cDNA and deduced amino acid sequence of rat copper-zinc-containing superoxide dismutase

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Superoxide dismutases are involved in cellular antioxidant defense mechanisms against the oxidative damage mediated by superoxide anion radicals which are generated intracellularly during normal metabolism (1). Two distinct types of superoxide dismutases have been found in mammalian cells. The copper-zinc-containing superoxide dismutase consisting of two identical subunits is found principally in the cytosol (2), and the manganese-containing superoxide dismutase consisting of four identical subunits is found predominately in the mitochondrial matrix (3,4). We have isolated cDNA clones coding for the rat copper-zinc-containing superoxide dismutase from a Sprague-Dawley rat liver λ gtll cDNA library (Clontech Laboratories, Inc., Palo Alto, CA) by cross hybridization with the corresponding human cDNA (5,6). Sixty hybridized clones were recovered from ~ 350,000 recombinant bacteriophages. The nucleotide and deduced amino acid sequences of one of these clones containing the entire coding region are shown in Figure 1. The deduced amino acid sequence of the 154 amino acids of rat copper-zinc-containing superoxide dismutase 83.1% identity with the corresponding human enzyme (5,6).

Figure 1. Nucleotide and deduced amino acid sequences of rat copper-zinc-containing superoxide dismutase cDNA.

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