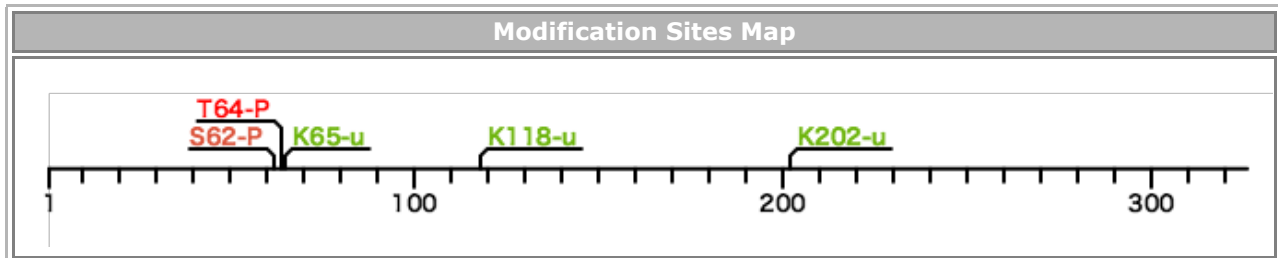


ID	Accession	GeneName	Chr.No.		Description
ALG5_HUMAN	Q9Y673	ALG5	13q13.3	37523912..37574398	Dolichyl-phosphate beta-glucosyltransferase



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
64	T	P	Lab	130327_HEK_ME_pphos.mgf[F015008]

**Protein Sequence**

MAPLLQLAV LGAALAAAAL VLISIVAFTT ATKMPALHRH EEEKFFLNAK GQKETLPSIW DSPTKQLSVV VPSYN  
 EEKRL PVMMDEALSY LEKRQKRDP AFTYEIVVDD GSKDQTSKVA FKYCQKYGSD KVRVITLVKN RGKGGAIR  
 MG IFSSRGEKIL MADADGATKF PDVEKLEKGL NDLQPWPNQM AIACGSRAHL EKESIAQRSY FRTLLMYGFH F  
 LVWFLCVKG IRDTQCGFKL FTREASRTF SSLHVERWAF DVELLYIAQF FKIPAEIAV NWTEIEGSKL VPFWSW  
 LQMG KDLLFIRLRY LTGAWRLEQT RKM

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