

Nucleotide ▾

Display Settings: GenBank

Klebsiella pneumoniae metallo-beta-lactamase VIM-like gene, partial sequence

GenBank: HQ588906.1

[FASTA](#) [Graphics](#)[Go to:](#)

LOCUS HQ588906 892 bp DNA linear BCT 12-APR-2011
 DEFINITION Klebsiella pneumoniae metallo-beta-lactamase VIM-like gene, partial sequence.

ACCESSION HQ588906
 VERSION HQ588906.1 GI:328496247

KEYWORDS .
 SOURCE Klebsiella pneumoniae
 ORGANISM [Klebsiella pneumoniae](#)

Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae; Klebsiella.

REFERENCE 1 (bases 1 to 892)
 AUTHORS Parveen,M., Harish,B.N. and Parija,S.C.
 TITLE Direct Submission
 JOURNAL Submitted (07-OCT-2010) Department of Microbiology, Jawaharlal Institute of Postgraduate Medical Education and Research, Dhanvanthari Nagar, Pondicherry, Pondicherry 605006, India

COMMENT GenBank staff are unable to verify sequence and/or annotation provided by the submitter.

FEATURES
 Location/Qualifiers
 source 1..892
 /organism="Klebsiella pneumoniae"
 /mol_type="genomic DNA"
 /isolate="A"
 /isolation_source="blood"
 /db_xref="taxon:573"
 /country="India"
 misc_feature <1..408
 /note="similar to metallo-beta-lactamase VIM"

ORIGIN

```

1  tgagttgctt ttgattgata cgggagtaga gtggcatagc cctaccacc cgtggttctc
61  atcaatcaag cgtggggtgc gaaaaacaca gcggcacttc tcgcgagat tgagaagcaa
121 tttagcctgt aacgcgtgca gtctccacgc ctagcaccct attcgtactt tcatatcatc
181 ccgggcagtg tcttaacgac gaccgcgtcg gcggcgttga tgtccttcgg gcggctgggg
241 tggcaacgta cgcatacacc tcgacacgcc ggctagccga ggtagagggg aacgagattc
301 ccacgcactc tctagaagga ctctcatcga gcggggacgc agtgtccgaa gatccgaaga
361 tgtactcgct tcggtccagt agaactcttc tatcctgtgc ctgcgatac acttctcgcc
421 tcaatcactt ctactcgcgc tgctgaatca aatacagggg gtacgggtag cctcccagga
481 agtcaggccc actttccatg gtaaaccaat cgccccgagc ggatcttacc ccccaaaca
541 attaaggata gagcctgcct ccttaatatg accgcgggtg cggactcgta cccagcttat
601 gcttctgcta acaaggacca tccatatggc tcgcacgctc gccttcatac cgctagtggg
661 atttgctcag gctccctcac gtgttctacc cttcggcagt ccgcccctca gagtccgagc
721 ccggggcggc gtctcactct cgcgggccgt catcntgtcg tcgcattaga tcaaggccta
781 tgtcgcggta tgcctcataa ccacagaggg ggataagcat caaaaatcat ctcacactgc
841 attacgacat tagggaccta ttagtcctcc tcttatacct gttggtcaac gg

```

//