

Murine rotavirus strain EB-C8/G16P[16] VP1 (VP1) gene, complete cds

GenBank: KJ477127.1

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LOCUS KJ477127 3302 bp RNA linear VRL 29-MAY-2014
DEFINITION Murine rotavirus strain EB-C8/G16P[16] VP1 (VP1) gene, complete
cds.
ACCESSION KJ477127
VERSION KJ477127.1
KEYWORDS .
SOURCE Murine rotavirus
ORGANISM [Murine rotavirus](#)
Viruses; Riboviria; Orthornavirae; Duplornaviricota;
Resentoviricetes; Reovirales; Sedoreoviridae; Rotavirus.
REFERENCE 1 (bases 1 to 3302)
AUTHORS Tsugawa,T., Tatsumi,M. and Tsutsumi,H.
TITLE Virulence-associated genome mutations of murine rotavirus
identified by alternating serial passages in mice and cell
cultures
JOURNAL J. Virol. 88 (10), 5543-5558 (2014)
PUBMED [24599996](#)
REFERENCE 2 (bases 1 to 3302)
AUTHORS Tsugawa,T., Tatsumi,M. and Hoshino,Y.
TITLE Direct Submission
JOURNAL Submitted (21-FEB-2014) Rotavirus Vaccine Development Section,
Laboratory of Infectious Disease, National Institute of Allergy
and
Infectious Diseases, National Institutes of Health, Building 50,
Room 6308, 50 South Drive, MSC 8026, Bethesda, MD 20892-8026, USA
COMMENT GenBank Accession Numbers KJ477127-KJ477137 represent sequences
from the 11 segments of Murine rotavirus EB-C8/G16P[16].

##Assembly-Data-START##
Assembly Method :: Sequencher v. 5.2
Sequencing Technology :: dye-terminator
##Assembly-Data-END##
FEATURES Location/Qualifiers
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ORIGIN

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