

32-6916: ST6GAL1 Human, sf9

Alternative Name :

ST6 Beta-Galactoside Alpha-2,6-Sialyltransferase 1, ST6 Beta-Galactosamide Alpha-2,6-Sialyltransferase 1, ST6Gal I, CMP-N-Acetylneuraminate-Beta-Galactosamide-Alpha-2,6-Sialyltransferase 1, B-Cell Antigen CD75, Alpha 2,6-ST 1, EC 2.4.99.1, ST6GalI, SIAT1, CMP-N-Acetylneuraminate Beta-Galactosamide Alpha-2,6-Sialyltransferase, Sialyltransferase 1 (Beta-Galactoside Alpha-2,6-Sialyltransferase) , ST6 N-Acetylgalactosaminide Alpha-2,6-Sialyltransferase 1, Sialyltransferase 1, ST6N.

Description

Source: Sf9, Baculovirus cells.

Sterile Filtered colorless solution.

ST6GAL1 also known as ST6 Beta-Galactosamide Alpha-2,6-Sialyltransferase 1, is part of the glycosyltransferase family 29. ST6GAL1 is a type II membrane protein which catalyzes the transfer of sialic acid from CMP-sialic acid to galactose-containing substrates. Furthermore, ST6GAL1 is normally found in the Golgi however it can be proteolytically processed to a soluble form, ST6GAL1 is also involved in the generation of the cell-surface carbohydrate determinants as well differentiation antigens HB-6, CD75, and CD76.

ST6GAL1 produced in Sf9 Baculovirus cells is a single, glycosylated polypeptide chain containing 389 amino acids (27-406 a.a.) and having a molecular mass of 44.6kDa (Migrates at 40-57kDa on SDS-PAGE under reducing conditions).

Product Info

Amount :	2 µg / 10 µg
Purification :	Greater than 90.0% as determined by SDS-PAGE.
Content :	ST6GAL1 protein solution (0.5mg/ml) contains Phosphate Buffered Saline (pH 7.4) and 10% glycerol.
Storage condition :	Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°C for longer periods of time. For long term storage it is recommended to add a carrier protein (0.1% HSA or BSA). Avoid multiple freeze-thaw cycles.
Amino Acid :	ADPKEKKKGS YYDSFKLQTK EFQVLKSLGK LAMGSDSQSV SSSSTQDPHR GRQTLGSLRG LAKAKPEASF QVWNKDSSSK NLIPRLQKIW KNYLSMKNKYK VSYKGGPGPI KFSAEALRCH LRDHVNVMV EVTDFPFNTS EWEGYLPKES IRTKAGPWGR CAVVSSAGSL KSSQLGREID DHDVAVLRFNG APTANFQQDV GTKTTIRLMN SQLVTTEKRF LKDSLNEGI LIVWDPSVYH SDIPKWYQNP DYNFFNNTYKT YRKLHPNQPF YILKPQMPWE LWDILQEISP EEIQPNPPSS GMLGIIMMT LCDQVDIYEF LPSKRKTDVC YYYQKFFDSA CTMGAYHPLL YEKNLVKHLN QGTDEDIYLL GKATLPGFRT IHCHHHHHH.