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COVID-19 is an emerging, rapidly evolving situation.
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Get the latest research from NIH: <https://www.nih.gov/coronavirus>.

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NCBI Virus

Retrieve, view, and download SARS-CoV-2 coronavirus genomic and protein sequences.

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Severe acute respiratory syndrome coronavirus 2 isolate Wuhan-Hu-1, complete genome

GenBank: MN908947.3

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COMMENT

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source

5'UTR

gene

CDS

MN908947

29903 bp ss-RNA linear

VRL 18-MAR-2020

Severe acute respiratory syndrome coronavirus 2 isolate Wuhan-Hu-1, complete genome.

MN908947

MN908947.3

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Severe acute respiratory syndrome coronavirus 2 (SARS-CoV2)

[Severe acute respiratory syndrome coronavirus 2](#)

Viruses; Riboviria; Nidovirales; Coronavirineae; Coronaviridae; Orthocoronavirinae; Betacoronavirus; Sarbecovirus.

1 (bases 1 to 29903)

Wu,F., Zhao,S., Yu,B., Chen,Y.M., Wang,W., Song,Z.G., Hu,Y., Tao,Z.W., Tian,J.H., Pei,Y.Y., Yuan,M.L., Zhang,Y.L., Dai,F.H., Liu,Y., Wang,Q.M., Zheng,J.J., Xu,L., Holmes,E.C. and Zhang,Y.Z.

A new coronavirus associated with human respiratory disease in China

Nature 579 (7798), 265-269 (2020)

[32015508](#)

2 (bases 1 to 29903)

Wu,F., Zhao,S., Yu,B., Chen,Y.-M., Wang,W., Hu,Y., Song,Z.-G., Tao,Z.-W., Tian,J.-H., Pei,Y.-Y., Yuan,M.L., Zhang,Y.-L., Dai,F.-H., Liu,Y., Wang,Q.-M., Zheng,J.-J., Xu,L., Holmes,E.C. and Zhang,Y.-Z.

Direct Submission

Submitted (05-JAN-2020) Shanghai Public Health Clinical Center & School of Public Health, Fudan University, Shanghai, China

On Jan 17, 2020 this sequence version replaced [MN908947.2](#).

##Assembly-Data-START##

Assembly Method :: Megahit v. V1.1.3

Sequencing Technology :: Illumina

##Assembly-Data-END##

Location/Qualifiers

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