

Canine adenovirus type 2, complete genome

NCBI Reference Sequence: AC_000020.1

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LOCUS AC_000020 31323 bp DNA linear VRL 13-AUG-2018

DEFINITION Canine adenovirus type 2, complete genome.

ACCESSION AC_000020 BK000403

VERSION AC_000020.1

DBLINK BioProject: [PRJNA485481](#)

KEYWORDS RefSeq.

SOURCE Canine adenovirus 2

ORGANISM [Canine adenovirus 2](#)

Viruses; Varidnaviria; Bamfordvirae; Preplasmiviricota;
Tectiliviricetes; Rowavirales; Adenoviridae; Mastadenovirus;

Canine

mastadenovirus A.

REFERENCE 1 (bases 1 to 31323)

AUTHORS Davison,A.J., Benko,M. and Harrach,B.

TITLE Genetic content and evolution of adenoviruses

JOURNAL J. Gen. Virol. 84 (PT 11), 2895-2908 (2003)

PUBMED [14573794](#)

REFERENCE 2 (bases 1 to 31323)

AUTHORS Shibata,R., Shinagawa,M., Iida,Y. and Tsukiyama,T.

TITLE Nucleotide sequence of E1 region of canine adenovirus type 2

JOURNAL Virology 172 (2), 460-467 (1989)

PUBMED [2800332](#)

REFERENCE 3 (bases 1 to 31323)

CONSRTM NCBI Genome Project

TITLE Direct Submission

JOURNAL Submitted (01-DEC-2004) National Center for Biotechnology
Information, NIH, Bethesda, MD 20894, USA

REFERENCE 4 (bases 1 to 31323)

AUTHORS Davison,A.J.

TITLE Direct Submission

JOURNAL Submitted (03-MAY-2002) MRC Virology Unit, Church Street, Glasgow
G11 5JR, U.K.

COMMENT INFERRED [REFSEQ](#): This record is predicted by genome sequence
analysis and is not yet supported by experimental evidence. The
reference sequence was derived from [BK000403](#).
On Oct 3, 2005 this sequence version replaced [BK000403.1](#).
This record represents an alternative annotation for U77082. It is
included in the NCBI RefSeq collection as an alternative to the
reference sequence NC_001734.
COMPLETENESS: full length.

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	1-31323	U77082.1	1-31323	
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