

Nucleotide sequence of two proline tRNA (AGG and CGG) genes from chicken

D.Weill and T.Heyman

Institut Curie, Section de Biologie, Centre Universitaire, 91405 Orsay, France

Submitted September 17, 1990

EMBL accession no. X55888

We present the nucleotide sequence of two chicken tRNA proline genes. From base 277 to 348: tRNA^{Pro} (AGG), from base 748 to 819: tRNA^{Pro} (CGG). The latter is on the complementary strand. The only difference between the two tRNA^{Pro} is in the first position of the anticodon. Coding sequences have been identified on the basis of their homology (100%) to the corresponding mouse genes (1, 2). The first gene would correspond to the chicken tRNA^{Pro} (AGG) (3); tRNA^{Pro} with a CGG anticodon has not yet been found in chicken.

REFERENCES

1. Russo, T., Duilio, A., Ammendola, R., Costanzo, F. and Cimino, F. (1986) *Eur. J. Biochem.* **158**, 437-442.
2. Hu, J.C., Cote, B.D., Lund, E. and Dahlberg, J.E. (1983) *Nucl. Acids Res.* **11**, 4809-4821.
3. Ouenzar, B., Agoutin, B., Reinisch, F., Weill, D., Périn, F., Keith, G. and Heyman, T. (1988) *Biochem. Biophys. Res. Commun.* **150**, 148-155.

```

TCGTAGCCTG GTTTGGCCAG GTGAAGCCAC GCGATGACCC AGCTCAGCAA GCAGCAGCAC TGGTCAGCAG 70
AGGGCACAGG GGAGGGCACC AGGTCCGCTT AGCACAAACC TTTTGCACAT CACGATGACT GAATTAGTGT 140
CTTCTGCTGC TTCTGAAGAC TTGGCACCGC CCCAACAGCC GCCAGCCAAC GGCCACTGCA ACAAGGCAAA 210
CTATGCGGGC AGGACGCCTT CGGCCAGCCG TTCGTCTTCG AGCGACGGAA AACCGCCTCT CGCGCTGGCT 280
CGTTGGTCTA GGGGTATGAT TCTCGCTTAG GGTGCGAGAG GTCCCGGGTT CAAATCCCGG ACGAGCCCAA 350
GTTTTTTTTT TCTTCTTCTT TTTTTTCGTT TATCTTCGCC TCCTTGCCGN NCCNNNCCCA CCTACGCGTG 420
CATCGCACAG CGGCCGCCAC CGCGCAAAC AGAGGGACGA GCGACGCGCT GCTGCGTTCT AAGAGCAGGA 490
TTAACGTGAG AAAATGGCGA GGAAAAGGAC CCCTTCTTCC CCTTGCCGCC TTCTCANNNC CCACCGTGAG 560
CCCGGTTCCC TCGCACCGCC CTTTGCATCC CCAAGGCCTC CCTACCGCCG GCATGCCCGT CGCATCCTTG 630
AGCCCCGCAA CCTCGCTCCG CTCTATCAAT CCGCCGCGGG ACGGTCAAGA CGACAAAAAC GCTCCGAAGC 700
GCCACACGGG CGCAAAAACA GGAGGAGTTC TCCCCTCCGT TCACCGAGGG CTCGTCCGGG ATTTGAACCC 770
GGGACCTCTC GCACCCGAAG CGAGAATCAT ACCCCTAGAC CAACGAGCCG CACACAGTTA GGTTTTCCCA 840
GCTGAATACT TGACTCATGT CTCTCTCGCT TCAACATGTC TATGCTA 887

```