

Dactylogyrus sphyrna 18S ribosomal RNA gene, partial sequence; internal transcribed spacer 1, complete sequence; and 5.8S ribosomal RNA gene, partial sequence

GenBank: KJ605446.1

[FASTA Graphics](#)

[Go to:](#)

LOCUS KJ605446 1046 bp DNA linear INV 01-MAY-2016
DEFINITION Dactylogyrus sphyrna 18S ribosomal RNA gene, partial sequence;
internal transcribed spacer 1, complete sequence; and 5.8S
ribosomal RNA gene, partial sequence.
ACCESSION KJ605446
VERSION KJ605446.1
KEYWORDS .
SOURCE Dactylogyrus sphyrna
ORGANISM [Dactylogyrus sphyrna](#)
Eukaryota; Metazoa; Spiralia; Lophotrochozoa; Platyhelminthes;
Monogenea; Monopisthocotylea; Dactylogyridea; Dactylogyridae;
Dactylogyrus.
REFERENCE 1 (bases 1 to 1046)
AUTHORS Hao,C. and Yue,C.
TITLE Direct Submission
JOURNAL Submitted (21-MAR-2014) College of Veterinary Medicine, Xinjiang
Agricultural University, No. 311 Nongda East Road, Urumqi,
Xinjiang
830052, China
COMMENT ##Assembly-Data-START##
Sequencing Technology :: Sanger dideoxy sequencing
##Assembly-Data-END##
FEATURES Location/Qualifiers
source 1..1046
/organism="Dactylogyrus sphyrna"
/mol_type="genomic DNA"
/db_xref="taxon:[231338](#)"
[rRNA](#) <1..514
/product="18S ribosomal RNA"
[misc RNA](#) 515..1010
/product="internal transcribed spacer 1"
[rRNA](#) 1011..>1046
/product="5.8S ribosomal RNA"
ORIGIN
1 ctaacctgct aaatagtagc attgcaatth ttatgtggac ctctgtcctt gtgcctctgg
61 tacttggata gaaccacata agcaggtgaa acttcttaga ggaactggcg ctcgaaaagg
121 cgcacgagaa agagcaataa caggtctgtg atgccctaag atgtccgggg ccgcacgcgt
181 gctacaatga cgatgtact gagcatgatt aacttgtccg aaaggatcgg tgaaactgtt
241 caatcatcgt cgtgactggg attgggggtt gcaattgtcc cccatgaacc aggaattcct

301 agtaagcaca agtcataagc ttgtgctgat tacgtccctg ccctttgtac acaccgcccg
361 tcgctactac cgattgaatg gtttagcaag gacattggat tgagcccttg cggtttcggc
421 tgcttggggtt tgagaagatg tccgaacttg atcatttaga ggaagtaaaa gtcgtaacaa
481 ggtttccgta ggtgaacctg cggaaggatc attatcaagt ttccaataca actctctgac
541 tatgccgcat gtatcatttt gtctgggctt gtatctgaat cccacagatc gttcagcttg
601 gtatgtgtgt gatgttagcg gtttgtgaat gagatccatg cccgggtaat atggcccact
661 gcatggatcg cctatgatac agaaatgtgc ctggcggttc ggattgtaac actcttacgg
721 tctgcgctta gccttcacaca tctgccgact ctgattggag cggcctatga gttgcgcgca
781 ggtggtaacg ttctgcgcgt atgctcttgg tcgtacccat gtaatgcatt cattcagtca
841 gagatccggt gcgttacaaa cattctccct tcgggtgagt cccatgtgta cttttcaaac
901 gttttaaaca atctggctaa gcgcagtgct gagtgtcgtt atgcaaatac tgacgaggtg
961 ctgttctggc tcggtgcgag ttgccttcgg gtgataagta ccaacctata caactgcaag
1021 tggtgatca ctcggtcgt gtgtcg

//