**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/MAR/CNRST-HM36/2021, complete genome**

GenBank: OM011974.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OM011974.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OM011974.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OM011974.1" \l "goto2170127341_0)

LOCUS OM011974 29804 bp RNA linear VRL 25-DEC-2021

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/MAR/CNRST-HM36/2021, complete genome.

ACCESSION OM011974

VERSION OM011974.1

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 29804)

AUTHORS Chouati,T., Hemlali,M., melloul,M., Alaoui,S.A., Ghoulam,S.,

Ghamaz,H., Ouarab,M., Askandar,O., Belayachi,L., Touil,N., El

Mchichi,B., Elannaz,H., Laraqui,A., Elouennass,M., Ennibi,K. and El

Fahime,E.

TITLE Genome Sequence of Omicron variant detected in Morocco

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29804)

AUTHORS Chouati,T., Hemlali,M., melloul,M., Alaoui,S.A., Ghoulam,S.,

Ghamaz,H., Ouarab,M., Askandar,O., Belayachi,L., Touil,N., El

Mchichi,B., Elannaz,H., Laraqui,A., Elouennass,M., Ennibi,K. and El

Fahime,E.

TITLE Direct Submission

JOURNAL Submitted (25-DEC-2021) UATRS/FGP, CNRST, Angle Allal

ELfassi/Avenue des far, Hay Ryad, BP8027, RABAT, MOROCCO 10102,

Morocco

COMMENT ##Assembly-Data-START##

Assembly Method :: SPADES v. MAY-2018; IRMA Thermofisher v.

MARS-2021

Sequencing Technology :: IonTorrent

##Assembly-Data-END##

FEATURES Location/Qualifiers

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

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