

The sequence of the Cu-Zn superoxide dismutase gene of *Drosophila*

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The sequence shown below is a 1.8 Kb EcoR1 fragment from a genomic library made from an isogenic stock of *Drosophila melanogaster* (Oregon R) (J. Leung, unpublished). The sequence contains the transcribed region for a Cu-Zn superoxide dismutase (SOD) gene (in capitals), 413 bp of the 5'-untranslated region, and 247 bp of the 3'-untranslated region. The amino acid sequence of the Cu-Zn SOD of *Drosophila* has been reported (1) as well as the sequence of a corresponding cDNA (2). Transcription start and stop sites of the gene are assumed to be the limits of the sequence of the cDNA (2). Two additional triplets in the translated DNA sequence indicate that the initial translation product probably contains not only an N-terminal methionine residue but also a C-terminal valine. Both amino acids would have to be removed to give the amino acid sequence reported for SOD (1). In addition, the gene contains a single 725 bp intron which separates the triplets for amino acids 21 and 22 of SOD. This intron is in the same position as the first intron of the human Cu-Zn SOD gene (3). The transcribed region of the gene has three transitions (overlined) when compared with the sequence of the corresponding cDNA(2) from a different *D. melanogaster* source (4). These changes would not affect the amino acid sequence of the SOD. Possible control sequences are underlined. As with the cDNA, the genomic DNA hybridized to 68A4-9 on *Drosophila* polytene chromosomes.

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gaattcctggattcggttttttattatacaaaacaaagtaaatgaaatgctgcgccactgtcattggaataaa -271
tgaaggcctccaagtgaccaccctcggtggaacagctcaaaaaattcaacgccatttgcagtaaaattgttccgttttcaaat -181
tgaattttagcttttctcttaaaaatgtaaaataaagtcccaataaaacatgagcttgaatattacaagaataaaatgctccagt -91
accgccataatagtggaacgaaactttgcacaactgaacactaacagtaaaagtccgcattgtttgtaagctgctctgcacgctc -1
ACCCATAGAAGATACCTGGAAAGTCTCAACTTTTTTCGTTTTGATAAATTGATTAATTCATTGCGAAATGGTGGTTAAAGCTGTCTGGC 90
M V V K A V C
TAATTAACGGCGATGCCAAGGGCACGGTTTTCTTTCGAAACAGGAGGTGAGAATCCAAAATCATTGAACTTCTCTGCTCGGCAAAATGTAC 180
V I N G D A K G T V F F E Q E
GAAAAACAGAAGTTCTAAAGGTCAAATAGCCGGCTGCACC CGCGCCCTCTTCCACTTCAATATGCTGCTTAAATTTGTCGAGCAT 270
TTTAAATTAAGTCGATTGAGTTTACGCCTAGTCACCCAGCAAGTGACCTTTATATTTAATAAGCCGCACAAAATGGCAGATATGTGT 360
GTGGCCTCAAGTGCCTACAGCAAAGGTACAGAAATAGTACTGGACATAAAAAGGAGTTAAGATATAAAGCTCACTTGTTCGTAAGATAT 450
CGTAAATATCAACAAATATTTGTTTGAATAAAGCAATAGGAATATGGGAATAATAGAATGATGCTGTTCAATTAATTTGTACATC 540
AAAGTCAAAGCAGCAATGTCAAAGTCAAAGTAAACGATTATAAATGATGATTACAGGTATGTTTCAGTGCAGGAAATTTATGTTT 630
TTAATCTAAAGATAAACCAGTGTACTTTGCTGCCATATAAATATTTCCGTTTAAACGTGTGCTATTAACAAATGTTATTTTCAGTATA 720
TAACCTATTATCATATGAAGTTGGCCACGCTCGTTATCATAATCAGTGCTTCTGCTCACTATTATACACAACTTGTGCTTATCAGTATT 810
CGAGTATTATCTGAAGCGTTATAACCAATCCCTTCATCCCGTCCACAGAGCAGCGGTACGCCGTGAAGGCTCCCGTGAGGTGTGCCG 900
S S G T P V K V S G E V C G
CCTGGCCAAAGGTTCTGCACGGATTCCACGTGCACGAGTTCGGTGACAACACCAATGGCTGCATGTGCTCGGACCGCACTTCAATCCGTA 990
L A K G L H G F H V H E F G D N T N G C M S S G P H F N P Y
TGCCAAAGGAGCATGGCGCTCCCGTCGACGAGAAATCGTACCTGGCGATCTGGCAACATTGAGGCCACCGCGACTGCCACCAAGGT 1080
G K E H G A P V D E N R H L G D L G N I E A T G D C P T K V
CAACATCACCGACTCAAGATTACGCTCTTCCGGCCGACAGCATCATCGACGACCCTGTTGCTGCACGCCGATCCGATGATCTTGG 1170
N I T D S K I L F G A D S I I G R T V V V H A D A D D L G
CCAGGGTGGACACGACTGAGCAAGTCAACGGSCAACGCTGGTGCCCGCATCGGATCGGCGTATTGGCATGCCAAGGTCTAAGCGAT 1260
O G G H E L S K S T G N A G A R I G C G V I G I A K V *
AATCTATTCCGATGTCCGCCACTGTGCTGATCTACTCTATTTAGCACTACCCACTGGAGATATGCAACGATATACATCTCTAAACAT 1350
AAATACATACGCTGTGGTCTGTAGTTGATGACGCAACCTTTGAGGTTCAATAAAATGGTGTTTGAAATGCCCCATAAACaaagttat 1440
agtttcaattgagttgagtggtgaagtgaaatatacactgttgcctgcagcaattc 1499
    
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