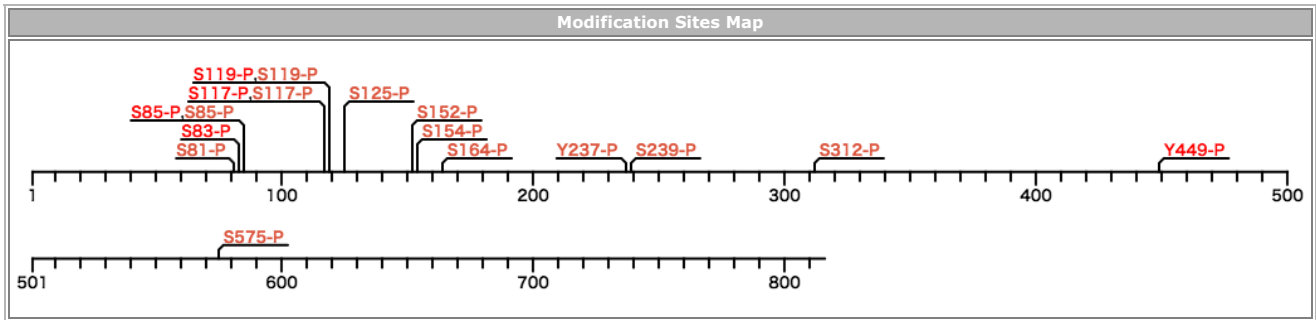


ID	Accession	GeneName	Chr.No.		Description
EIF3B_HUMAN	P55884	EIF3B	7p22.3	2393721..2420380	Eukaryotic translation initiation factor 3 subunit B



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
83	S	P	Lab	130327_HEK_ME_pphos.mgf[F015008]
83	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
83	S	P	Lab	130415_HEK_ME_tphos.mgf[F015010]

Protein Sequence

MQDAENVAVP EAAEERAEPG QQQPAAEPPP AEGLLRPAGP GAPEAAAGTEA SSEEVGIAEA GPESVTRTEP AAEAEAASGP SESPPAAE ELPGSHPAEP VPAQGEAPGE QA
 RDERSDSR AQAVSEDAGG NEGRAAEAE P RALENGDADE PSFSDPEDFV DDVSEELLG DVLKDRPQEA DGIDSVIVVD NVPQVGPDLR EKLKNVIHKI FSKFGKITND FYPE
 EDGKTK GYIFLEYASP AHAVDAVKNA DGYKLDKQHT FRVNLFTDFD KYMTISDEWD IPEKQPFKDL GNLRYWLEEA ECRDQYSVIF ESGDRTSIFW NDVKDPVSIE ERARWT
 ETYV RWSPKGTyla TFHQRGIALW GGEKFKQIQR FSHQGVQLID FSPCERYLVT FSPLMDTQDD PQAIIIWDIL TGHKKRGFHC ESSAHWPFIK WSHDGGKFFAR MTLDTLSI
 YE TPSMGLLDKK SLKISGIKDF SWSPGGNIIA FWVPEDKDIP ARVTLMLQPT RQEIRVRNLF NVVDCKLHWQ KNGDYLCVKV DRTPKGTQGV VTNFEIFRMR EKQVPVDVVE
 MKETIIAFaw EPNGSKFAVL HGEAPRISVS FYHVKNNGKI ELIKMFDKQQ ANTIFWSPQG QFVVLAGLRS MNGALAFVDT SDCTVMNIAE HYMASDVEWD PTGRYVVTsv S
 WWSHKVDNA YLWTFQGRl LQKNNKDRFC QLLWRPRPPT LLSQEIQIKI KKDlKKYSKI FEQKDRLSQS KASKELVERR RTMMEDFRKY RKMAQELyME QKNERLELRG G
 VDTDELDSN VDDWEEETIE FFVTEEIIPL GNQE

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