Acinetobacter baumannii strain E0528 16S ribosomal RNA gene, partial sequence; 16S-23S ribosomal RNA intergenic spacer, complete sequence; and 23S ribosomal RNA gene, partial sequence

GenBank: KR732283.1

FASTA Graphics PopSet

Go to:

```
LOCUS       KR732283               765 bp    DNA     linear   BCT 25-OCT-2015
DEFINITION  Acinetobacter baumannii strain E0528 16S ribosomal RNA gene,
            partial sequence; 16S-23S ribosomal RNA intergenic spacer,
complete
            sequence; and 23S ribosomal RNA gene, partial sequence.
ACCESSION   KR732283
VERSION     KR732283.1
KEYWORDS    .
SOURCE      Acinetobacter baumannii
  ORGANISM  Acinetobacter baumannii
            Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
            Moraxellaceae; Acinetobacter; Acinetobacter
calcoaceticus/baumannii
            complex.
REFERENCE   1  (bases 1 to 765)
  AUTHORS   Lin,C.-H., Lee,M.-C., Tzen,J.T.C., Lee,H.-M., Chang,S.-M., Tu,W.-
C.
            and Lin,C.-F.
  TITLE     Efficacy of mastoparan-AF alone and in combination with clinically
            used antibiotics on nosocomial multidrug-resistant Acinetobacter
            baumannii
  JOURNAL   Unpublished
REFERENCE   2  (bases 1 to 765)
  AUTHORS   Lin,C.-H., Lee,M.-C., Tzen,J.T.C., Lee,H.-M., Chang,S.-M., Tu,W.-
C.
            and Lin,C.-F.
  TITLE     Direct Submission
  JOURNAL   Submitted (13-MAY-2015) Central Taiwan University of Science and
            Technology, Institute of Biotechnology, No. 666, Buzih Road,
Beitun
            District, Taichung City 40601, Taiwan
COMMENT     ##Assembly-Data-START##
            Sequencing Technology :: Sanger dideoxy sequencing
            ##Assembly-Data-END##
FEATURES             Location/Qualifiers
     source          1..765
                     /organism="Acinetobacter baumannii"
                     /mol_type="genomic DNA"
```

/strain="E0528"
/isolation_source="blood"
/db_xref="taxon:470"
/collection_date="12-Aug-2008"
misc_RNA <1..>765
/note="contains 16S ribosomal RNA, 16S-23S ribosomal RNA intergenic spacer, and 23S ribosomal RNA"

ORIGIN

```
      1 aagcggaggg cagggaaact ccttacgaaa gattgacgat tggtaagaat ccacaacaag
     61 ttgttcttca tagatgtatc tgagggtctg tagctcagtt ggttagagca cacgcttgat
    121 aagcgtgggg tcacaagttc aagtcttgtc agacccacca tgactttgac tggttgaagt
    181 tatagataaa agatacatga ttgatgatgt aagctgggga cttagcttag ttggtagagc
    241 gcctgctttg cacgcaggag gtcaggagtt cgactctcct agtctccacc agaacttaag
    301 ataagttcgg attacagaaa ttagtaaata aagattgaga tcttggttta ttaacttctg
    361 tgatttcatt atcacggtaa ttagtgtgat ctgacgaaga cacattaact cattaacaga
    421 ttggcaaaat tgagtctgaa ataaattgtt cactcaagag tttaggttaa gcaattaatc
    481 tagatgaatt gagaactagc aaattaactg aatcaagcgt tttggtatgt gaatttagat
    541 tgaagctgta cagtgcttaa gtgcacagtg ctctaaactg aaatgttgaa gttactaact
    601 tgtaggtaac atcgactgtt tggggttgta tagtcaagta attaagtgca tgtggtggat
    661 gccttggcag tcagaggcga tgaaagacgt gatagcctgc gaaaagctcc ggggaggcgg
    721 caaatatcct ttgatccgga gatttccgac gggggggaaa ccaaa
```