

Human coronavirus OC43 strain ATCC VR-759, complete genome

NCBI Reference Sequence: NC_006213.1

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LOCUS NC_006213 30741 bp RNA linear VRL 16-SEP-2020

DEFINITION Human coronavirus OC43 strain ATCC VR-759, complete genome.

ACCESSION NC_006213

VERSION NC_006213.1

DBLINK BioProject: [PRJNA485481](#)

KEYWORDS RefSeq.

SOURCE Human coronavirus OC43 (HCoV-OC43)

ORGANISM [Human coronavirus OC43](#)

Viruses; Riboviria; Orthornavirae; Pisuviricota; Pisoniviricetes; Nidovirales; Cornidovirineae; Coronaviridae; Orthocoronavirinae; Betacoronavirus; Embecovirus.

REFERENCE 1 (bases 1 to 30741)

AUTHORS St-Jean,J.R., Jacomy,H., Desforges,M., Vabret,A., Freymuth,F. and Talbot,P.J.

TITLE Human Respiratory Coronavirus OC43: Genetic Stability and Neuroinvasion

JOURNAL J. Virol. 78 (16), 8824-8834 (2004)

PUBMED [15280490](#)

REFERENCE 2 (bases 1 to 30741)

CONSRM NCBI Genome Project

TITLE Direct Submission

JOURNAL Submitted (21-FEB-2019) National Center for Biotechnology Information, NIH, Bethesda, MD 20894, USA

REFERENCE 3 (bases 1 to 30741)

AUTHORS St-Jean,J.R. and Talbot,P.J.

TITLE Direct Submission

JOURNAL Submitted (29-MAR-2004) INRS-Institut Armand-Frappier, 531 boul. des Prairies, Laval, Qc H7V 1B7, Canada

COMMENT REVIEWED [REFSEQ](#): This record has been curated by NCBI staff. The reference sequence is identical to [AY585228](#).
Annotation was based on information found in PMID: 16537637, PMID: 27381385 and the annotation of KF530089.1.
COMPLETENESS: full length.

FEATURES Location/Qualifiers

source 1..30741

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