Avian orthoreovirus segment S4, complete genome

NCBI Reference Sequence: NC_015135.1

FASTA Graphics

Go to:

LOCUS NC_015135 1104 bp RNA linear VRL 13-AUG-2018

DEFINITION Avian orthoreovirus segment S4, complete genome.

ACCESSION NC_015135 REGION: 24..1127

VERSION NC_015135.1

DBLINK BioProject: PRJNA485481

KEYWORDS RefSeq.

SOURCE Avian orthoreovirus

ORGANISM Avian orthoreovirus

Viruses; Riboviria; Orthornavirae; Duplornaviricota; Resentoviricetes; Reovirales; Spinareoviridae; Orthoreovirus; Orthoreovirus avis.

REFERENCE 1

AUTHORS Banyai,K., Dandar,E., Dorsey,K.M., Mato,T. and Palya,V.

TITLE The genomic constellation of a novel avian orthoreovirus strain associated with runting-stunting syndrome in broilers

JOURNAL Virus Genes 42 (1), 82-89 (2011)

PUBMED 21116842

REFERENCE 2 (bases 1 to 1104)

CONSRTM NCBI Genome Project

TITLE Direct Submission

JOURNAL Submitted (11-FEB-2011) National Center for Biotechnology Information, NIH, Bethesda, MD 20894, USA

REFERENCE 3 (bases 1 to 1104)

AUTHORS Banyai,K.

TITLE Direct Submission

JOURNAL Submitted (29-SEP-2010) Banyai K., Hungarian Academy of Sciences, Veterinary Medical Research Institute, Hungaria krt. 21., Budapest, H-1143, HUNGARY

COMMENT PROVISIONAL REFSEQ: This record has not yet been subject to final NCBI review. The reference sequence is identical to FR694200.
COMPLETENESS: full length.

FEATURES Location/Qualifiers

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gene 1..1104

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ORIGIN

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