Nucleotide •

The Nucleotide database will include EST and GSS sequences in early 2019. Read more.

GenBank

Bacillus cereus strain SMK86 16S ribosomal RNA gene, partial sequence

GenBank: MH511615.1 FASTA Graphics

```
Go to:
LOCUS
                                     726 bp
            MH511615
                                               DNA
                                                       linear BCT 27-JUN-2018
DEFINITION Bacillus cereus strain SMK86 16S ribosomal RNA gene, partial
            sequence.
ACCESSION
            MH511615
VERSION
            MH511615.1
KEYWORDS
SOURCE
            Bacillus cereus
 ORGANISM
            Bacillus cereus
            Bacteria; Firmicutes; Bacilli; Bacillales; Bacillaceae; Bacillus;
            Bacillus cereus group.
REFERENCE
            1 (bases 1 to 726)
 AUTHORS
            K,S., V,M. and D,K.
  TITLE
            Direct Submission
            Submitted (22-JUN-2018) Biotechnology, Adhiyamaan College of
 JOURNAL
            Engineering, MGR Nagar, Hosur, Tamilnadu 635130, India
COMMENT
            ##Assembly-Data-START##
            Sequencing Technology :: Sanger dideoxy sequencing
            ##Assembly-Data-END##
FEATURES
                     Location/Qualifiers
     source
                     1..726
                     /organism="Bacillus cereus"
                     /mol_type="genomic DNA"
                     /strain="SMK86"
                     /isolation_source="Waste water"
                     /db_xref="taxon: 1396"
     rRNA
                     <1...>726
                     /product="16S ribosomal RNA"
ORIGIN
        1 aggacatgaa tgcagtgcta gtgttagagg gtttccagcc ctttagtgct cgaagttaac
       61 gcattaagca ctccgcctgg ggagtacggc cgcaaggctg aaactcaaag gaattgacgg
      121 gggcccgcac aagcggtgga gcatgtggtt taattcgaag caacgcgaag aaccttacca
      181 ggtcttgaca tcctctgaca accctagaga tagggcttct ccttcgggag caaagtgaca
      241 ggtggtgcat ggttgtcgtc agctcgtgtc gtgagatgtt gggttaagtc ccgcaacgag
      301 cgcaaccctt gatcttagtt gccatcattt agttgggcac tctaaggtga ctgccggtga
      361 caaaccggag gaaggtgggg atgacgtcaa atcatcatgc cccttatgac ctgggctaca
      421 cacgtgctac aatggacggt acaaagagct gcaagaccgc gaggtggagc taatctcata
      481 aaaccgttct cagttcggat tgtaggctgc aactcgccta catgaagctg gaatcgctag
      541 taatcgcgga tcagcatgcc gcggtgaata cgttcccggg ccttgtacac accgcccgtc
      601 acaccacgag agtttgtaac acccgaagtc ggtggggtaa cctttttgaa gccagccgcc
      661 aaagggggga caaataattg gggtgaagtc gtaacaggtt aacccgaaaa cgattaattc
      721 aataat
```