

Nucleotide ▼

[GenBank](#)

Explanatum explanatum isolate Mithun 5.8S ribosomal RNA gene, partial sequence; internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence

GenBank: KY068130.1

[FASTA](#) [Graphics](#)[Go to:](#)

LOCUS KY068130 484 bp DNA linear INV 03-APR-2017
 DEFINITION Explanatum explanatum isolate Mithun 5.8S ribosomal RNA gene, partial sequence; internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence.
 ACCESSION KY068130
 VERSION KY068130.1
 KEYWORDS .
 SOURCE Explanatum explanatum
 ORGANISM [Explanatum explanatum](#)
 Eukaryota; Metazoa; Platyhelminthes; Trematoda; Digenea; Plagiorchiida; Pronocephalata; Paramphistomoidea; Paramphistomidae; Explanatum.
 REFERENCE 1 (bases 1 to 484)
 AUTHORS Chamuah,J.K., Jacob,S.S., Sakhrie,A., Hazarika,S.B., Jamir,I.Y., Borkotoky,D., Dutta,P.R. and Raina,O.K.
 TITLE Direct Submission
 JOURNAL Submitted (01-NOV-2016) Veterinary Parasitology, ICAR- National Research Centre on Mithun, Nagaland, Jharnapani, Medziphema, Dimapur, Nagaland 797106, India
 COMMENT ##Assembly-Data-START##
 Sequencing Technology :: Sanger dideoxy sequencing
 ##Assembly-Data-END##
 FEATURES Location/Qualifiers
 source 1..484
 /organism="Explanatum explanatum"
 /mol_type="genomic DNA"
 /isolate="Mithun"
 /host="Bos frontalis (Mithun)"
 /db_xref="taxon:1224815"
 /dev_stage="adult parasite"
 /country="India"
[misc RNA](#) <1..>484
 /note="contains 5.8S ribosomal RNA, internal transcribed spacer 2, and large subunit ribosomal RNA"
 ORIGIN
 1 tgtgtcgtatg aagagcgcag ccaactgtgt gaattaatgt gaactgcata ctgctttgaa
 61 catcgacatc ttgaacgcac attgcggcca cgggttttcc tgtggccacg cctgtccgag
 121 ggtcggccta taaactatca cgacgccc aaagtcgtgg cttgggatct gccagctggc
 181 gtgatttcct ctgtggttcg ccatgtgagg tgccagatct atggcgtttt cctaattgct
 241 ccggacacaa ctgcgtcttg ctggtagcgc agacgagggt gtggcggtag agtcgtggct
 301 cagtgaactg taatggtagc acgctctgct gttgtgcctt tgttagtgta actggtttga
 361 gatgctattg ctgtccgtcc aatcatgata acctactatg gtgttctgtt acctgacctc
 421 ggatcagacg tgaatacccg ctgaacttaa gcatatcact aagcggagga aaagaaacta
 481 acca

//