

Genome Warehouse

GWH

Find an assembly



e.g., panda (https://bigd.big.ac.cn/search?dbld=gwh&q=panda); TKS (https://bigd.big.ac.cn/search?dbld=gwh&q=TKS); Manihot esculenta (https://bigd.big.ac.cn/search?dbld=gwh&q=Manihot esculenta); SAMC012932 (https://bigd.big.ac.cn/search?dbld=gwh&q=SAMC012932); PRJCA000437 (https://bigd.big.ac.cn/search?dbld=gwh&q=PRJCA000437); GWHAAAA00000000 (https://bigd.big.ac.cn/search?dbld=gwh&q=GWHAAAA00000000)

📌 Severe acute respiratory syndrome coronavirus 2 (https://bigd.big.ac.cn/gwh/Genome/210/show) / BetaCoV_HZ-1

Scientific Name : Severe acute respiratory syndrome coronavirus 2

Common Names : -

Bioproject : PRJCA002185 (https://bigd.big.ac.cn/bioproject/browse/PRJCA002185)

Biosample : SAMC133663 (https://bigd.big.ac.cn/biosample/browse/SAMC133663)

Accession No. : GWHABKS00000000

Submitter Organization : Microbiology Detection, Hangzhou center for disease control and prevention

Contacts : Yu Hua, indexofire@gmail.com

Released Date : 2020-02-08

Assembly Level : Completed genome

Genome Representation : Full Genome

Reference Genome : yes

Assembly method : SPAdes 3.11

Sequencing & coverage : Illumina Miseq 5037.0

Download :

DNA (ftp://download.big.ac.cn/gwh/Viruses/Severe_acute_respiratory_syndrome_coronavirus_2_BetaCoV_HZ-1_GWHABKS00000000)

GFF (ftp://download.big.ac.cn/gwh/Viruses/Severe_acute_respiratory_syndrome_coronavirus_2_BetaCoV_HZ-1_GWHABKS00000000)

RNA (ftp://download.big.ac.cn/gwh/Viruses/Severe_acute_respiratory_syndrome_coronavirus_2_BetaCoV_HZ-1_GWHABKS00000000)

Protein (ftp://download.big.ac.cn/gwh/Viruses/Severe_acute_respiratory_syndrome_coronavirus_2_BetaCoV_HZ-1_GWHABKS00000000)

Publication(s) : -

History

Statistics of Genome Assembly

Genome size (bp)	29,833
GC content	38.02%

Annotation of Whole Genome Assembly

	Protein	Total
Total	12	12

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