

## Product datasheet for **MR210156**

### **Stt3a (NM\_008408) Mouse Tagged ORF Clone**

#### **Product data:**

Product Type:	Expression Plasmids
Product Name:	Stt3a (NM_008408) Mouse Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	Stt3a
Synonyms:	AA408947; BB081708; ltm1
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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ORF Nucleotide  
Sequence:

>MR210156 ORF sequence  
Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**CGGATCGCC**

ATGACTAAGCTTGGATTTTTGCGATTGTCCTATGAGAAGCAGGACACACTTCTAAAGCTTCTCATCCTGT  
CGATGGCTGCTGTGTTATCTTTTTCTACTCGTCTTTTTGCTGTGCTGAGATTTGAAAAGTGCATCCATGA  
GTTTGATCCGTACTTTAATTATCGGACTACCCGGTTTCTGGCTGAGGAGGGGTTTTATAAATCCATAAC  
TGGTTTGTGACCGGGCTTGGTACCCTTTGGGCCGAATCATTGGAGGAACAATTTACCCAGGTTTAAATGA  
TCACTTCTGCTGCAATCTACCATGTACTCCATTCTTCCATATCACTATTGACATTCCGGAATGTCTGTGT  
TTTCTGGCCCCACTTTTCTCTCTTCCACCACCATCGTTACGTACCACCTTACCAAAGAGCTCAAGGAT  
GCAGGAGCTGGGCTTCTGCTGCTGCCATGATTGCTGTAGTTCCTGGGTATATTTCTCGATCTGTAGCTG  
GCTCCTATGATAATGAAGGAATTGCTATCTTTTGCATGCTGCTTACTTACTACATGTGGATCAAGGCAGT  
GAAGACTGGTCCATCTATTGGGCTGCCAAGTGTGCCCTCGCTTATTTCTACATGGTCTCTTCATGGGGA  
GGCTATGTGTTCCCTGATCAACTTGATTCCCTCTACATGTCCCTGGTCTAATGCTGACAGGCCGTTTTTCTC  
ACCGGATCTACGTAGCCTACTGTACTGTTACTGCCTGGGCACCATTCTTTCTATGCAGATTTCTTTTGT  
TGGTTTCCAGCCCGTCTTTTCATCAGAACATGGCAGCCTTTGGAGTGTGGTCTCTGTGAGTCCAT  
GCTTTTCGTAGATTACCTGCGCAGCAAGTTGAATCCACAGCAATTCGAAGTCTTTTCCGGAGTGTATCT  
CCCTGGTTGGCTTTGCTCCTCACTGTGGGAGCTCCTCATGCTAACAGGAAAAATTTCTCCCTGGAC  
AGGGCGTTTCTACTCTCTGCTGGATCCCTCTTATGCTAAGAATAACATTTCCATTTATGCATCTGTTTCT  
GAGCACCAGCCCACAACCTGGTCTTCTACTATTTTGTACTACAGCTCCTTGTCTTCATGTTTCCAGTTG  
GCCTCTATTACTGCTTTAGCAACCTGTCTGATGCTCGGATTTTTATCATCATGTATGGTGTGACCAGGAT  
GTACTTTTTCAGCTGTAATGGTGCGTCTAATGCTGGTATTGGCACCTGTTATGTGCATTCTTCTGGCATT  
GGTGTTCCTCAGGTGCTGTCCACATATATGAAAAATCTGGACATAAGTCGCCAGACAAGAAGAGCAAGA  
AGCAACAGGATTCTACTTACCCTATTAAGAATGAGGTGGCGAGTGGGATGATACTGGTCAATGGCTTTTTT  
TCTCATCACCTACACGTTTCATTCGACTTGGGTGACCAGTGAAGCCTATTCTTCTCCCTCCATTGTACTG  
TCTGCTCGTGGTGGGATGGCAGTAGGATCATTTTTGATGACTCCGAGAAGCGTATTATTGGCTCCGTC  
ACAATACTCCAGAGGATGCAAAAGTCATGTCATGGTGGGATTATGGCTACCAAATTAAGTCAATGGCAAA  
TCGGACAATTTAGTGGACAATAACACATGGAATAATACCCATATTTCTCGAGTAGGGCAGGCAATGGCA  
TCCACAGAAGAAAAAGCCTATGAAATCATGAGGGAGCTTGTATGTCAGCTATGTGCTTGTCAATTTTGGAG  
GCCTTACTGGGTATTCTTCGGATGATATCAACAAGTTTCTTTGGATGGTCCGGATTGGAGGAAGCACAGA  
GACAGGAAGACATTAAGGAGAATGACTACTATACTCCTACTGGGAATTCCTGTTGATCGTGAGGGT  
TCTCCGGTGTGCTCAACTGCCTTATGTACAAAATGTGTTACTACCGCTTTGGGCAGGTCTACACAGAAG  
CCAAGCGTCCACCAGGCTTTGACCGTGTTCGAAATGCTGAGATTGGTAATAAAGACTTTGAGCTTGATGT  
CCTGGAGGAAGCGTATACCACAGAACACTGGCTAGTCAGGATATACAAGGTAAGGACCTGGATAATCGA  
GGCTTGCAAGGACA

**ACGCGT**ACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
ACAAGGATGACGACGATAAGGTTTAA

**Protein Sequence:** >MR210156 protein sequence  
 Red=Cloning site Green=Tags(s)

MTKLGFLRLSYEKQDTLLKLLILSMAAVLSFSTRLFAVLRFEFSVIHEFDPYFNRYTRTRFLAEEGFYKFHN  
 WFDDRAWYPLGRIIGGTIYPGLMITSAAIYHVLHFFHITIDIRNVCVFLAPLFSSTTIVTYHLTKELKD  
 AGAGLLAAMIIVPGYISRSVAGSYDNEGIAIFCMLLTYYMWIKAVKGTGSIYAAKCALAYFYMVSSWG  
 GYVFLINLIPLHVLVLMLTGRFSHRIYVAYCTVYCLGTILSMQISFVGFQPVLSSEHMAAFGVFGLCQIH  
 AFVDYLRSKLNPQQFEVLFRSVISLVGFVLLTVGALLMLTGKISPWTGRFYSLLDPSYAKNNIPIIASVS  
 EHQPPTWSSYYFDLQLLVFMFPVGLYYCFSNLSDARIFIIMYGVTSMYFSAVMVRLMLVLAPVMCILSGI  
 GVSQVLSTYMNKLDISRDKSKKQDSTYPIKNEVASGMILVMAFFLITYTFHSTWVTSEAYSSPSIVL  
 SARGGDGSRIFDDFREAYYWL RHNTPEDAKVMSSWDYGYQITAMANRTILVDNNTWNNTHSRVQGAMA  
 STEEKAYEIMRELDVSYVLVIFGGLTGYSSDDINKFLWMVRIGGSTETGRHIKENDYYTPTGEFRVDREG  
 SPVLLNCLMYKMCYYRFQVYTEAKRPPGFDRVRNAEIGNKDFELDVLEEAYTTEHWLVRIYKVKDLNDR  
 GLSRT

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

**Restriction Sites:** SgfI-MluI

**Cloning Scheme:**

Cloning sites used for ORF Shuttling:



\* The last codon before the Stop codon of the ORF

**ACCN:** NM\_008408

**ORF Size:** 2118 bp

**OTI Disclaimer:** The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

**OTI Annotation:** This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

**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:** [NM\\_008408.1](#)

**RefSeq Size:** 2697 bp

**RefSeq ORF:** 2118 bp

**Locus ID:** 16430

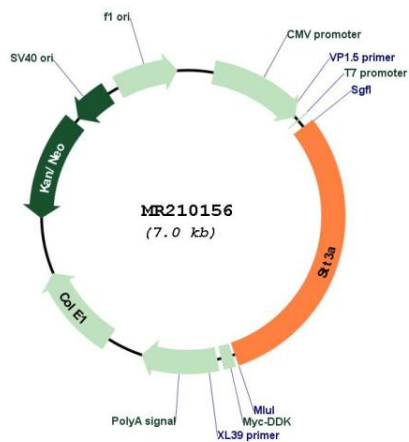
**UniProt ID:** [P46978](#)

**Cytogenetics:** 9 20.67 cM

**MW:** 80.6 kDa

**Gene Summary:** Catalytic subunit of the oligosaccharyl transferase (OST) complex that catalyzes the initial transfer of a defined glycan (Glc(3)Man(9)GlcNAc(2) in eukaryotes) from the lipid carrier dolichol-pyrophosphate to an asparagine residue within an Asn-X-Ser/Thr consensus motif in nascent polypeptide chains, the first step in protein N-glycosylation. N-glycosylation occurs cotranslationally and the complex associates with the Sec61 complex at the channel-forming translocon complex that mediates protein translocation across the endoplasmic reticulum (ER). All subunits are required for a maximal enzyme activity. This subunit contains the active site and the acceptor peptide and donor lipid-linked oligosaccharide (LLO) binding pockets (By similarity). STT3A is present in the majority of OST complexes and mediates cotranslational N-glycosylation of most sites on target proteins, while STT3B-containing complexes are required for efficient post-translational glycosylation and mediate glycosylation of sites that have been skipped by STT3A (By similarity).[UniProtKB/Swiss-Prot Function]

Product images:



Circular map for MR210156