Human adenovirus 5, complete genome

NCBI Reference Sequence: AC_000008.1

FASTA Graphics

Go to:

LOCUS AC_000008 35938 bp DNA linear VRL 13-AUG-2018

DEFINITION Human adenovirus 5, complete genome.

ACCESSION AC_000008

VERSION AC_000008.1

DBLINK BioProject: PRJNA485481

KEYWORDS RefSeq.

SOURCE Human adenovirus 5

ORGANISM Human adenovirus 5

Viruses; Varidnaviria; Bamfordvirae; Preplasmiviricota; Polisuviricotina; Pharingeaviricetes; Rowavirales; Adenoviridae; Mastadenovirus; Mastadenovirus caesari.

REFERENCE 1 (bases 1 to 35938)

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 35938)

AUTHORS Chroboczek,J., Bieber,F. and Jacrot,B.

TITLE The sequence of the genome of adenovirus type 5 and its comparison with the genome of adenovirus type 2

JOURNAL Virology 186 (1), 280-285 (1992)

PUBMED 1727603

REFERENCE 3 (bases 1 to 35938)

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 35938)

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 BK000408.

On Oct 3, 2005 this sequence version replaced BK000408.1.
This record represents an alternative annotation for M73260 M29978.
It is included in the NCBI RefSeq collection as an alternative to the reference sequence NC_001405.

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          COMPLETENESS: full length.
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