**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/MWI/MLW000918/2022, complete genome**

GenBank: ON099844.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/ON099844.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/ON099844.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/ON099844.1" \l "goto2213892225_0)

LOCUS ON099844 29676 bp RNA linear VRL 29-MAR-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/MWI/MLW000918/2022, complete genome.

ACCESSION ON099844

VERSION ON099844.1

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 29676)

AUTHORS Feasey,N.

TITLE Direct Submission

JOURNAL Submitted (29-MAR-2022) Salmonella Group, Malawi Liverpool Wellcome

Clinical Research Program, Queen Elizabeth Central Hospital,

Blantyre, Malawi

COMMENT ##Assembly-Data-START##

Assembly Method :: ARTIC v. 1.2.1

Sequencing Technology :: Oxford Nanopore Technologies

##Assembly-Data-END##

FEATURES Location/Qualifiers

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stem-loop 1"

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/ON099844.1?from=13423&to=13477) 13423..13477

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NSVTSSIVITSGDGTTSPISEHDYQIGGYTEKWESGVKDCVVLHSYFTSDYYQLYSTQ

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LLVQ"

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/gene="ORF8"

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SFYEDFLEYHDVRVVLDFI"

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

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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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