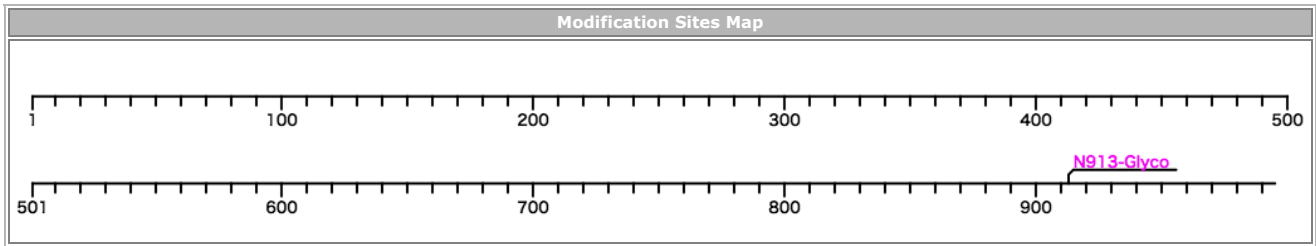


ID	Accession	GeneName	Chr.No.		Description
EMC1_HUMAN	Q8N766	EMC1	1p36.13	19542158..19578046	ER membrane protein complex subunit 1



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
913	N	Glyco	Lab	130626_Gly_S_MGFPeaklist.mgf[F015371]
913	N	Glyco	Lab	130626_Gly_Y_MGFPeaklist.mgf[F015373]
913	N	Glyco	Lab	130703_Gly_Y_re_MGFPeaklist.mgf[F015388]

Protein Sequence	
MAAEWASRFW LWATLLIPAA AVYEDQVGKF DWRQQYVGKV KFALEFSPG SKKLVVATEK NVIAALNSRT GEILWRHVDK GTAEGAVDAM LLHGQDVITV SNGGRIMRSW ETNIGGLNWE ITLDSGSFQA LGLVGLQESV RYIAVLKTT LALHHLSSGH LKWVHELPE DSIHYQMVYS YGSGVWVALG VVPFHVNI V KFNVEDGEIV QQVRVSTPWL Q HLSGACGVV DEAVLCPDP SSRSLQTLAL ETEWELRQIP LQSLDLEFGS GFQPRVLTQ PNPVDASRAQ FFLHLSPSHY ALLQYHYGTL SLLKNFPQTA LVSFATTGEK TAAAV MACRN EVQKSSSED GSMGSFSEKS SSKDSLACFN QTYTINLYLV ETGRRLLDTT ITFSLEQSGT RPERLYIQVF LKKDDSVGYR ALVQTEDHLL LFLQQLAGKV VLWSREES LA EVVCLEMVDL PLTGAQAELE GEFGKKADGL LGMFLKRLSS QLILLQAWTS HLWKMFYDAR KPRSQIKNEI NIDTLARDEF NLQKMMVMVT ASGKLFGIES SSGTILWKQY LPNVKPDSSF KLMVQRRTAH FPHPPQCTLL VKDKESGMSS LYVFNPIFGK WSQVAPPVLK RPILQSLLLP VMDQDYAKVL LLIDDEYKVT AFPATRNVLRL QHHELAPSIF FYLVD AEQGR LCGYRLRKDL TTELSWELTI PPEVQRIVKV KGRSSEHVH SQGRVMGDRS VLYKSLNPNL LAVVTESTDA HHERTFIGIF LIDGVTGRII HSSVQKKAKG PVHIVHSE NW VVYQYWNTKA RRNEFTVLEL YEGTEQYNAT AFSSLDRPQL PQVLQQSYIF PSSISAMEAT ITERGITSRH LLIGLPSGAI LSLPKALLDP RRPEIPEQS REENLIPYSP DVQ IHAERFI NYNQTVSRMR GIYTAPSGLE STCLVVAYGL DIYQTRVYPS KQFDVLKDDY DYVLSSVLF GLVFATMITK RLAQVKLLNR AWR	

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