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<https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=2697049>



**Taxonomy Browser**

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## Severe acute respiratory syndrome coronavirus 2

Taxonomy ID: 2697049 (for references in articles please use NCBI:txid2697049)

current name

**Severe acute respiratory syndrome coronavirus 2**  
 acronym: **SARS-CoV-2**  
 equivalent: **2019-nCoV**

NCBI BLAST name: **viruses**  
 Rank: **no rank**  
 Genetic code: [Translation table 1 \(Standard\)](#)  
 Host: human/vertebrates

[Lineage](#) (full )  
[Viruses](#); [Riboviria](#); [Orthornavirae](#); [Pisuviricota](#); [Pisoniviricetes](#); [Nidovirales](#); [Coronavirineae](#); [Coronaviridae](#); [Orthocoronavirinae](#); [Betacoronavirus](#); [Sarbecovirus](#); [Severe acute respiratory syndrome-related coronavirus](#)

Entrez records	
Database name	Direct links
<a href="#">Nucleotide</a>	<a href="#">6,735,594</a>
<a href="#">Protein</a>	<a href="#">39,147,346</a>
<a href="#">Structure</a>	<a href="#">2,914</a>
<a href="#">Genome</a>	<a href="#">1</a>
<a href="#">Popset</a>	<a href="#">75</a>
<a href="#">Conserved Domains</a>	<a href="#">2</a>
<a href="#">GEO Datasets</a>	<a href="#">1,489</a>
<a href="#">PubMed Central</a>	<a href="#">15,570</a>
<a href="#">Gene</a>	<a href="#">22</a>
<a href="#">SRA Experiments</a>	<a href="#">6,215,096</a>
<a href="#">Identical Protein Groups</a>	<a href="#">4,055,033</a>
<a href="#">BioProject</a>	<a href="#">994</a>
<a href="#">BioSample</a>	<a href="#">7,578,731</a>
<a href="#">Assembly</a>	<a href="#">116</a>
<a href="#">Genetic Testing Registry</a>	<a href="#">78</a>

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https://www.ncbi.nlm.nih.gov/sra/docs/submitbio/

An official website of the United States government [Here's how you know](#)

**NIH** National Library of Medicine  
National Center for Biotechnology Information

SRA SRA Advanced

Getting Started Submission Quick Start Search and Download SRA in the Cloud Archive Documentation

## Submit your project and biological samples

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- [BioSample submissions guidelines](#)
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- [Contact SRA staff](#)

### Introduction

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Registering project and biological samples at the NCBI BioProject and the BioSample databases is a prerequisite for any public SRA submission. The BioProject and BioSample databases store data that relate to organizational and biological aspects of sequencing experiments.

### Getting Started

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### Submitting Data

#### General

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### SRA Submission

- [Submitting](#)

NCBI Requires Bioproject and Sample Submissions for High Throughput Sequencing submissions (SRA)