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# Zika virus strain H/PF/2013, complete genome

GenBank: KJ776791.2

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LOCUS KJ776791 10807 bp RNA linear VRL 31-AUG-2016  
DEFINITION Zika virus strain H/PF/2013, complete genome.  
ACCESSION KJ776791  
VERSION KJ776791.2  
KEYWORDS .  
SOURCE Zika virus  
ORGANISM [Zika virus](#)  
Viruses; Riboviria; Orthornavirae; Kitrinoviricota; Flasuviricetes; Amarillovirales; Flaviviridae; Flavivirus.  
REFERENCE 1 (bases 1 to 10807)  
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 10807)  
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  
REFERENCE 3 (bases 1 to 10807)  
AUTHORS Baronti,C. and Piorkowski,G.  
TITLE Direct Submission  
JOURNAL Submitted (31-AUG-2016) UMR190-Unite des Virus Emergents, IRD, 27 bd Jean Moulin, Marseille, Bouches du Rhone 13005, France  
REMARK Sequence update by submitter  
COMMENT On Aug 31, 2016 this sequence version replaced [KJ776791.1](#).

##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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