| Nucleotide ▾ | PQ133629:PQ133630[accn] | ⊗ |

Summary   Sort by Default order

Items: 2

☐ 1. [Proteus sp. (in: enterobacteria) strain S_Y 16S ribosomal RNA gene, partial sequence](#)
464 bp linear DNA
Accession: PQ133630.1   GI: 2782796410

[GenBank](#)   [FASTA](#)   [Graphics](#)

☐ 2. [Proteus sp. (in: enterobacteria) strain S_C 16S ribosomal RNA gene, partial sequence](#)
451 bp linear DNA
Accession: PQ133629.1   GI: 2782796409

[GenBank](#)   [FASTA](#)   [Graphics](#)

Nucleotide ▼

GenBank

Proteus sp. (in: enterobacteria) strain S_C 16S ribosomal RNA gene, partial sequence

GenBank: PQ133629.1

FASTA   Graphics

Go to:

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VERSION     PQ133629.1
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  ORGANISM  Proteus sp. (in: enterobacteria)
            Bacteria; Pseudomonadota; Gammaproteobacteria; Enterobacterales;
            Morganellaceae; Proteus.
REFERENCE   1  (bases 1 to 451)
  AUTHORS   Khalid,K.M.
  TITLE     Microbiology Lab
  JOURNAL   Unpublished
REFERENCE   2  (bases 1 to 451)
  AUTHORS   Khalid,K.M.
  TITLE     Direct Submission
  JOURNAL   Submitted (02-AUG-2024) Biology, Soran University, Azadi Street,
            Soran 44008, Iraq
COMMENT     ##Assembly-Data-START##
            Sequencing Technology :: Sanger dideoxy sequencing
            ##Assembly-Data-END##
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Nucleotide ▾ [   ]

GenBank

Proteus sp. (in: enterobacteria) strain S_Y 16S ribosomal RNA gene, partial sequence

GenBank: PQ133630.1

FASTA   Graphics

Go to:

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VERSION     PQ133630.1
KEYWORDS    .
SOURCE      Proteus sp. (in: enterobacteria)
  ORGANISM  Proteus sp. (in: enterobacteria)
            Bacteria; Pseudomonadota; Gammaproteobacteria; Enterobacterales;
            Morganellaceae; Proteus.
REFERENCE   1  (bases 1 to 464)
  AUTHORS   Khalid,K.M.
  TITLE     Microbiology Lab
  JOURNAL   Unpublished
REFERENCE   2  (bases 1 to 464)
  AUTHORS   Khalid,K.M.
  TITLE     Direct Submission
  JOURNAL   Submitted (02-AUG-2024) Biology, Soran University, Azadi Street,
            Soran 44008, Iraq
COMMENT     ##Assembly-Data-START##
            Sequencing Technology :: Sanger dideoxy sequencing
            ##Assembly-Data-END##
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                     /collection_date="18-May-2024"
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      301 agatacaaag agaagcgacc tcgcgagagc aagcggaact cataaagtct gtcgtagtcc
      361 ggattggagt ctgcaactcg actccatgaa gtcggaatcg ctagtaatcg tagatcagaa
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