Salmonella sp. WGH-01 chromosome, complete genome

GenBank: CP096171.1

FASTA Graphics

Go to:

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LOCUS       CP096171               1001 bp    DNA     linear   BCT 16-MAY-2022
DEFINITION  Salmonella sp. WGH-01 chromosome, complete genome.
ACCESSION   CP096171 REGION: 3643480..3644480
VERSION     CP096171.1
DBLINK      BioProject: PRJNA827233
            BioSample: SAMN27606337
KEYWORDS    .
SOURCE      Salmonella sp. WGH-01
  ORGANISM  Salmonella sp. WGH-01
            Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacterales;
            Enterobacteriaceae; Salmonella.
REFERENCE   1  (bases 1 to 1001)
  AUTHORS   Zhao,J., Lu,X., Tie,A., Ngegba,E., Wang,L., Sun,L., Liang,Y.,
            Abdulai,M.K., Bah,S., Wang,G., Dong,X., Harding,D. and Kan,B.
  TITLE     Molecular diagnostics and next-generation sequencing reveal real
            etiological characteristics of invasive Salmonella infection in
            febrile illness in Freetown, Sierra Leone
  JOURNAL   Emerg Microbes Infect, 1-20 (2022) In press
   PUBMED   35537043
  REMARK    Publication Status: Available-Online prior to print
REFERENCE   2  (bases 1 to 1001)
  AUTHORS   Liu,J.
  TITLE     Direct Submission
  JOURNAL   Submitted (19-APR-2022) Chinese Center for Disease Control and
            Prevention, Chinese Center for Disease Control and Prevention, 155
            Changbai Road, Changping District, Beijing 102200, China
COMMENT     The annotation was added by the NCBI Prokaryotic Genome
Annotation
            Pipeline (PGAP). Information about PGAP can be found here:
            https://www.ncbi.nlm.nih.gov/genome/annotation_prok/
            NCBI analysis of this genome has determined that >30% of the
            predicted CDS features contain frameshifts, which is well above
the
            usually observed value of <10%.
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                     /isolation_source="blood samples from outpatients"
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ORIGIN

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