GenBank flat file:

LOCUS KJ149294 601 bp DNA linear INV 12-MAY-2014

DEFINITION Entamoeba chattoni isolate WOSMR1 small subunit ribosomal RNA gene,

 partial sequence.

ACCESSION KJ149294

VERSION KJ149294

KEYWORDS .

SOURCE Entamoeba chattoni

 ORGANISM Entamoeba chattoni

 Eukaryota; Amoebozoa; Archamoebae; Entamoebidae; Entamoeba.

REFERENCE 1 (bases 1 to 601)

 AUTHORS Elsheikha,H.M.

 TITLE Entamoeba species in captive primates in zoological gardens in the

 UK

 JOURNAL Unpublished

REFERENCE 2 (bases 1 to 601)

 AUTHORS Elsheikha,H.M.

 TITLE Direct Submission

 JOURNAL Submitted (09-JAN-2014) Veterinary Medicine and Science, University

 of Nottingham, College Road, Loughborough, Leicestershire LE12 5RD,

 United Kingdom

COMMENT ##Assembly-Data-START##

 Assembly Method :: Geneious Pro v. 5.4.6

 Sequencing Technology :: Illumina

 ##Assembly-Data-END##

FEATURES Location/Qualifiers

 source 1..601

 /organism="Entamoeba chattoni"

 /mol\_type="genomic DNA"

 /isolate="WOSMR1"

 /isolation\_source="host feces"

 /host="Leontopithecus chrysomelas (golden headed lion

 tamarin)"

 /db\_xref="taxon:110765"

 /country="United Kingdom"

 /collection\_date="17-Aug-2010"

 /collected\_by="Carl Regan"

 /identified\_by="Hany Elsheikha"

 rRNA <1..>601

 /product="small subunit ribosomal RNA"

ORIGIN

 1 atagcttttt gagaagaagg ttaaattgag aattgatgaa acttaaatgt ttttacaagt

 61 aactgtttta aatatctgac ctatcaactt gaaggtatga tagaggcata cccaagtgat

 121 aacgggtaac gagaaataag ggtttgattt cggagaggga gctttaaaaa tggctaccac

 181 ttctaaggaa ggcagcaggc gcgcaaatta cccactttta atttagagag gtagtgacga

 241 taattaatag gatttgtttt ataacaagtt caattgtaat gattagagtg taaaaaaatt

 301 taagaaatcg attggagggc aagtctggtg ccagcagccg cggtaattcc agctccaata

 361 gtgtatatta aagttgttgt gtttaaaaag ctcgtagtcg aattataggg tttttatgtt

 421 agtaattcta ttactaggat agaaaaagga gaaaggttat ttataatctt ttatttactt

 481 tgaaaaaaat agagtgttca aagcaaaagt caattaatgt ataatgaagc ataggataat

 541 aatgaggaga agaaattttt gagaacagta ttaaaaaggg aaattggggt aatttagaca

 601 a

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LOCUS KJ149295 604 bp DNA linear INV 12-MAY-2014

DEFINITION Entamoeba chattoni isolate WOSMR2 small subunit ribosomal RNA gene,

 partial sequence.

ACCESSION KJ149295

VERSION KJ149295

KEYWORDS .

SOURCE Entamoeba chattoni

 ORGANISM Entamoeba chattoni

 Eukaryota; Amoebozoa; Archamoebae; Entamoebidae; Entamoeba.

REFERENCE 1 (bases 1 to 604)

 AUTHORS Elsheikha,H.M.

 TITLE Entamoeba species in captive primates in zoological gardens in the

 UK

 JOURNAL Unpublished

REFERENCE 2 (bases 1 to 604)

 AUTHORS Elsheikha,H.M.

 TITLE Direct Submission

 JOURNAL Submitted (09-JAN-2014) Veterinary Medicine and Science, University

 of Nottingham, College Road, Loughborough, Leicestershire LE12 5RD,

 United Kingdom

COMMENT ##Assembly-Data-START##

 Assembly Method :: Geneious Pro v. 5.4.6

 Sequencing Technology :: Illumina

 ##Assembly-Data-END##

FEATURES Location/Qualifiers

 source 1..604

 /organism="Entamoeba chattoni"

 /mol\_type="genomic DNA"

 /isolate="WOSMR2"

 /isolation\_source="host feces"

 /host="Trachypithecus auratus auratus (Eastern Javan

 langur)"

 /db\_xref="taxon:110765"

 /country="United Kingdom"

 /collection\_date="28-Aug-2010"

 /collected\_by="Carl Regan"

 /identified\_by="Hany Elsheikha"

 rRNA <1..>604

 /product="small subunit ribosomal RNA"

ORIGIN

 1 gaatagcctt tttgagaaga aggttaaatt gagaattgat gaaacttaaa tgtttttaca

 61 agtaactgtt ttaaatatct gacctatcaa cttgaaggta tgatagaggc atacccaagt

 121 gataacgggt aacgagaaat aagggtttga tttcggagag ggagctttaa aaatggctac

 181 cacttctaag gaaggcagca ggcgcgcaaa ttacccactt ttaatttaga gaggtagtga

 241 cgataattaa taggatttgt tttataacaa gttcaattgt aatgattaga gtgtaaaaaa

 301 atttaagaaa tcgattggag ggcaagtctg gtgccagcag ccgcggtaat tccagctcca

 361 atagtgtata ttaaagttgt tgtgtttaaa aagctcgtag tcgaattata gggtttttat

 421 gttagtaatt ctattactag gatagaaaaa ggagaaaggt tatttataat cttttattta

 481 ctttgaaaaa aatagagtgt tcaaagcaaa agtcaattaa tgtataaaga agcataggat

 541 aataatgagg agaagaaatt tttgagaaca gtattaaaaa gggaaattgg ggtaatttag

 601 acaa

//

LOCUS KJ149296 900 bp DNA linear INV 12-MAY-2014

DEFINITION Entamoeba dispar isolate PRIR1 trnR-trnR intergenic spacer, partial

 sequence.

ACCESSION KJ149296

VERSION KJ149296

KEYWORDS .

SOURCE Entamoeba dispar

 ORGANISM Entamoeba dispar

 Eukaryota; Amoebozoa; Archamoebae; Entamoebidae; Entamoeba.

REFERENCE 1 (bases 1 to 900)

 AUTHORS Elsheikha,H.M.

 TITLE Entamoeba species in captive primates in zoological gardens in the

 UK

 JOURNAL Unpublished

REFERENCE 2 (bases 1 to 900)

 AUTHORS Elsheikha,H.M.

 TITLE Direct Submission

 JOURNAL Submitted (09-JAN-2014) Veterinary Medicine and Science, University

 of Nottingham, College Road, Loughborough, Leicestershire LE12 5RD,

 United Kingdom

COMMENT ##Assembly-Data-START##

 Assembly Method :: Geneious Pro v. 5.4.6

 Sequencing Technology :: Illumina

 ##Assembly-Data-END##

FEATURES Location/Qualifiers

 source 1..900

 /organism="Entamoeba dispar"

 /mol\_type="genomic DNA"

 /isolate="PRIR1"

 /isolation\_source="feces"

 /host="Trachypithecus auratus auratus (Eastern Javan

 langur)"

 /db\_xref="taxon:46681"

 /country="United Kingdom"

 /collection\_date="09-May-2011"

 /collected\_by="Carl Regan"

 /identified\_by="Hany Elsheikha"

 misc\_feature <1..>900

 /note="trnR-trnR intergenic spacer"

ORIGIN

 1 catatataga tctgtcttag ataaaatcat tatatttcat gagacaccta tcgtttagca

 61 tatcttttat ttttcggtct tttatatgtc tgtttctctg agagacaaag gcaatacctc

 121 actatgcaaa aagtcgactg gaggtccatc ttccgactgc ataataatag tatagtcgta

 181 tatccctata atttacaatt tccatactcc aatattacaa tctgcttaaa ttcgcaattt

 241 aagagagcag aggcgtgaat gttgaaaatt atgaggcgcc tttttatcat cggcactgtc

 301 ggtgcggcaa tatatctaaa tcccctatac ggacttatgc tgtacataat acatatcatt

 361 tcatctgtta tagtaggaat attattccga ttttacggca aaaaccgcca caactcaccg

 421 cccacaaaca taaacaccgc agagcttgaa ctgccacgtg ttattactac agctcttgaa

 481 accgcatcaa aaaatatatt aaccgtctgt ttttcgagtg tgttcaatat catcatcccc

 541 tatttgagca gcttcctata ctaaattcgc ccttgattga aatattttca gatccactaa

 601 aatattctcc gtgcttgtaa atgatgcatg ttcgctcgtg aaatatgtaa ccccaccctt

 661 tgagtgttcc cactcgtata actcctatta tttatgttca tatttttgaa ttaatttagt

 721 cgaatatttt ataaaattta ctatcccaga aagtgtgcta atatgttcat cctatataat

 781 catattaggt catcacgtta ttgtgttatt aggatattat tctataatga atgaatgtgt

 841 cttatatcat tatcccctaa tgttattatg ttcttatgag tgtgttcaat atcatcatcc //