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Nucleotide

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Bat SARS-like coronavirus Rs3367, complete genome

GenBank: KC881006.1

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Go to:

LOCUS KC881006 29792 bp ss-RNA linear VRL 22-NOV-2013

DEFINITION Bat SARS-like coronavirus Rs3367, complete genome.

ACCESSION KC881006

VERSION KC881006.1

KEYWORDS .

SOURCE Bat SARS-like coronavirus Rs3367

ORGANISM [Bat SARS-like coronavirus Rs3367](#)
Viruses; Riboviria; Orthornavirae; Pisuviricota; Pisoniviricetes;
Nidovirales; Cornidovirineae; Coronaviridae; Orthocoronavirinae;
Betacoronavirus; Sarbecovirus.

REFERENCE 1 (bases 1 to 29792)

AUTHORS Ge,X.Y., Li,J.L., Yang,X.L., Chmura,A.A., Zhu,G., Epstein,J.H.,
Mazet,J.K., Hu,B., Zhang,W., Peng,C., Zhang,Y.J., Luo,C.M., Tan,B.,
Wang,N., Zhu,Y., Crameri,G., Zhang,S.Y., Wang,L.F., Daszak,P. and
Shi,Z.L.

TITLE Isolation and characterization of a bat SARS-like coronavirus that
uses the ACE2 receptor

JOURNAL Nature 503, 535-538 (2013)

PUBMED [24172901](#)

REFERENCE 2 (bases 1 to 29792)

AUTHORS Ge,X.-Y., Li,J.-L., Yang,X.-L., Hu,B., Zhang,W., Peng,C.,
Zhang,Y.-J., Luo,C.-M., Tan,B., Wang,N., Zhu,Y., Crameri,G.,
Wang,L.-F., Daszak,P. and Shi,Z.-L.

TITLE Direct Submission

JOURNAL Submitted (08-APR-2013) Wuhan Institute of Virology, Chinese
Academy of Sciences, 44 Xiao Hong Shan, Wuhan, Hubei 430071, China

COMMENT ##Assembly-Data-START##
Sequencing Technology :: Sanger dideoxy sequencing
##Assembly-Data-END##

FEATURES

source Location/Qualifiers

1..29792

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[CDS](#)

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