

Strongylus vulgaris isolate ERU-Svul-1 18S ribosomal RNA gene, partial sequence; internal transcribed spacer 1 and 5.8S ribosomal RNA gene, complete sequence; and internal transcribed spacer 2, partial sequence
GenBank: MF489225.1

FASTA Graphics

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LOCUS MF489225 778 bp DNA linear INV 04-JUN-2018

DEFINITION Strongylus vulgaris isolate ERU-Svul-1 18S ribosomal RNA gene, partial sequence; internal transcribed spacer 1 and 5.8S ribosomal RNA gene, complete sequence; and internal transcribed spacer 2, partial sequence.

ACCESSION MF489225

VERSION MF489225.1

KEYWORDS .

SOURCE Strongylus vulgaris

ORGANISM Strongylus vulgaris

Eukaryota; Metazoa; Ecdysozoa; Nematoda; Chromadorea; Strongylida; Strongyloidea; Strongylidae; Strongylinae; Strongylus.

REFERENCE 1 (bases 1 to 778)

AUTHORS Yildirim,A. and Bilgin,M.

TITLE Molecular prevalence and phylogenetic characterization of Strongylus vulgaris in horses from Nevsehir Province of Turkey

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 778)

AUTHORS Yildirim,A. and Bilgin,M.

TITLE Direct Submission

JOURNAL Submitted (14-JUL-2017) Parasitology Department, Erciyes University Faculty of Veterinary Medicine, Talas yolu, Melikgazi, Kayseri 38039, Turkey

FEATURES Location/Qualifiers

source 1..778
/organism="Strongylus vulgaris"
/mol_type="genomic DNA"
/isolate="ERU-Svul-1"
/isolation_source="faeces"
/host="Equus caballus"
/db_xref="taxon:40348"
/dev_stage="egg"
/country="Turkey"

rRNA <1..132
/product="18S ribosomal RNA"

misc_RNA 133..408

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rRNA      409..561
        /product="5.8S ribosomal RNA"
misc_RNA  562..>778
        /product="internal transcribed spacer 2"

ORIGIN
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  661 tattattcaa taatatacac agtatgtcat ggattattc tcactactta attgtttcgc
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