Human herpesvirus 5 strain Merlin, complete genome

NCBI Reference Sequence: NC_006273.2

FASTA Graphics

Go to:

LOCUS NC_006273 1686 bp DNA linear VRL 13-AUG-2018

DEFINITION Human herpesvirus 5 strain Merlin, complete genome.

ACCESSION NC_006273 REGION: complement(120656..122341)

VERSION NC_006273.2

DBLINK Project: 14559

BioProject: PRJNA485481

KEYWORDS RefSeq.

SOURCE Human betaherpesvirus 5 (HHV-5; HCMV)

ORGANISM Human betaherpesvirus 5

Viruses; Duplodnaviria; Heunggongvirae; Peploviricota;

Herviviricetes; Herpesvirales; Orthoherpesviridae;

Betaherpesvirinae; Cytomegalovirus; Cytomegalovirus humanbeta5.

REFERENCE 1 (bases 1 to 1686)

AUTHORS Gatherer,D., Seirafian,S., Cunningham,C., Holton,M., Dargan,D.J.,

Baluchova,K., Hector,R.D., Galbraith,J., Herzyk,P., Wilkinson,G.W.

and Davison,A.J.

TITLE High-resolution human cytomegalovirus transcriptome

JOURNAL Proc. Natl. Acad. Sci. U.S.A. 108 (49), 19755-19760 (2011)

PUBMED 22109557

REFERENCE 2 (bases 1 to 1686)

AUTHORS Dolan,A., Cunningham,C., Hector,R.D., Hassan-Walker,A.F., Lee,L.,

Addison,C., Dargan,D.J., McGeoch,D.J., Gatherer,D., Emery,V.C.,

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cell cytotoxicity; binds NKp30; DURP family"

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