

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

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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; Moraxellales;
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JOURNAL Unpublished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##
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intergenic spacer, and 23S ribosomal RNA"
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ORIGIN

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121 aagcgtgggg tcacaagttc aagtcttgtc agacccacca tgactttgac tggttgaagt  
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661 gccttggcag tcagaggcga tgaaagacgt gatagcctgc gaaaagctcc ggggaggcgg  
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