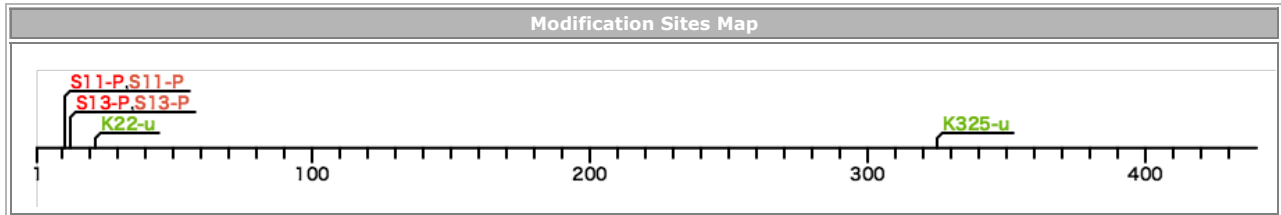


ID	Accession	GeneName	Chr.No.		Description
ALG3_HUMAN	Q92685	ALG3	3q27.1	183960089..183967336	Dol-P-Man:Man(5)GlcNAc(2)-PP-Dol alpha-1,3-mannosyltransferase



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
11	S	P	Lab	110218_pOVMANA_2.mgf[F017467]
11	S	P	Lab	140320_HEK_ME_.mgf[F017427]
11	S	P	Lab	140320_OVISE_ME_.mgf[F017429]
11	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
11	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence

MAAGLRKRGR **S**GSAAQAEGL **C**KQWLQRAWQ ERRLLLREPR YTLVAACLC LAEVGITFWV IHRVAYTEID WKAYMAEVEG VINGTYDYTQ LQDGTGPL VY PAGFVYIFMG LYYATSRGTD IRMAQNIFAV LYLATLLLVF LIYHQTCKVP PFVFFMCCA SYRVHSIFVL RLFNDPVAMV LLFLSINLLL AQRWGWGCC F FSLAVSVKMN VLLFAPGLLF LLLTQFGFRG ALPKLGICAG LQVVLGLPFL LENPSGYLSR SFDLGRQFLF HWTVNWRFLP EALFLHRAFH LALLTAHLTL LLLFALCRWH RTGESILSLL RDPS**K**RKVPP QPLTPNQIVS TLFTSNFIGI CFSRSLHYQF YVWYFHTLPY LLWAMPARWL THLLRLLVLG LIELSWNTYP S TSCSSAALH ICHAVILLQL WLGPPFPKKS TQHSKKA

Backcolor of amino acid : Yellow -> site of modification, gray -> in front of processing