

# Middle East respiratory syndrome-related coronavirus isolate Merscov/Egypt/Camel/AHRI-FAO-1/2018, complete genome

GenBank: MK967708.1

[FASTA Graphics](#)

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LOCUS MK967708 30106 bp RNA linear VRL 15-DEC-2019

DEFINITION Middle East respiratory syndrome-related coronavirus isolate Merscov/Egypt/Camel/AHRI-FAO-1/2018, complete genome.

ACCESSION MK967708

VERSION MK967708.1

KEYWORDS .

SOURCE Middle East respiratory syndrome-related coronavirus (MERS-CoV)  
ORGANISM [Middle East respiratory syndrome-related coronavirus](#)  
Viruses; Riboviria; Orthornavirae; Pisuviricota; Pisoniviricetes; Nidovirales; Cornidovirineae; Coronaviridae; Orthocoronavirinae; Betacoronavirus; Merbecovirus.

REFERENCE 1 (bases 1 to 30106)  
AUTHORS Shaheen,M.A., Hagag,N.M., Hanafy,S., Ibrahim,E., Mansour,A., Nubie,O., Lubroth,J., Von Dobschuetz,S. and Makonnen,Y.  
TITLE Genetic characterization of complete genome of Middle East respiratory syndrome-related coronavirus (MERS-Cov) recently circulated in Egypt  
JOURNAL Unpublished

REFERENCE 2 (bases 1 to 30106)  
AUTHORS Shaheen,M.A., Hagag,N.M., Hanafy,S., Ibrahim,E., Mansour,A., Nubie,O., Lubroth,J., Von Dobschuetz,S. and Makonnen,Y.  
TITLE Direct Submission  
JOURNAL Submitted (22-MAY-2019) Genome, Animal Health Research institute, nadi el said, Dokki, Giza 12611, Egypt

COMMENT ##Assembly-Data-START##  
Assembly Method :: CLC genomic workbench v. 12  
Assembly Name :: MERS Cov  
Coverage :: 95%  
Sequencing Technology :: 454  
##Assembly-Data-END##

FEATURES Location/Qualifiers  
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