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Genome Warehouse

GWH	⬍

Find an assembly

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♟ *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_GWHABKS000000**
>
> **GFF ⊕ (ftp://download.big.ac.cn/gwh/Viruses/Severe_acute_respiratory_syndrome_coronavirus_2_BetaCoV_HZ-1_GWHABKS000000**
>
> **RNA ⊕ (ftp://download.big.ac.cn/gwh/Viruses/Severe_acute_respiratory_syndrome_coronavirus_2_BetaCoV_HZ-1_GWHABKS000000**
>
> **Protein ⊕ (ftp://download.big.ac.cn/gwh/Viruses/Severe_acute_respiratory_syndrome_coronavirus_2_BetaCoV_HZ-1_GWHABKS000**

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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