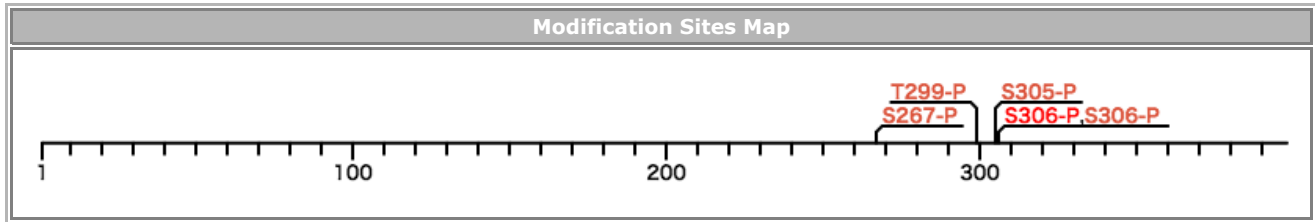


ID	Accession	GeneName	Chr.No.		Description
FL2D_HUMAN	Q15007	WTAP	6q25.3	160146617..160177351	Pre-mRNA-splicing regulator WTAP



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
306	S	P	Lab	140326_OVISE_NES_tita_2_.mgf[F017520]
306	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
306	S	P	Lab	140320_Agarose_.mgf[F017423]
306	S	P	Lab	110218_pRMUGS_2.mgf[F017480]
306	S	P	Lab	110218_pRMUGS_3.mgf[F017481]
306	S	P	Lab	110218_pRMUGS_4.mgf[F017482]
306	S	P	Lab	100628_akimura_pOVSAHO_1.mgf[F017460]
306	S	P	Lab	100628_akimura_pOVSAHO_2.mgf[F017461]
306	S	P	Lab	100628_akimura_pOVSAHO_3.mgf[F017462]
306	S	P	Lab	110218_pOVKATE_1.mgf[F017463]
306	S	P	Lab	110218_pOVKATE_2.mgf[F017464]
306	S	P	Lab	110218_pOVMANA_1.mgf[F017466]
306	S	P	Lab	110218_pOVMANA_2.mgf[F017467]
306	S	P	Lab	110218_pOVMANA_3.mgf[F017468]
306	S	P	Lab	110218_pOVSAHO_1.mgf[F017469]
306	S	P	Lab	110218_pOVSAHO_2.mgf[F017470]
306	S	P	Lab	110218_pOVSAHO_3.mgf[F017471]
306	S	P	Lab	110218_pRMG1_1.mgf[F017472]
306	S	P	Lab	110218_pRMG1_2.mgf[F017473]
306	S	P	Lab	110218_pRMG1_3.mgf[F017474]
306	S	P	Lab	110218_pRMG2_1.mgf[F017475]
306	S	P	Lab	110218_pRMG2_2.mgf[F017476]
306	S	P	Lab	110218_pRMG2_3.mgf[F017477]
306	S	P	Lab	110218_pRMG2_4.mgf[F017478]
306	S	P	Lab	110218_pRMUGS_1.mgf[F017479]
306	S	P	Lab	100627_akimura_pOVISE_1.mgf[F017437]
306	S	P	Lab	100627_akimura_pOVISE_2.mgf[F017440]
306	S	P	Lab	100627_akimura_pOVISE_3.mgf[F017443]
306	S	P	Lab	100627_akimura_pOVTOKO_1.mgf[F017447]
306	S	P	Lab	100627_akimura_pOVTOKO_2.mgf[F017449]
306	S	P	Lab	100627_akimura_pRMG1_1.mgf[F017451]
306	S	P	Lab	100627_akimura_pRMG1_2.mgf[F017452]
306	S	P	Lab	100627_akimura_pRMG1_3.mgf[F017453]
306	S	P	Lab	100628_akimura_pMCAS_1.mgf[F017454]
306	S	P	Lab	100628_akimura_pMCAS_2.mgf[F017455]
306	S	P	Lab	100628_akimura_pMCAS_3.mgf[F017456]
306	S	P	Lab	100628_akimura_pOVCAR3_1.mgf[F017457]
306	S	P	Lab	100628_akimura_pOVCAR3_2.mgf[F017458]
306	S	P	Lab	100628_akimura_pOVCAR3_3.mgf[F017459]

306	S	P	Lab	140320_tita_C18_.mgf[F017426]
306	S	P	Lab	140320_HEK_SCE_.mgf[F017428]
306	S	P	Lab	140320_tita_SDB_.mgf[F017430]
306	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
306	S	P	Lab	100520-GIST-IM1.mgf[F017509]
306	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
306	S	P	Lab	100520-GIST-IM2.mgf[F017512]
306	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
306	S	P	Lab	100520-GIST-IM3.mgf[F017514]
306	S	P	Lab	100520-GIST-R2.mgf[F017517]
306	S	P	Lab	140326_OVISE_NES_tita_.mgf[F017518]
306	S	P	Lab	100520-GIST-R3.mgf[F017519]
306	S	P	Lab	100520-GIST-W1.mgf[F017521]
306	S	P	Lab	100520-GIST-W2.mgf[F017522]
306	S	P	Lab	100520-GIST-W3.mgf[F017524]
306	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence

MTNEEPLPKK VRLSETDFKV MARDELILRW KQYEAYVQAL EGKYTDLNSN DVTGLRESEE KLKQQQQESA RRENILVMRL ATKEQEMQE
 C TTQIQYLKQV QQPSVAQLRS TMVDPAINLF FLKMKGELEQ TKDKLEQAQN ELSAWKFTPD SQTGKKLMAK CRMLIQENQE LGRQLSQ
 GRI AQLEAELALQ KKYSEELKSS QDELNDFIIQ LDEEVEGMQS TILVLQQQLK ETRQQLAQYQ QQSQASAPS TSRTTASEPV EQSEAT
 SKDC SRLTNGPSNG SSSRQRTSGS GFHREGNTIE DDFPSSPGNG NKSSNSSEER TGRGGSGYVN QLSAGYESVD SPTGSENSLT HQ
 SNDTDSSH DPQEEKAVSG KGNRTVGSRH VQNGLDSSVN VQGSVL

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