Candidozyma auris isolate CMIL_01 small subunit ribosomal RNA gene, partial sequence; internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence

GenBank: MZ323988.1

FASTA Graphics

Go to:

LOCUS MZ323988 412 bp DNA linear PLN 08-JUN-2021

DEFINITION Candidozyma auris isolate CMIL_01 small subunit ribosomal RNA gene, partial sequence; internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence.

ACCESSION MZ323988

VERSION MZ323988.1

KEYWORDS .

SOURCE Candidozyma auris (Candida auris)

ORGANISM Candidozyma auris

Eukaryota; Fungi; Dikarya; Ascomycota; Saccharomycotina; Pichiomycetes; Metschnikowiaceae; Candidozyma.

REFERENCE 1 (bases 1 to 412)

AUTHORS Adhikary,P.C., Chowdhury,A.H., Rahman,T., Islam,M.N., Sadique,T., Sen,D., Tabassum,A., Halim,F., Chowdhury,A., Abdullah,A., Mondal,D. and Ahmed,D.

TITLE Direct Submission

JOURNAL Submitted (02-JUN-2021) Clinical Microbiology and Immunology, icddr, b, 68, Shaheed Tajuddin Ahmed Sarani, Mohakhali, Dhaka 1212, Bangladesh

COMMENT ##Assembly-Data-START##

Sequencing Technology :: Sanger dideoxy sequencing

##Assembly-Data-END##

FEATURES Location/Qualifiers

source 1..412

/organism="Candidozyma auris"

/mol_type="genomic DNA"

/isolate="CMIL_01"

/db_xref="taxon:498019"

misc RNA <1..>412

/note="contains small subunit ribosomal RNA, internal transcribed spacer 1, 5.8S ribosomal RNA, internal transcribed spacer 2, and large subunit ribosomal RNA"

ORIGIN

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     301 tcacaaaatt acagcttgca cgaaaaaaat ctacgctttt tttttcgttt tgttgtcgcc
     361 tcaaatcagg taggactacc cgctgaactt aagcatatca attaagcgga gg
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