Salmonella enterica subsp. enterica serovar Pullorum plasmid pSPUV, complete sequence

GenBank: JN885081.1

FASTA Graphics

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LOCUS       JN885081            86686 bp    DNA     circular BCT 06-APR-2020
DEFINITION  Salmonella enterica subsp. enterica serovar Pullorum plasmid pSPUV,
            complete sequence.
ACCESSION   JN885081
VERSION     JN885081.1
DBLINK      BioSample: SAMN14226166
KEYWORDS    .
SOURCE      Salmonella enterica subsp. enterica serovar Pullorum
  ORGANISM  Salmonella enterica subsp. enterica serovar Pullorum
            Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacterales;
            Enterobacteriaceae; Salmonella.
REFERENCE   1  (bases 1 to 86686)
  AUTHORS   Feng,Y., Liu,J., Li,Y.G., Cao,F.L., Johnston,R.N., Zhou,J.,
            Liu,G.R. and Liu,S.L.
  TITLE     Inheritance of the Salmonella virulence plasmids: Mostly vertical
            and rarely horizontal
  JOURNAL   Infect. Genet. Evol. 12 (5), 1058-1063 (2012)
   PUBMED   22449772
REFERENCE   2  (bases 1 to 86686)
  AUTHORS   Feng,Y. and Liu,S.-L.
  TITLE     Direct Submission
  JOURNAL   Submitted (20-OCT-2011) Genome Research Center, Harbin Medical
            University, XueFu Road No. 194, Harbin, Heilongjiang 150081, China
FEATURES             Location/Qualifiers
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FSALDVSDVAVVAARRSRVEWENQQRKKQNLEPLEMDELIAKAWRFVRERFRSYQSER
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gene complement(25470..25583)
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CDS complement(25470..25583)
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/locus_tag="pSPUV_040"
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gene complement(25661..25906)
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/locus_tag="pSPUV_041"
CDS complement(25661..25906)
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/locus_tag="pSPUV_041"
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gene complement(26038..26418)
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CDS complement(26038..26418)
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/EC_number="3.1.-.-"
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CDS complement(26714..27166)
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gene complement(27283..27846)
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/locus_tag="pSPUV_044"
CDS complement(27283..27846)
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/locus_tag="pSPUV_044"
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gene complement(27901..28641)
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gene complement(33919..36252)
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CDS complement(33919..36252)
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gene complement(36629..37360)
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/locus_tag="pSPUV_048"

CDS complement(36629..37360)
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/locus_tag="pSPUV_048"
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gene complement(37606..38088)
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/locus_tag="pSPUV_049"
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gene complement(38110..40596)
/gene="traG"
/locus_tag="pSPUV_050"

CDS complement(38110..40596)
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gene complement(40586..41134)
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/locus_tag="pSPUV_051"

CDS complement(40586..41134)
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gene complement(41121..41414)
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/locus_tag="pSPUV_052"

CDS complement(41121..41414)
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gene complement(41628..42371)
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/locus_tag="pSPUV_053"

CDS complement(41628..42371)

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/locus_tag="pSPUV_053"

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MLAHPELDYNLQYSHYNGTVRNQLAADQTQQRQAISKLAEHYGIMFFYRGQDPIDGQL
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gene 42424..42603

/gene="trbE"

/locus_tag="pSPUV_054"

CDS 42424..42603

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/locus_tag="pSPUV_054"

/codon_start=1

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gene complement(43097..44923)

/gene="traN"

/locus_tag="pSPUV_055"

CDS complement(43097..44923)

/gene="traN"

/locus_tag="pSPUV_055"

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/transl_table=11

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ATITGDWRETSEVRTYTLTAFSFSRSGKQIVFSVTVPEAGTISSASLNVTTQNYLWNS
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GRTTFTLTLVMQVKDREWIPRVEWVESCPFNKVDGVLKDTECTEPGGTKTGVMEGKPW
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SKTAGKVMICGGDVFCLDGECDKAQSGKSNDFGEAVSQLAALAAAGKDVAALNGIDVR

AFTGQAKFCKKAAAGYSNCCKDSGWGQDIGLAKCSSDEKALAKAKTNKLTVSVGEFCS
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gene complement(44920..45339)
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CDS complement(44920..45339)
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/protein_id="AFC61073.1"
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RGRRSSVNFRTAERFGLISVVNKPVVFWFEQYEGENK"

gene complement(45376..46014)
/gene="trbC"
/locus_tag="pSPUV_057"

CDS complement(45376..46014)
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/locus_tag="pSPUV_057"
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gene complement(45999..46307)
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CDS complement(45999..46307)
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gene complement(46326..47318)
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/locus_tag="pSPUV_059"

CDS complement(46326..47318)
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gene complement(47315..47947)
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/locus_tag="pSPUV_060"
CDS complement(47315..47947)
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/locus_tag="pSPUV_061"
CDS complement(47944..48330)
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gene complement(48305..50952)
/gene="traC"
/locus_tag="pSPUV_062"
/pseudo

gene complement(50968..51351)
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CDS complement(50968..51351)
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/note="FIG01045198"
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gene complement(51348..51815)
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CDS complement(51348..51815)
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/locus_tag="pSPUV_065"
CDS complement(51808..52029)
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CDS complement(52164..52679)
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protein TraV"
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CDS complement(52878..53474)
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gene complement(53464..54885)
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GAVPPPTAFYPGNGVTPPPQVTYQSVPVPNRIQRKTFSYDAAAKKDPSLPYIPSGSFA
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gene complement(55612..56178)
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gene complement(56200..56511)
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CDS complement(56930..57325)
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/protein_id="AFC61090.1"
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gene complement(57424..58104)
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VMGIINNC"

gene complement(58308..58688)
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CDS complement(58308..58688)
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/locus_tag="pSPUV_076"
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gene 58973..59581
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CDS 58973..59581
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CDS complement(59892..60062)
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gene complement(60813..60983)
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CDS complement(60813..60983)
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CDS complement(61220..61825)
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gene complement(61813..62001)
/gene="psiB"
/locus_tag="pSPUV_083"
CDS complement(61813..62001)

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AVEHVCPPCHMRVLFCAC"
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CDS complement(64096..64209)
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CDS complement(64196..64552)
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/locus_tag="pSPUV_086"
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IVNGRRSGDAPSLGCA"
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CDS complement(64583..64711)
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gene 64879..65001
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CDS             complement(65325..65693)
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gene            complement(65884..66318)
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CDS             complement(65884..66318)
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gene            complement(66385..66819)
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CDS complement(66385..66819)
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CDS complement(67315..68079)
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GNWRLLADEYDHDNWLDEEDEDGLDKVLDAVLVRNARFCPVLLTLVNEPEEEIEAAGV
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gene complement(68151..68564)
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ADAILREIGRKGMGEHTGDGISDSTVQALYAGLRTEIPTVSVPADLLARVTNIAGRAL
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CDS complement(68620..68841)
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gene            complement(69666..69857)
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CDS             complement(69666..69857)
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gene            complement(69903..70259)
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CDS             complement(69903..70259)
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gene            complement(70252..70722)
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CDS             complement(70252..70722)
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gene complement(70859..70987)
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CDS complement(70859..70987)
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gene 71233..71655
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CDS 71233..71655
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/locus_tag="pSPUV_101"
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/protein_id="AFC61115.1"
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gene 71655..72929
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/locus_tag="pSPUV_102"

CDS 71655..72929
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/locus_tag="pSPUV_102"
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(UmuC)"
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GVLALTLHNQKRTEKLLSLQPVEEIWGVGRRISKKLNTMGITTALQLARANPTFIRKN
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gene complement(73011..73985)
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CDS complement(73011..73985)
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FNQKEIAELEGLSQAKVTRALQAAAVPQELISLFPVQSELSFSDYKILLEVNEKLSEK
GLTSEGLIQSVSDQHDAILSDYERPDDEQKSSILKLISQASQALIAPPPKEKSVISAL
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CDS complement(73985..75190)
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/locus_tag="pSPUV_104"
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CDS 75605..76546
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TLKLESGSFVEDDLRQYFSDVLYSLKTTAGDGYIHVLVEHQSTPDKHMAFRLIRYAVA

AMQRHLEAGHKKLPLVIPVLFYTGKRSPYPYSTRWLDEFDDTALADKLYSSAFPLVDV
TVIPDDEIAGHRSMAALTLLQKHIHQRDLAELVDRLAPILLAGYLSSSQVISLVHYIV
QAGETSDAEAFVRELAQRVPQHGDALMTIAQQLEQKGIEKGIQLGEQRGIEKGRSEGE
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CDS complement(76578..77144)
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/note="FIG01046430"
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gene complement(77201..77614)
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CDS complement(77201..77614)
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gene complement(77589..77708)
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CDS complement(77589..77708)
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/note="FIG01045535"

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FPVLSVRLA"

gene complement(78222..79337)
/gene="tlpA"
/locus_tag="pSPUV_110"

CDS complement(78222..79337)
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