Nucleotide 🗸

## COVID-19 is an emerging, rapidly evolving situation.

Public health information (CDC)

Research information (NIH)

SARS-CoV-2 data (NCBI)

Prevention and treatment information (HHS)

GenBank

## Mus musculus 7S RNA 2 (Rn7s2), small cytoplasmic RNA

NCBI Reference Sequence: NR\_052008.1

FASTA Graphics

exon

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Go to:
LOCUS
            NR 052008
                                      299 bp
                                                RNA
                                                         linear
                                                                  ROD 21-OCT-2020
DEFINITION
            Mus musculus 7S RNA 2 (Rn7s2), small cytoplasmic RNA.
            NR 052008
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VERSION
            NR_052008.1
KEYWORDS
            RefSea.
SOURCE
            Mus musculus (house mouse)
  ORGANISM
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            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;
            Muroidea; Muridae; Murinae; Mus; Mus.
REFERENCE
            1 (bases 1 to 299)
  AUTHORS
            Wilson CL, Heppner KJ, Rudolph LA and Matrisian LM.
  TITLE
            The metalloproteinase matrilysin is preferentially expressed by
            epithelial cells in a tissue-restricted pattern in the mouse
  JOURNAL
            Mol Biol Cell 6 (7), 851-869 (1995)
  PUBMED
            7579699
            2 (bases 1 to 299)
REFERENCE
  AUTHORS
            Balmain, A., Krumlauf, R., Vass, J.K. and Birnie, G.D.
  TITLE
            Cloning and characterisation of the abundant cytoplasmic 7S RNA
            from mouse cells
  JOURNAL
            Nucleic Acids Res 10 (14), 4259-4277 (1982)
  PUBMED
            6896909
COMMENT
            PROVISIONAL REFSEQ: This record has not yet been subject to final
            NCBI review. The reference sequence was derived from <a href="AC157822.2">AC157822.2</a>.
            Sequence Note: The RefSeq transcript and protein were derived from
            genomic sequence to make the sequence consistent with the reference
            genome assembly. The genomic coordinates used for the transcript
            record were based on alignments.
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PRIMARY
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