

Coccidioides immitis 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: MK577426.1

[FASTA Graphics](#)

[Go to:](#)

LOCUS MK577426 595 bp DNA linear PLN 12-APR-2019
DEFINITION Coccidioides immitis 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 MK577426
VERSION MK577426.1
KEYWORDS .
SOURCE Coccidioides immitis
ORGANISM [Coccidioides immitis](#)
Eukaryota; Fungi; Dikarya; Ascomycota; Pezizomycotina; Eurotiomycetes; Eurotiomycetidae; Onygenales; Onygenaceae; Coccidioides.
REFERENCE 1 (bases 1 to 595)
AUTHORS Motter,A.N., Canteros,C.E. and Suarez-Alvarez,R.O.
TITLE Epidemiology of coccidioidomycosis in Argentina
JOURNAL Unpublished
REFERENCE 2 (bases 1 to 595)
AUTHORS Motter,A.N., Canteros,C.E. and Suarez-Alvarez,R.O.
TITLE Direct Submission
JOURNAL Submitted (28-FEB-2019) UOCCB, ANLIS Dr. Carlos G. Malbran, Av Velez Sarsfield 563, CABA 1281, Argentina
COMMENT ##Assembly-Data-START##
Sequencing Technology :: Sanger dideoxy sequencing
##Assembly-Data-END##
FEATURES Location/Qualifiers
source 1..595
/organism="Coccidioides immitis"
/mol_type="genomic DNA"
/strain="DMic 185677"
/host="Homo sapiens"
/db_xref="taxon:[5501](#)"
/country="Mexico"
[misc RNA](#) <1..>595
/note="contains small subunit ribosomal RNA, internal transcribed spacer 1, 5.8S ribosomal RNA, internal transcribed spacer 2, and large subunit ribosomal RNA"
ORIGIN

1 cttccgtagg tgaacctgcg gaaggatcat taaagtggcg tccggctgcg cacctcccc
61 gcggggggtcg cgcggtccgt acctcccacc cgtgtttact gaactattgt tgccttggca
121 ggectgccgg gcctctggct gccggggatc gccgccttg cgcggcgtcc cgggcgcgcg
181 cctgccagtg gatcaattga actcttatgt gaagattgtc agtctgagca tcatagcaaa
241 aatcaaacia aactttcaac aacggatctc ttggttccgg catcgatgaa gaacgcagcg
301 aaatgcgata agtaatgtga attgcagaat tccgtgaatc atcgaatctt tgaacgcaca
361 ttgcgccctc tggattccg gggggcatgc ctgttcgagc gtcattgcaa acccttcaag
421 cacggcttgt gtgttgggcc aacgtccccg cttgtgtgga cgggcctgaa atgcagtggc
481 ggcaccgagt tcttggtgtc tgagtgtatg ggaaatcaat tcatcgctca aagaccgat
541 cggggccgat cttttttttt ttttaatatc cggtttgacc tcggatcagg tagga

//