

The nucleotide sequence of gene rplJ encoding ribosomal protein L10 of *Salmonella typhimurium*

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Gene rplJ of *S. typhimurium* LT2 was isolated from recombinant pNL1 (1) plasmid in the EcoRI-E- fragment and further cloned on vector plasmid pUC19 (2). The rplJ gene carried by the resultant pMW12 plasmid was sequenced according to the procedure (3). Maintenance of pMW12 showed it to be growth-detrimental for normal JM101 (2) *E. coli* host cells due to superexpression of the cloned rplJ gene. This feature demonstrated thus the ability of the heterologous L10 protein of *S. typhimurium* to feedback rplJL genes expression in *E. coli*. Nucleotide sequence determined confirmed the striking homology of *S. typhimurium* and *E. coli* (4) rplJ genes. The secondary structure providing for the coupled translation of L10 and L12 cistrons in *E. coli* L10–L12 mRNA (5) according to the nucleotide sequence determined is restored in *S. typhimurium*. Three a.a. (Val-62, Gln-67 and Thr-74) differ in L10 of *S. typhimurium* from that of *E. coli* (4).

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ATGGCTTTAAATCTTCAAGACAAACAAGCGATTGTTGCTGAAGTCAGCGAAGTAGCCAAAGGCGCGCTG  70
 M  A  L  N  L  Q  D  K  Q  A  I  V  A  E  V  S  E  V  A  K  G  A  L

TCTGCAGTAGTTGCGGATTCCCGTGGCGTAACTGTAGATAAAATGACTGAACTGCGTAAAGCAGGTCGT 140
 S  A  V  V  A  D  S  R  G  V  T  V  D  K  M  T  E  L  R  K  A  G  R

GAAGCTGGCGTATACATGCGTGTTGTTCGTAACACCCTGCTGCGCCGCGTCGTTGAAGGTACTCAGTTC 210
 E  A  G  V  Y  M  R  V  V  R  N  T  L  L  R  R  V  V  E  G  T  Q  F

GAGTGCCTGAAAGACACGTTTGTTGGTCCGACCCTGATTGCATACTCTATGGAACACCCGGGCGCTGCT 280
 E  C  L  K  D  T  F  V  G  P  T  L  I  A  Y  S  M  E  H  P  G  A  A

GCTCGTCTGTTCAAAGAGTTCGCGAAAGCGAATGCAAAATTTGAGGTCAAAGCTGCAGCCTTTGAAGGT 350
 A  R  L  F  K  E  F  A  K  A  N  A  K  F  E  V  K  A  A  A  F  E  G

GAGCTGATCCCGGCGTCTCAGATCGACCGCCTGGCAACTCTGCCGACCTACGAAGAAGCAATTGCACGC 420
 E  L  I  P  A  S  Q  I  D  R  L  A  T  L  P  T  Y  E  E  A  I  A  R

CTGATGGCAACCATGAAAGAAGCTTCGGCTGGCAAACTGGTTCGCACACTGGCTGCTGTACGCGATGCA 490
 L  M  A  T  M  K  E  A  S  A  G  K  L  V  R  T  L  A  A  V  R  D  A

AAAGAAGCTGCGTAA                                                         505
 K  E  A  A
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