

Product datasheet for SC335030

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UGT2B10 (NM_001290091) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: UGT2B10 (NM_001290091) Human Untagged Clone

Tag: Tag Free

Symbol: UGT2B10

Synonyms: UDPGT2B10

Mammalian Cell

Neomycin

Selection: Vector:

pCMV6-Entry (PS100001)

E. coli Selection: Kanamycin (25 ug/mL)

Fully Sequenced ORF: >NCBI ORF sequence for NM_001290091, the custom clone sequence may differ by one or

more nucleotides

TAG

Restriction Sites: Sgfl-Mlul

ACCN: NM 001290091

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).





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Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method: 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of

shipping when stored at -20°C.

RefSeq: <u>NM 001290091.1</u>, <u>NP 001277020.1</u>

 RefSeq Size:
 2187 bp

 RefSeq ORF:
 843 bp

 Locus ID:
 7365

 UniProt ID:
 P36537

 Cytogenetics:
 4q13.2

Protein Families: Transmembrane

Protein Pathways: Androgen and estrogen metabolism, Ascorbate and aldarate metabolism, Drug metabolism -

cytochrome P450, Drug metabolism - other enzymes, Metabolic pathways, Metabolism of xenobiotics by cytochrome P450, Pentose and glucuronate interconversions, Porphyrin and

chlorophyll metabolism, Retinol metabolism, Starch and sucrose metabolism

Gene Summary: UDPGT is of major importance in the conjugation and subsequent elimination of potentially

toxic xenobiotics and endogenous compounds.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (3) differs in its 5' UTR and uses a downstream start codon, compared to variant 1. The encoded isoform (3) has a shorter N-terminus, compared to isoform 1. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.