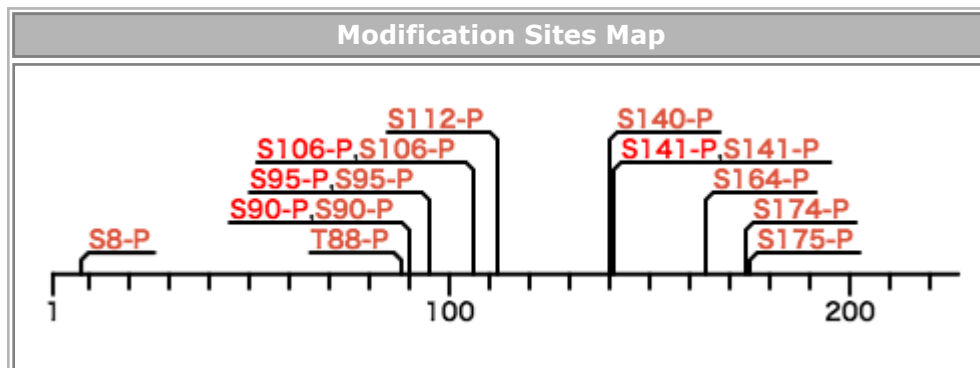


ID	Accession	GeneName	Chr.No.		Description
EF1B_HUMAN	P24534	EEF1B2	2q33.3	207024309..207027652	Elongation factor 1-beta



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
106	S	P	Lab	130327_HEK_ME_pphos.mgf[F015008]
106	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
106	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
106	S	P	Lab	140326_OVISE_NES_tita_2_.mgf[F017520]
106	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
106	S	P	Lab	100627_akimura_pRMG1_1.mgf[F017451]
106	S	P	Lab	100627_akimura_pRMG1_2.mgf[F017452]
106	S	P	Lab	140320_tita_C18_.mgf[F017426]
106	S	P	Lab	140320_tita_SDB_.mgf[F017430]
106	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
106	S	P	Lab	100520-GIST-IM1.mgf[F017509]
106	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
106	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
106	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
106	S	P	Lab	140326_OVISE_NES_tita_.mgf[F017518]
106	S	P	Lab	100520-GIST-W3.mgf[F017524]
106	S	P	Paper	Sci Signal 2009, 2(84), ra46
106	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence

MGFGDLK^SPA GLQVLNDYLA DKSYIEGYVP SQADVAVFEA VSSPPPADLC HALRWYNHIK SYEKEKASLP
GVKKALGKYG PADVEDT^TIG^S GATDS^SKDDDD IDLFG^SDD^EES^EEEAKRLRE ERLAQYESKK AKKPALVAK
^S^SILLDVKPWD DETDMAKLEE CVRS^SIQADGL VWG^SSS^SKLVPV GYGIKKLQIQ CVVEDDKVGT DML^EEEQI
TAF EDYVQSM^DVA AFNKI

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