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Nucleotide V

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GenBank: X01714.1

E. coli dut gene for dUTPase (EC 3.6.1.23) (deoxyuridine 5'-triphosphate nucleotidohydrolase)

FASTA Graphics Go to: LOCUS DNA linear BCT 23-0CT-2008 X01714 1609 bp E. coli dut gene for dUTPase (EC 3.6.1.23) (deoxyuridine DEFINITION 5'-triphosphate nucleotidohydrolase). ACCESSION X01714 X01714.1 VERSION KEYWORDS dUTPase; unidentified reading frame. SOURCE Escherichia coli ORGANISM Escherichia coli Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacterales; Enterobacteriaceae; Escherichia. REFERENCE 1 (bases 1 to 1609) AUTHORS Lundberg, L.G., Thoresson, H.O., Karlstrom, O.H. and Nyman, P.O. Nucleotide sequence of the structural gene for dUTPase of TITLE Escherichia coli K-12 TOURNAL. EMBO J. 2 (6), 967-971 (1983) PUBMED 6139280 Data kindly reviewed (25-NOV-1985) by L. Lundberg. COMMENT FEATURES Location/Qualifiers source 1..1609 /organism="Escherichia coli" /mol_type="genomic DNA" /db_xref="taxon:<u>562</u>" 286..291 regulatory /regulatory class="promoter" /note="-35 region" 310..316 regulatory /regulatory_class="promoter" /note="-10 region" 322..324 misc_feature /note="put. transcription start region" 330..333 regulatory $/ {\tt regulatory_class="ribosome_binding_site"}$ /note="put. rRNA binding site" 343..798 CDS /note="unnamed protein product; dUTP-ase (aa 1-151)" /codon_start=1 /transl table=11 /protein_id="<u>CAA25859.1</u>" /db_xref="GOA: P06968" /db_xref="InterPro: IPR008180" /db_xref="InterPro:<u>IPR008181</u> /db_xref="PDB:<u>1DUD</u>" /db_xref="PDB:1DUP" /db_xref="PDB: 1EU5 /db_xref="PDB: <u>1EUW</u>" /db_xref="PDB:<u>1RN8</u>" /db_xref="PDB:<u>1RNJ</u> /db_xref="PDB: ISEH" /db_xref="PDB: ISYL' /db_xref="PDB: 2HR6 /db_xref="PDB: 2HRM" /db_xref="UniProtKB/Swiss-Prot: P06968" /translation="MKKIDVKILDPRVGKEFPLPTYATSGSAGLDLRACLNDAVELAP GDTTLVPTGLAIHIADPSLAAMMLPRSGLGHKHGIVLGNLVGLIDSDYQGQLMISVWN RGQDSFTIQPGERIAQMIFVPVVQAEFNLVEDFDATDRGEGGFGHSGRQ" misc feature 831..851 /note="put.stem-loop structure"

831..838

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