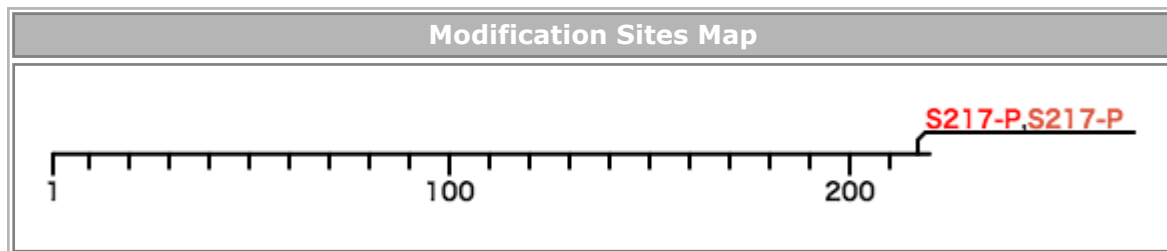


ID	Accession	GeneName	Chr.No.		Description
EIF3K_HUMAN	Q9UBQ5	EIF3K	19q13.2	39109735..39127595	Eukaryotic translation initiation factor 3 subunit K



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
217	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
217	S	P	Lab	140320_Agarose_.mgf[F017423]
217	S	P	Lab	140320_Agarose_.mgf[F017423]
217	S	P	Lab	110218_pRMUGS_2.mgf[F017480]
217	S	P	Lab	110218_pRMUGS_3.mgf[F017481]
217	S	P	Lab	110218_pRMUGS_4.mgf[F017482]
217	S	P	Lab	110218_pOVKATE_1.mgf[F017463]
217	S	P	Lab	110218_pOVKATE_3.mgf[F017465]
217	S	P	Lab	110218_pOVMANA_1.mgf[F017466]
217	S	P	Lab	110218_pOVMANA_3.mgf[F017468]
217	S	P	Lab	110218_pOVSAYO_2.mgf[F017470]
217	S	P	Lab	110218_pRMG1_2.mgf[F017473]
217	S	P	Lab	110218_pRMG2_2.mgf[F017476]
217	S	P	Lab	110218_pRMG2_3.mgf[F017477]
217	S	P	Lab	100627_akimura_pOVISE_3.mgf[F017443]
217	S	P	Lab	100628_akimura_pMCAS_2.mgf[F017455]
217	S	P	Lab	100628_akimura_pOVCAR3_2.mgf[F017458]
217	S	P	Lab	100628_akimura_pOVCAR3_2.mgf[F017458]
217	S	P	Lab	140320_tita_C18_.mgf[F017426]
217	S	P	Lab	140320_tita_C18_.mgf[F017426]
217	S	P	Lab	140320_HEK_SCE_.mgf[F017428]
217	S	P	Lab	140320_tita_SDB_.mgf[F017430]
217	S	P	Lab	140320_tita_SDB_.mgf[F017430]
217	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
217	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
217	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
217	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]

217	S	P	Lab	100520-GIST-W3.mgf[F017524]
217	S	P	Paper	Sci Signal 2009, 2(84), ra46

Protein Sequence

MAMFEQMRAN VGKLLKGIDR YNPENLATLE RYVETQAKEN AYDLEANLAV LKLYQFNPAF FQTTVTAQIL
LKALTNLPHT DFTLCKCMID QAHQEERPIR QILYLGDLLE TCHFQAFWQA LDENMDLLEG ITGFEDSVRK
FICHVVGITY QHIDRWLLAE MLGDLSDSQL KVMMSKYGWS ADESGQIFIC SQEESIKPKN IVEKIDFDSV
SSIMASSQ

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