

COVID-19 is an emerging, rapidly evolving situation.
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Get the latest research from NIH: <https://www.nih.gov/coronavirus>.

GenBank

Rhinolophus bat coronavirus BtCoV/4991 RNA-dependent RNA polymerase (RdRp) gene, partial cds

GenBank: KP876546.1

FASTA Graphics PopSet

Go to:

LOCUS KP876546 370 bp RNA linear VRL 02-MAR-2016
DEFINITION Rhinolophus bat coronavirus BtCoV/4991 RNA-dependent RNA polymerase (RdRp) gene, partial cds.
ACCESSION KP876546
VERSION KP876546.1
KEYWORDS .
SOURCE Rhinolophus bat coronavirus BtCoV/4991
ORGANISM [Rhinolophus bat coronavirus BtCoV/4991](#)
Viruses; Riboviria; Orthornavirae; Pisuviricota; Pisoniviricetes; Nidovirales; Coronidovirineae; Coronaviridae; Coronavirinae; unclassified Coronavirinae.
REFERENCE 1 (bases 1 to 370)
AUTHORS Ge,X.Y., Wang,N., Zhang,W., Hu,B., Li,B., Zhang,Y.Z., Zhou,J.H., Luo,C.M., Yang,X.L., Wu,L.J., Wang,B., Zhang,Y., Li,Z.X. and Shi,Z.L.
TITLE Coexistence of multiple coronaviruses in several bat colonies in an abandoned mineshaft
JOURNAL Virol Sin 31 (1), 31-40 (2016)
PUBMED [26920708](#)
REFERENCE 2 (bases 1 to 370)
AUTHORS Ge,X., Wang,N., Zhang,W., Hu,B., Li,B., Zhang,Y., Zhou,J., Luo,C., Yang,X., Wu,L., Wang,B., Zhang,Y., Li,Z. and Shi,Z.
TITLE Direct Submission
JOURNAL Submitted (04-MAR-2015) Key Laboratory of Special Pathogens and Biosafety, Wuhan Institute of Virology, Chinese Academy of Sciences, Xiaohongshan 44, Wuhan, Hubei 430071, China
COMMENT ##Assembly-Data-START##
Sequencing Technology :: Sanger dideoxy sequencing
##Assembly-Data-END##
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ORIGIN
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