

Japanese encephalitis virus, genome

NCBI Reference Sequence: NC_001437.1

FASTA Graphics

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LOCUS NC_001437 10976 bp ss-RNA linear VRL 24-MAY-2019
DEFINITION Japanese encephalitis virus, genome.
ACCESSION NC_001437
VERSION NC_001437.1
DBLINK BioProject: [PRJNA485481](#)
KEYWORDS RefSeq.
SOURCE Japanese encephalitis virus
ORGANISM [Japanese encephalitis virus](#)
Viruses; Riboviria; Orthornavirae; Kitrinoviricota;
Flasuviricetes;
Amarillovirales; Flaviviridae; Flavivirus.
REFERENCE 1 (bases 2478 to 3688)
AUTHORS Moureau,G., Cook,S., Lemey,P., Nougairede,A., Forrester,N.L.,
Khasnatinov,M., Charrel,R.N., Firth,A.E., Gould,E.A. and de
Lamballerie,X.
TITLE New insights into flavivirus evolution, taxonomy and biogeographic
history, extended by analysis of canonical and alternative coding
sequences
JOURNAL PLOS ONE 10 (2), E0117849 (2015)
PUBMED [25719412](#)
REMARK Publication Status: Online-Only
REFERENCE 2 (sites)
AUTHORS Melian,E.B., Hinzman,E., Nagasaki,T., Firth,A.E., Wills,N.M.,
Nouwens,A.S., Blitvich,B.J., Leung,J., Funk,A., Atkins,J.F.,
Hall,R. and Khromykh,A.A.
TITLE NS1' of flaviviruses in the Japanese encephalitis virus serogroup
is a product of ribosomal frameshifting and plays a role in viral
neuroinvasiveness
JOURNAL J. Virol. 84 (3), 1641-1647 (2010)
PUBMED [19906906](#)
REFERENCE 3 (bases 1 to 10976)
AUTHORS Sumiyoshi,H., Mori,C., Fuke,I., Morita,K., Kuhara,S., Kondou,J.,
Kikuchi,Y., Nagamatu,H. and Igarashi,A.
TITLE Complete nucleotide sequence of the Japanese encephalitis virus
genome RNA
JOURNAL Virology 161 (2), 497-510 (1987)
PUBMED [3686827](#)
REFERENCE 4 (bases 1 to 10976)
CONSRTM NCBI Genome Project
TITLE Direct Submission

JOURNAL Submitted (01-AUG-2000) National Center for Biotechnology Information, NIH, Bethesda, MD 20894, USA

COMMENT REVIEWED [REFSEQ](#): This record has been curated by NCBI staff. The reference sequence was derived from [M18370](#).
Truncated version of JEV polyprotein and NS1' protein have been added and annotated with the kind help of Dr. David Karlin(Department of Zoology, University of Oxford, UK) and Dr. Andrew Firth (Department of Pathology, University of Cambridge, UK).
Draft entry and computer-readable sequence for [1] kindly submitted
by H.Sumiyoshi, 31-AUG-1989.
COMPLETENESS: not full length.

FEATURES Location/Qualifiers

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