



XP-002474326

EBI EMBL Nucleotide Sequence Database Sequence Version Archive

2)

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F.A.Q.

Accession Number or Sequence Version: AF516751 ☐ case sensitive

Snapshot at day-month-year (e.g. 30-11-1998 or 30-NOV-1998)

AF516751

This flat file was issued: 09-MAR-2004 Rel: 78

```
ID AF516751 standard; genomic DNA; PRO; 888 BP.
XX
AC AF516751;
XX
SV AF516751.2
XX
DT 06-AUG-2002 (Rel. 72, Created)
DT 28-JUN-2003 (Rel. 76, Last updated, Version 2)
XX
DE Rhodoferax sp. P230 R-2,4-dichlorophenoxypropionate dioxygenase (rdpA)
DE gene, complete cds.
XX
KW
XX
OS Rhodoferax sp. P230
OC Bacteria; Proteobacteria; Betaproteobacteria; Burkholderiales;
OC Comamonadaceae; Rhodoferax.
XX
RN [1]
RP 1-888
RA Schleinitz K.M., Kleinstein S., Babel W.;
RT "Two novel genes, rdpA and sdpa, encode enantiospecific cleavage of
RT 2,4-dichlorophenoxypropionate in Delftia acidovorans MC1";
RL Unpublished.
XX
RN [2]
RP 1-888
RA Schleinitz K.M.;
RT ;
RL Submitted (31-MAY-2002) to the EMBL/GenBank/DDBJ databases.
RL Environmental Microbiology, Centre for Environmental Research,
RL Permoserstrasse 15, Leipzig 04318, Germany
XX
RN [3]
RC Sequence update by submitter
RP 1-888
RA Schleinitz K.M.;
RT ;
RL Submitted (25-JUN-2003) to the EMBL/GenBank/DDBJ databases.
RL Environmental Microbiology, Centre for Environmental Research,
RL Permoserstrasse 15, Leipzig 04318, Germany
XX
DR GOA; Q8KSC9.
DR TrEMBL; Q8KSC9; Q8KSC9.
XX
CC On Jun 26, 2003 this sequence version replaced gi:22087565.
XX
FH Key Location/Qualifiers
FH
FT source 1..888
FT /db_xref="taxon:103036"
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FT      similar to TauD and TfdA"
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FT      LDAFHTYQVIYFPGQAITNEQHIAFSRRFGPVDVPLLLKSIEGYPEVQMIRREANESGR
FT      VIGDDWHTDSTFLDAPPAAVVMRAIDVPEHGGDTGFLSMYTAWETLSPTMQATIEGLNV
FT      VHSATRVFGSLYQAQNRFRFSNTSVKVMVDVDAGDRETVHPLVVTHPGSGRKGLYVNVQVYC
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FT      YLRTTTVGGVRPAR"
XX
SQ

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Sequence 888 BP; 168 A; 307 C; 272 G; 141 T; 0 other;
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