK757581; S, MK757582); TTV, Tacheng tick virus 1 (L, NC_031284; M, NC_031285; S, NC_031286); HTV, Huangpi tick virus 1 (L, NC_031135; M, NC_031136; S, NC_031137); WTV, Wenzhou tick virus (L, NC_031291; M, NC_031288; S, NC_031289); SGLV, Songling virus (L, MT328776; M, MT328775; S, MT328777); ATV, Antu virus.