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Nucleotide

GenBank

Zika virus strain H/PF/2013 polyprotein gene, complete cds

GenBank: KJ776791.1

This sequence has been updated. [See current version.](#)

[FASTA](#) [Graphics](#)

Go to:

LOCUS KJ776791 10617 bp RNA linear VRL 13-JUN-2014
DEFINITION Zika virus strain H/PF/2013 polyprotein gene, complete cds.
ACCESSION KJ776791
VERSION KJ776791.1
KEYWORDS .
SOURCE Zika virus
ORGANISM [Zika virus](#)
Viruses; ssRNA viruses; ssRNA positive-strand viruses, no DNA stage; Flaviviridae; Flavivirus.
REFERENCE 1 (bases 1 to 10617)
AUTHORS Baronti,C., Piorkowski,G., Charrel,R.N., Boubis,L., Leparc-Goffart,I. and de Lamballerie,X.
TITLE Complete coding sequence of zika virus from a French polynesia outbreak in 2013
JOURNAL Genome Announc 2 (3), e00500-14 (2014)
PUBMED [24903869](#)
REMARK Publication Status: Online-Only
REFERENCE 2 (bases 1 to 10617)
AUTHORS Baronti,C. and Piorkowski,G.
TITLE Direct Submission
JOURNAL Submitted (30-APR-2014) UMR190-Unite des Virus Emergents, IRD, 27 bd Jean Moulin, Marseille, Bouches du Rhone 13005, France
COMMENT [WARNING] On Aug 31, 2016 this sequence was replaced by [KJ776791.2](#).

##Assembly-Data-START##
Assembly Method :: CLC genomics workbench v. 7.0.3
Assembly Name :: Zika PF
Sequencing Technology :: IonTorrent
##Assembly-Data-END##

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[CDS](#)

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