

Product datasheet for SC207711

CYP26A1 (NM_000783) Human 3' UTR Clone

Product data:

OriGene Technologies, Inc.

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Clones A1 (NM_000783) Human 3' UTR Clone
2xxxxx (DC1000C2)
arget (PS100062)
A1
CYP26; P450RAI; P450RAI1
00783
711 3'UTR clone of NM_000783 quence shown below is from the reference sequence of NM_000783. The complete ce of this clone may contain minor differences, such as SNPs. top Codon Red=Cloning site
TTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TTGGCAGAGCTCAGAATTCAAGCGATCGCC ACCCATTTCCATGGGGAAATCTGATGAGCTTGAATGTTCAAACCTGAGACTTATTGGAAGTGT GAGTTTTTAAGGAGTGTTGTGTTGACTTTATATTTAATTTCTAAATGTATATTATAATATTTA TTGACTATACTACCACAATCTTTAAATATTAAAATAATGAATTTGTATCATTTCCAAATAAAG TTGAAGGTACTTTTCTGGTATTTTAAGATTCCTGTTGGGTAAAACTCACCAGTTTAGTATTTT GTATTTAACCAGATTTTACAATGCCTACCTGGACTTATTTGTCATCTTTGCATCTGTTTTCTG GAAATCTTAGCTGTTTTTAAGATTCCTGTGGGAAAATATATGCATCTGTGTGTG
lui
olecular clone sequence data has been matched to the sequence identifier above as a of reference. Note that the complete sequence of this clone is largely the same as the nce sequence but may contain minor differences , e.g., single nucleotide orphisms (SNPs).
DNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The ge also includes 100 pmols of both the corresponding 5' and 3' vector primers in ate vials.



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	CYP26A1 (NM_000783) Human 3' UTR Clone – SC207711
RefSeq:	<u>NM 000783.4</u>
Summary:	This gene encodes a member of the cytochrome P450 superfamily of enzymes. The cytochrome P450 proteins are monooxygenases which catalyze many reactions involved in drug metabolism and synthesis of cholesterol, steroids and other lipids. This endoplasmic reticulum protein acts on retinoids, including all-trans-retinoic acid (RA), with both 4-hydroxylation and 18-hydroxylation activities. This enzyme regulates the cellular level of retinoic acid which is involved in regulation of gene expression in both embryonic and adult tissues. Two alternatively spliced transcript variants of this gene, which encode the distinct isoforms, have been reported. [provided by RefSeq, Jul 2008]
Locus ID:	1592
MW:	23.9

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