

Nucleotide sequence of the Chinese hamster ornithine decarboxylase gene

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Ornithine decarboxylase (ODC; EC 4.1.1.17) is the first enzyme in the pathway of polyamine biosynthesis in eukaryotic cells, and it plays a key role in the regulation of intracellular polyamine levels. Genomic and cDNA sequences have been reported for this gene from several species (1-7). Srinivasan *et al* have reported a partial cDNA sequence for ODC from the Chinese hamster lung fibroblast V79 line (8). This sequence contains the 3' half of the coding sequence and the entire 3' UTR. In this report we present the sequence of the entire coding region and the 5' UTR of an ODC cDNA from Chinese hamster ovary cells (Figure 1). Our sequence of the 3' UTR confirms that which has been reported previously. The coding sequence is strongly conserved among all mammalian species examined so far, while the UTRs are considerably more divergent.

CGCAGCGGGCGCCGCCGCTCCGCCGCCCCCTCAGCCAGCTGCCACTCGGGCAGCGTCTGCCGCCGCTCGACGA
GGCGCTGACGGGGCGGGCGGAGCTCTCGGGTTCTCGCGGCCACTAGTTCTCATGGGGCTGGCG
AGTCATTGCGTCCGCGTGTGAGGACGTTCACATTCAAGGAGTCTGGAAAGTCTGGATAGTGTGCTGGAGGAA
CTGCCATAACTGGATTCCATCTCTAGAGTTTGAGCACACCGAGGCCATGAAACAGCTCAATAAGGACGAGTTG
ACTGCCATACCTCGATGAAGGCTTACGGCCAAGGACATTCTGGACCAAAAAAATTAAATGAAGTATCCTCTGATGACA
AGGATGCTTTATGTCGGCGGACCTTGGAGACGTTCTGAAGAACGACCTAAAGATGGCTAAAAGCTCTCCCCTGACTC
CCTTTATCCAGTCAAATGTAATGACAGCAGCGTTAGTGAACACCTAGCTGCCATTACAGTGTGACTGTGCAAGCA
AGACTGAGATACAGTTGGTACAGGGCTTGGAGTGCTCTCCGAGAGAGTCATCTATGCAAATCCATGAAAGCAAGTGT
CTCAGATCAAGTATGCCGCCAGCAATGGAGTCCAGATGATGACTTTGACAGTGAATTGAGTTAATGAAGGTCGCA
GAGCACATCCAAAAGTTACCAAGTGGTTTGCGGATGCCACTGACGATTCTAAAGCAGTGTGACTCAGTGTAA
AGTTGGTCCACACTCGAACACGAGCTTCTCTTGGACCGGCAAAAGAGCTAAATATTGATGTCATTGGTGTCA
GCTTCCACCTGGGGACTGGATGACTGACCTTGAGACCTTCGTCAGGCCCTTGCGGATGCCGCTGTCTTGACA
TGGGAACAGAAGTTGGTTTCAGCATGTATGCTGTGATATTGGTGGTGGCTTCCGATCTGAGGATACGAAGCTTA
AATTGAGAGATCACCAAGTGTATCAACCCAGCTTGGACAAGTACTCCCGCAGACTTGGAGTGAAGGAGTATAG
CCGAGCCAGGCAGACTACAGTGGCTCAGCTTCCACACTGGCAGTCATATCATAGCCAAGAAAATCGTATCGAAGG
GCTCTGACCATGAAGATGAGTCCAGTGAACCTTATGTTGATGAGTGGAGTGTATGGGTCTTAACT
GCATTCTTACGATCATGCACATGTGAAGCCCTGCTGCCGAAGAGACCCAAGCCAGATGAGAAGTATTACTCATCCA
GCATCTGGGGACCAACATGCGATGGCCTTGACCGGATTGAGGCGCTGTAATCTGCTGAAATGCATGTGGTGATT
GGATGCTCTTGAGAACATGGGTCATACACTTGGCTCTGCACTACTTCAACGGGTTCCAGAGGCCCTTCTATCT
ACTATGTTGATGTCGAGGCCAATGTGGCAGCTTGAAGCAGATCCAGAACCATGGCTCCCACAGAAGTGGAGAGC
AGGATGTTGGCACTCTGCCCATCTTGTGCCAGGAGAGCGGGATGGACCGTCATCCAGCAGCTGTGCTCTGCTA
GTATCAATGTCAG

Figure 1. Chinese hamster ornithine decarboxylase cDNA sequence: nucleotide 1 is the transcription start, the translation initiation (ATG, nt 285) and termination (TAG, nt 1650) sites are underlined.

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REFERENCES

1. Fonzi, W.A. and Sypherd, P.S. (1987) *J. Biol. Chem.* **262**, 10127.
2. Phillips, M.A., Coffino, P. and Wang, C.C. (1987) *J. Biol. Chem.* **262**, 8721.
3. Hickok, N.J., Seppanen, P.J., Gunsalus, G.L., and Janne, O.A. (1987) *DNA* **6**, 179.
4. McConlogue, L., Gupta, M., Wu, L. and Coffino, P. (1984) *Proc. Natl. Acad. Sci. USA* **81**, 540.
5. Brabant, M., McConlogue, L., Van Daalen Wetters, T. and Coffino, P. (1988) *Proc. Natl. Acad. Sci. USA* **85**, 2200.
6. Kontula, K.K., Torkkeli, T.K., Bardin, C.W. and Janne, O.A. (1984) *Proc. Natl. Acad. Sci. USA* **81**, 731.
7. Van Steeg, H., Van Oostrom, C.T.M., Van Kranen, H.J. and Van Kreijl, C.F. (1988) *Nucl. Acids Res.* **16**, 8173.
8. Srinivasan, P.R., Tonin, P.N., Wensing, E.J. and Lewis, W.H. (1987) *J. Biol. Chem.* **262**, 12871.