**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/JPN/HiroC317c/2022, complete genome**

GenBank: OP024162.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OP024162.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OP024162.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OP024162.1" \l "goto2274373191_0)

LOCUS OP024162 29843 bp RNA linear VRL 21-JUL-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/JPN/HiroC317c/2022, complete genome.

ACCESSION OP024162

VERSION OP024162.1

KEYWORDS .

SOURCE Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)

ORGANISM [Severe acute respiratory syndrome coronavirus 2](https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=2697049)

Viruses; Riboviria; Orthornavirae; Pisuviricota; Pisoniviricetes;

Nidovirales; Cornidovirineae; Coronaviridae; Orthocoronavirinae;

Betacoronavirus; Sarbecovirus.

REFERENCE 1 (bases 1 to 29843)

AUTHORS Kurose,M., Kanda,M., Fukunaga,S., Nomura,T., Tahara,H., Ohge,H.,

Ohdan,H., Kubo,T., Kishita,E., Kuwabara,M., Takahashi,K., Tanaka,J.

and Sakaguchi,T.

TITLE Molecular epidemiology of SARS-CoV-2 in Hiroshima Prefecture in

Japan

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29843)

AUTHORS Nomura,T., Hirakawa,R., Morita,T., Kurose,M. and Sakaguchi,T.

TITLE Direct Submission

JOURNAL Submitted (21-JUL-2022) Department of Virology, Hiroshima

University, 1-2-3 Kasumi, Hiroshima 734-8551, Japan

COMMENT ##Assembly-Data-START##

Assembly Method :: One Codex v. Jun-2022

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

source 1..29843

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stem-loop 1"

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/OP024162.1?from=13477&to=13531) 13477..13531

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stem-loop 2"

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/product="ORF3a protein"

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/gene="ORF8"

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/gene="ORF8"

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SFYEDFLEYHDVRVVLDFI"

[gene](https://www.ncbi.nlm.nih.gov/nuccore/OP024162.1?from=28260&to=29510) 28260..29510

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/gene="N"

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[gene](https://www.ncbi.nlm.nih.gov/nuccore/OP024162.1?from=29535&to=29651) 29535..29651

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/note="Coronavirus 3' UTR pseudoknot stem-loop 1"

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/OP024162.1?from=29606&to=29634) 29606..29634

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/note="Coronavirus 3' UTR pseudoknot stem-loop 2"

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/note="Coronavirus 3' stem-loop II-like motif (s2m)"

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