

**Sequence of the rabbit  $\alpha_{s1}$ -casein cDNA**

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Two overlapping cDNA clones isolated from two rabbit mammary gland cDNA libraries constructed with pBR322 (1) and  $\lambda$ gt11 (kindly provided by M. Edery) were sequenced by the method of F. Sanger (2).

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ATCGCTAGATCATCAACCCAACTTGCCTCTTTCACTGAGTTAACGGCTGGACCCAC  ATG AAQ CTT CTC ATC CTC ACT TGC CTT GTG GCT ACT GCT CTT OCC  105
                                                               M K L L I L T C L V A T A L A   +
AGG CAT AAA TTT CAT TTA GGA AAC CTC ACT CAG GAG CCT GAG AGC AGT GAG CAG GAA ATC TTA AAA GAA AGA AAQ CTC CTC  195
R H K F H L G H L K L T O E P E S S E O E I L K E R K L L +30
AGG TTT GTC CAG ACA GTA CCA CTA GAA TTA AGA GAG GAA TAT GTC AAT GAA CTG AAC AGG CAG AGA GAA CTT CTG AGA GAA AAA GAG AAT  285
R F V Q T V P L E L R E E Y V N E L N R O E L L R E K E N  60
GAG GAA ATC AAG GAA ACT AGA AAT GAA GAA ACT GAG GAA CAT GTT TTO GCA GAC COT GAG ACA GAA GCT AGC ATC AGC TCA TCA AGT GAG  375
E E I K G T R N E V T E E H V L A D R E T E A S I S S S E  90
GAA ATT GTT CCC AGC AGC ACC AAC CAQ AAC AGG TAC GTG CCA AGG GAA GAC CTG GCT TAC CAA CCT TCG AGT CAG CAG CTT CTC AGA ATG  465
E I V P S S T K Q K Y V P R E D L A Y O P Y V Q Q O O L R M  120
AAA GAA CGC TAC CAA ATC CAG GAG AGA GAG CCT ATG AGA GTG GTG AAT CAG GAA CTG GCT CAG CTC TAT CTT CAG CCT TTC GAA CAA CCC  555
K E R Y Q O I Q E R P E M R V N Q E L A O L Y L O P F E O P  150
TAC CAG CTT GAT GCC TAT CTC CCT GCT CCT TGG TAC TAT ACT CCG GAA GTG ATG CAG TAT GTT CTT TCC CCA CTG TTC TAT GAC CTC GTT  645
Y Q L D A Y L P A P W Y Y T P E V M Q Y V V L S P L F Y D L V  180
ACA CCC AGT GCC TTT GAG AGT OCT GAA AAA ACT GAC GTT ATT CCA GAG TGG TTG AAG AAT TAAGTCATTCTCAAGGAACTCCACAATTATGACCATGGT  745
T P S A F E S A E K T D V I P E W L K N   200
GTGACTGAAAATCAACCTTAATGAAATTCTCATCTTGTGTTATAATCTAAACCCACTTATCCTAAAGGACTTGTTCTAGGATAGQAATAGCAATTCTCATTTGAGGGCTATCTTCTTTTG  865
AGCTTACTCTTTAACAGAACATCATCTTACTCTTAAAGCAAAATTCTTAAAGCACTTATCTCAACTCAGTGTATCATGCCAGTATGQAAGCACCTTGTTAGGGGGT  985
ATTAAAGCTTGTAGTAAAGTTCTGTAACTGQAATTTCTGTTGGATGQATTTTCTGTAAAGTCAATCATGTCAAATTATGTGTGAGTGTGAGTGTGAGTTCTTCTTCT  1105
CTCAATTAAATCATTTAAAGGCA   1130

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The 1130 nucleotide sequence encodes a 215 amino acid polypeptide with a Mr of 25'499. The rabbit  $\alpha_{s1}$ -casein corresponds in size to the corresponding bovine and ovine polypeptides, but is shorter than the rat corresponding  $\alpha_{s1}$ -casein, which contains at position 123 a ten time repeat of a 6 amino acid sequence (3).

The 15 amino acids of the rabbit  $\alpha_{s1}$ -casein signal sequence determined by protein sequencing (4) matches exactly to the deduced amino acid sequence (nucleotides 61 to 105). They reveal 87 (93) and 89 % (93) similarities with the rat (3) and bovine (5) signal sequences, respectively, whereas for the coding region, the homology is only 42 (38) and 55 % (34), respectively. Similarity is higher at the protein level (number in brackets) in the signal sequences, and higher at the nucleotide level in the coding regions. Among the 7 potential phosphorylation sites deduced from the cDNA sequence, three are conserved in rat, but not in bovine or ovine caseins (6).

In the three species, the 3' untranslated region is rather long. This sequence, as well as the leader sequence are more conserved among species than the coding sequence suggesting that they might play a role in mRNA stabilization or in transcriptional regulation.

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