

African horse sickness virus 7 strain 7/E.caballus-tc/NAM/2012/Windhoek-E120014_3
segment 7 VP7 (VP7) gene, complete cds
GenBank: KT030566.1

FASTA Graphics

Go to:

LOCUS KT030566 1167 bp RNA linear VRL 01-JAN-2016

DEFINITION African horse sickness virus 7 strain
7/E.caballus-tc/NAM/2012/Windhoek-E120014_3 segment 7 VP7 (VP7)
gene, complete cds.

ACCESSION KT030566

VERSION KT030566.1

DBLINK BioProject: PRJNA271179
BioSample: SAMN03764418

KEYWORDS .

SOURCE African horse sickness virus 7

ORGANISM African horse sickness virus 7
Viruses; Riboviria; Orthornavirae; Duplornaviricota;
Resentoviricetes; Reovirales; Sedoreoviridae; Orbivirus.

REFERENCE 1 (bases 1 to 1167)

AUTHORS Guthrie,A.J., Coetzee,P., Venter,E.H., Joone,C., Lourens,C.W.,
MacLachlan,N.J., Weyer,C.T. and le Grange,M.

TITLE Characterization of African horse sickness virus from field cases
in southern Africa

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 1167)

AUTHORS Guthrie,A.J., Coetzee,P., Venter,E.H., Joone,C., Lourens,C.W.,
MacLachlan,N.J., Weyer,C.T. and le Grange,M.

TITLE Direct Submission

JOURNAL Submitted (08-JUN-2015) Equine Research Centre, University of
Pretoria, Faculty of Veterinary Science, Soutpan Rd, Onderstepoort,
Gauteng 0110, South Africa

COMMENT GenBank Accession Numbers KT030560-KT030569 represent the complete
genome of African horse sickness virus 7 strain
7/E.caballus-tc/NAM/2012/Windhoek-E120014_3.

##Assembly-Data-START##

Assembly Method :: Geneious v. 8.1.5

Coverage :: 16608

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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ORIGIN

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