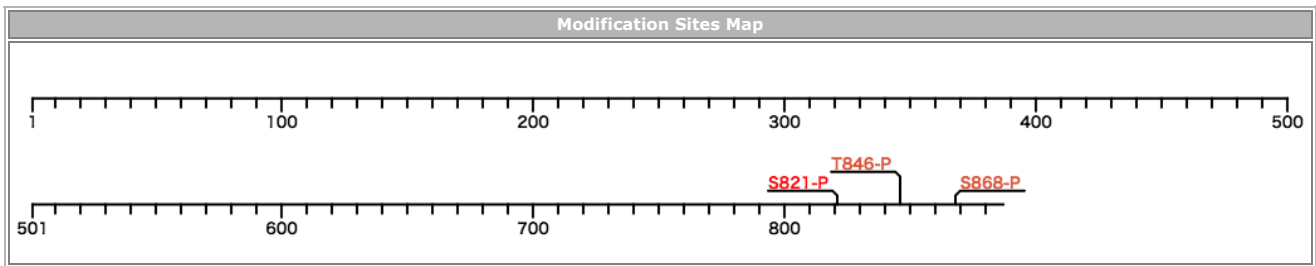


ID	Accession	GeneName	Chr.No.		Description
EXOSX_HUMAN	Q01780	EXOSC10	1p36.22	11126675..11159938	Exosome component 10



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
821	S	P	Lab	140320_tita_C18_.mgf[F017426]
821	S	P	Lab	140320_tita_SDB_.mgf[F017430]
821	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]

Protein Sequence

MAPPSTREPR VLSATSATKS DGEMVLPGF DADSFVKFAL GSVVAVTKAS GGLPQFGDEY DFYRSFPGFQ AFCETQGDRL LQCMSRVMQY HGCRSNIKDR SKVTELEDKF D LLVDANDVI LERVGILLDE ASGVNKNQQP VLPAGLQVPK TVVSSWNRKA AEYGGKAKSE TFRLLHAKNI IRPQLKFREK IDNSNTPFLP KIFIKPNAQK PLPQALSKER RERPQ DRPED LDVPPALADF IHQRTQQVE QDMFAHPYQY ELNHFTPADA VLQKPQPQLY RPIEETPCHF ISSLDELVEL NEKLLNCQEF AVDLEHHSYR SFLGLTCLMQ ISTRTEDFII DTLELRSDMY ILNESLTPA IVKVFHGADS DIEWLQKDFG LYVVMFDTH QAARLLNLGR HSLDHLKLY CNVDSNKQYQ LADWRIRPLP EEMLSYARDD THYLLIYDK MR LEMWERGN GQPVLQVWV QRSRDICLKK FIKPIFTDES YLELYRKQKK HLNTQQLTAF QLLFAWRDKT ARREDESYGY VLPNHMMLKI AEELPKPEQG IIACCNPVPP LVRQ QINEMH LLIQQAAREMP LLKSEVAAGV KKSGLPSAE RLENVLFQPH DCSHAPPDGY PIIPITSGSVP VQKQASLFPD EKEDNLLGTT CLIATAVITL FNEPSAEDSK KGPLTVAQ KK AQNIMESFEN PFRMFLPSLG HRAPVSQAQK FDPSTKIYEI SNRWKLAQVQ VQKDSKEAVK KKAQEQTAAE EQAKEACKAA AEQAISVRQQ VVLENAAKKR ERATSDPRT T EQKQEKRLK ISKPKDPEP PEKEFTPYDY SQSDFKAFAG NSKSKVSSQF DPNKQTPSGK KCIAAKKIKQ SVGNKSMSPF TGKSDRGFRY NWPQR

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