

Product datasheet for SC312777

OriGene Technologies, Inc.

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ST3GAL3 (NM_174966) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: ST3GAL3 (NM 174966) Human Untagged Clone

Tag: Tag Free Symbol: ST3GAL3

Synonyms: DEE15; EIEE15; MRT12; SIAT6; ST3GALII; ST3Gal III; ST3GalIII; ST3N

Mammalian Cell

Selection:

Neomycin

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >SC312777 representing NM_174966.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

ATGGGACTCTTGGTATTTGTGCGCAATCTGCTGCTAGCCCTCTGCCTCTTTCTGGTACTGGGATTTTTG
TATTATTCTGCGTGGAAGCTACACTTACTCCAGTGGGAGGAGGAGGACTCCAATTCAGTGGTTCTTTCCTTT
GACTCCGCTGGACAAACACTAGGCTCAGAGTATGATCGGTTGGGCTTCCTCCTGAATCTGGACTCTAAA
CTGCCTGCTGAATTAGCCACCAAGTACGCAAACTTTTCAGAGGGAGCTTGCAAGCCTGGCTATGCTTCA
GCCTTGATGACGGCCATCTTCCCCCGGTTCTCCAAGCCAGCACCCATGTTCCTGGATGACTCCTTTCGC
AAGTGGGCTAGAATCCGGGAGTTCGTGCCGCCTTTTTGGGATCAAAGGTCAAGACAATCTGATCAAAGCC
ATCTTGTCAGTCACCAAAGAGTACCGCCTGACCCCTTGGACAGCCTCCGCTGCCCCCGCTGCATC
ATCGTGGGCAATGGAGGCGTTCTTGCCAACAAGTCTCTGGGGTCACGAATTGACATTGTG
GTGAGACTGAATTCAGCACCAGTGAAAGGCTTTGAGAAAGGACGTGGGCAGCAAAACGACACTGCGCATC
ACCTACCCCGAGGGCGCCATGCAGCGGCCTGAGCAGTACGAGCGCGATTCTTCTTTTTCCTCGCCGGC
TTCAAGTGGCAGGAGAAAGAGTTTCTGCGGAAAGCTGGTGAAAGCTCCGCGTCATCACTGATCTAAGCAGTGGC

ATCTGA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT

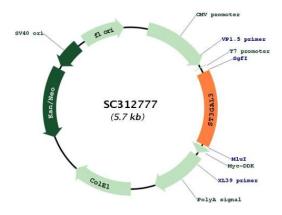
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

Restriction Sites: Sgfl-Mlul





Plasmid Map:



ACCN: NM_174966

Insert Size: 834 bp

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).

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning

into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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 174966.2</u>

RefSeq Size: 2000 bp RefSeq ORF: 834 bp



ST3GAL3 (NM_174966) Human Untagged Clone - SC312777

 Locus ID:
 6487

 UniProt ID:
 Q11203

 Cytogenetics:
 1p34.1

Protein Families: Secreted Protein, Transmembrane

Protein Pathways: Glycosphingolipid biosynthesis - lacto and neolacto series, Keratan sulfate biosynthesis,

Metabolic pathways

MW: 31.6 kDa

Gene Summary: The protein encoded by this gene is a type II membrane protein that catalyzes the transfer of

sialic acid from CMP-sialic acid to galactose-containing substrates. The encoded protein is normally found in the Golgi apparatus but can be proteolytically processed to a soluble form. This protein is a member of glycosyltransferase family 29. Mutations in this gene have been associated with a form of autosomal recessive nonsymdromic cognitive disability as well as infantile epileptic encephalopathy. Multiple transcript variants encoding several different

isoforms have been found for this gene. [provided by RefSeq, Jul 2017]

Transcript Variant: This variant (4) has multiple differences in the coding region but maintains the reading frame compared to variant 1. The resulting isoform (d, also called B4) is shorter compared to isoform a. 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.