

Nucleotide

Display Settings: GenBank

**Klebsiella pneumoniae strain P3560 metallo-beta-lactamase VIM-like gene, partial sequence**

GenBank: HQ588905.1

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

LOCUS HQ588905 900 bp DNA linear BCT 12-APR-2011  
 DEFINITION Klebsiella pneumoniae strain P3560 metallo-beta-lactamase VIM-like gene, partial sequence.

ACCESSION HQ588905

VERSION HQ588905.1 GI:328496246

KEYWORDS .

SOURCE Klebsiella pneumoniae

ORGANISM [Klebsiella pneumoniae](#)  
 Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;  
 Enterobacteriaceae; Klebsiella.

REFERENCE 1 (bases 1 to 900)

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..900  
 /organism="Klebsiella pneumoniae"  
 /mol\_type="genomic DNA"  
 /strain="P3560"  
 /db\_xref="taxon:573"  
 /country="India"  
 /collection\_date="11-Feb-2009"

[misc feature](#) <1..319  
 /note="similar to metallo-beta-lactamase VIM"

ORIGIN

```

1  tgagttgctt ttgattgata cagcgtgggg tgcgaaaaac acagcggcac ttctcgcgga
61  gattgagatg caaatggac ttctgtaac gcgtgcagtc tccacgcact ttcattgacga
121  ccgctcggc ggcttgatg tccttcgggc ggctggggtg gcaacgtacg catcacgcgc
181  gacacgccgg ctaccgagc tagaggggaa cgagattccc acgcactctc tagaaggact
241  ctcatcgagc ggggacgcag tgcgcttcgg tccagtagaa ctcttctatc ctgtgcctgc
301  gcatacactt ctgcctgat caatcacttc tactcgcgct gctgaatcaa atatcatccc
361  gggcagtgtc ttaacacagg gggtaggggt agcctcccag gaagtcaggc ccactttcca
421  tggtaaacca atcgccccga gcggatctta ccccccaaaa caattaagga tagagcctgc
481  ctcttaata tgaccgcccgg tccggactcg taccagctt atgcttctc taacaaggac
541  catccatgat gctgggagta gattggcata cggctaccac ccggtggttc tcatcaatca
601  cgcacgctcg ccttcatcac gctagttgga tttgttcagg ctccctcagc tgttctacc
661  ttcggcagtc cgcgcctcag agtccgagcc cggggcggcg tctcactctc gcgggcccgc
721  atcntgtcgt cgcattagat catcatctat gtcgcggtat gcctcataac cacagatcct
781  acataagcat caaaaatcat ctcacactcc gaagatccga agatgtactt gcattacgac
841  attagggacc tattagtctt cctcttatac ctgttggtca acggctagca ccctattcgt

```

//