
Nucleotide sequence of novel cDNAs generated by alternative splicing of a rat thyroid hormone receptor gene transcript

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Submitted April 26, 1988

Accession no.X07409

Two cDNAs (rTR α vI,vII) were obtained from a rat brain library by screening with the Pst I fragment of v-erbA. Comparative nucleotide sequence analysis revealed extensive similarities with the c-erbA sequences, reported to encode thyroid hormone receptors (1,2,3,4). Nucleotide sequences of these cDNAs are identical with that of rTR α , vI starting at nt 371 of rTR α and vII at nt 776, except for the substitution of the A in rTR α (nt 1165,1177) by the G in vI (804,816) and vII (392,404). The homology suddenly disappears at nt 1075 of vI (663 of vII), coinciding with nt 1438 of rTR α . Consequently the last 40 amino acids of the rTR α carboxy terminus (hormone binding domain) are replaced by 122 amino acids in vI, whereas vII contains only the last 83 amino acids of vI. Both variants share an identical 3' noncoding sequence. We have recently reported that rTR α and its variants are generated by alternative splicing of the same gene transcripts (5). Of particular interest is our observation that *in vitro* translation products of rTR α variants do not bind thyroid hormone and they are abundantly expressed in brain. The identical sequence in the DNA binding domains of the receptor and its variant forms suggests possible functions of the variants in thyroid hormone action. Shown is the cDNA sequence of vI starting at nt 1073 which corresponds to the last nucleotide of the common exon. The nucleotides absent in rTR α vII are underlined. The entire cDNA sequences of rTR α vI and vII have been submitted to the EMBL Library.

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1073 GGAGAGAGAAGTGCAGAGTTCCGATTCTGTACAAGGGCCAGCGCCACAAGCCGCCGGGGGGTCACTGGCGCTCCACCGGAAGGACAG
1163 CAGCTTCTCGGAATGCATCTTGTTCAGCGTCCCGACGCTCCCGCAGCTTGACCCAGCAGTTTCGTGAAGCCGGAAGTCTCCGAGGGCCGGTTC
1253 TTACGACCCAGAGCCCGAAGAGCCCGCAGCAGCGCTCTCTGGAGCTGCTCCACCGAAGCCGGAATTTCCGATTCCTCGAGCGGTCTGTGGGGA
1343 AGACGACAGCAGTGAAGCAAGTCCCTGAGCTCCTTCTCTGACGAGGACACGGAGGTCTTCGAGGACCTGGCAGGCAAGGCAGCCTCT
1423 CCCTGAGGCCCTTGGAGGGCGATGGGGAAGGAGAAGGAGCATGCTACTCTTCTCCAGGGCCCTGCGCCATAGAGCTGGGCACGCCACAT
1513 GCTGGGAGCAAGGCCACAGGCTGGCCGCTCCCACTCAAGACCACCCCTCTACCCCTGAGCACGCCACACGTTGGCCAAAGCTCCCTTGTAT
1603 TGTTCTGTAGTTCCTCTGCTGGGATGCCCTTCCCGCTCTATGCCTGGCAATACCTTGTCCCTTGGAGGCCCACTCAAGTGTCAAC
1693 TCCTTCCCGAGCTCTCCGAGGCAAAATAGTTGTCTGTGCTTACGTTGATGCTTCTCTGTGACACTTCACTGTTTTATAATTAGT
1783 CGGGCATGAGTCTCTTCCCAAGCTAGACTGTGTCTGAATCATGTCTGTAGCCCACTGCCAGTGGCCGGCTGGCATAGAGATAGGAAC
1873 TCCATAAAAGCCGAATTC

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References: 1) Sap, J. *et al.* (1986) *Nature* **324**, 635-640. 2) Weinberger, C. *et al.* (1986) *Nature* **324**, 641-646. 3) Thompson, C.C. *et al.* (1987) *Science* **237**, 1610-1614. 4) Benbrook, D., and Pfahl M. (1987) *Science* **238**, 788-791. 5) Mitsuhashi, T. *et al.* (1988) *Proc. Natl. Acad. Sci. USA* (in press).