

Product datasheet for SC210534

XYLT2 (NM 022167) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: XYLT2 (NM_022167) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: XYLT2

Synonyms: PXYLT2; SOS; XT-II; XT2; xylT-II

ACCN: NM_022167

Insert Size: 924 bp

Insert Sequence: >SC210534 3'UTR clone of NM_022167

The sequence shown below is from the reference sequence of NM_022167. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

CTCTGTGTTTTTCCTCCTGGGTGTCAG

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

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

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).



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XYLT2 (NM_022167) Human 3' UTR Clone - SC210534

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeq: <u>NM 022167.4</u>

Summary: The protein encoded by this gene is an isoform of xylosyltransferase, which belongs to a

family of glycosyltransferases. This enzyme transfers xylose from UDP-xylose to specific serine residues of the core protein and initiates the biosynthesis of glycosaminoglycan chains in proteoglycans including chondroitin sulfate, heparan sulfate, heparin and dermatan sulfate. The enzyme activity, which is increased in scleroderma nationts, is a diagnostic

sulfate. The enzyme activity, which is increased in scleroderma patients, is a diagnostic marker for the determination of sclerotic activity in systemic sclerosis. Alternatively spliced

transcript variants have been found for this gene. [provided by RefSeq, Dec 2013]

Locus ID: 64132 MW: 34.2