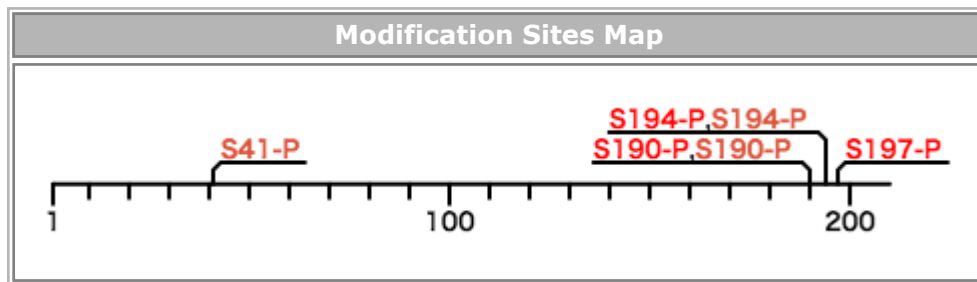


ID	Accession	GeneName	Chr.No.		Description
FADD_HUMAN	Q13158	FADD	11q13.3	70049269..70053496	FAS-associated death domain protein



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
194	S	P	Lab	130415_HEK_ME_tphos.mgf[F015010]
194	S	P	Lab	110218_pRMUGS_3.mgf[F017481]
194	S	P	Lab	110218_pRMUGS_4.mgf[F017482]
194	S	P	Lab	110218_pOVKATE_1.mgf[F017463]
194	S	P	Lab	110218_pOVKATE_3.mgf[F017465]
194	S	P	Lab	110218_pOVMANA_1.mgf[F017466]
194	S	P	Lab	110218_pOVMANA_2.mgf[F017467]
194	S	P	Lab	110218_pOVMANA_3.mgf[F017468]
194	S	P	Lab	110218_pOVSAYO_1.mgf[F017469]
194	S	P	Lab	110218_pOVSAYO_2.mgf[F017470]
194	S	P	Lab	110218_pOVSAYO_3.mgf[F017471]
194	S	P	Lab	110218_pRMG1_1.mgf[F017472]
194	S	P	Lab	110218_pRMG1_2.mgf[F017473]
194	S	P	Lab	110218_pRMG1_3.mgf[F017474]
194	S	P	Lab	110218_pRMG2_1.mgf[F017475]
194	S	P	Lab	110218_pRMG2_3.mgf[F017477]
194	S	P	Lab	110218_pRMG2_4.mgf[F017478]
194	S	P	Lab	100627_akimura_pOVICE_3.mgf[F017443]
194	S	P	Lab	100628_akimura_pOVCAR3_3.mgf[F017459]
194	S	P	Paper	Sci Signal 2009, 2(84), ra46

Protein Sequence
MDPFLVLLHS VSSSLSSSEL TELKFLCLGR VGKRKLERVQ S GLDLFSMLL EQNDLEPGHT ELLRELLASL R RHDLLRRVD DFEAGAAAGA APGEEDLCAA FNVICDNVGK DWRRRLARQLK VSDTKIDSIE DRYPRNLTER VRESLRIWKN TEKENATVAH LVGALRSCQM NLVADLVQEV QQARDLQNR S GAM S P S WNS DASTSEA S

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