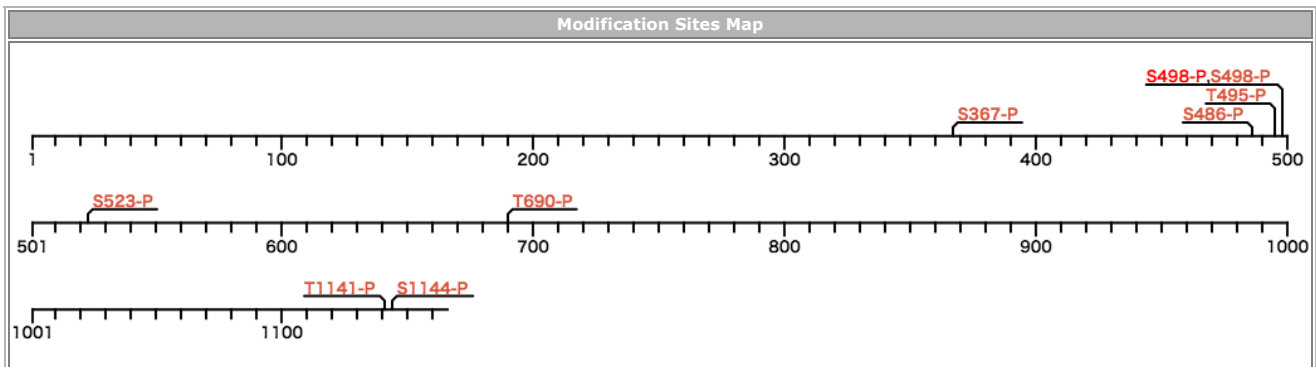


ID	Accession	GeneName	Chr.No.	Description
FHOD1_HUMAN	Q9Y613	FHOD1	16q22.1 67263290..67281561	FH1/FH2 domain-containing protein 1



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
498	S	P	Lab	100627_akimura_pOVICE_2.mgf[F017440]
498	S	P	Lab	100627_akimura_pOVTOKO_1.mgf[F017447]
498	S	P	Lab	100627_akimura_pRMG1_1.mgf[F017451]
498	S	P	Lab	100628_akimura_pMCAS_1.mgf[F017454]
498	S	P	Lab	100520-GIST-IM1.mgf[F017509]
498	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
498	S	P	Lab	100520-GIST-IM2.mgf[F017512]
498	S	P	Lab	100520-GIST-IM3.mgf[F017514]
498	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
498	S	P	Lab	100520-GIST-W1.mgf[F017521]
498	S	P	Lab	100520-GIST-W2.mgf[F017522]
498	S	P	Lab	100520-GIST-W3.mgf[F017524]
498	S	P	Paper	Cancer Res 2009, 69(6), 2663-2668
498	S	P	Paper	J Proteome Res 2008, 7(2), 526-534
498	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence

MAGGEDRDGDG EPVSVVTVRV QYLEDTPFA CANFPEPRA PTCSLDGALP LGAQIPAVHR LLGAPLKLED CALQVSPSGY YLDELSLEE QREMLEGFYE EISKGRKPTL ILRT QLSVRV NAILEKLYSS SGPELRRSLF SLKQIFQEDK DLVPEFVHSE GLSCLIRVGA AADHNYQSYI LRALGQLMLF VDGM LGVVAH SDTIQWLYTL CASLSRLVVK TALKLLLV FV EYSENNAPLF IRAVNSVAST TGAPPWANLV SILEEKNGAD PELLVYTVTL INKTLAALPD QDSFYDVTDA LEQQGMEALV QRHLGTAGTD VDLRTQLVLY ENALKLEDGD I EEAPGAGGR RERRKPSSEE GKRSRRS LEG GGCARAPPEP GPTGPASVPG PTSSTGPALL TGPASSVPGP PSGLQASVNL FPTISVAPSA DTSSERSIYK ARFLENVAAA ETEK QVALAQ GRAETLAGAM PNEAGGHPDA RQLWDS PETA PAART PQSPA PCVLLRAQRS LAPEPKEPLI PASKPAEPIW ELPTRAPRLS IGDLD FSDLG EDEDQDMLNV ESVEAG KDIP APSPLPLLS GVPPPPPLPP PPIKGFPP PPPLPLAAPL PHSVPDSSAL PTKRKT V KLF WRELKLAGGH GVSASRFGPC ATLWASLDPV SVDTARLEHL FESRAKEVLP SK KAGEGRR T MTTVLDPKRS NAINIGL T T L PPVHVIK AAL LNFDEFVSK DGIEKLLTMM PTEEERQKIE EAQLANPDIP LGPAENFLMT LASIGGLAAR LQLWAFKLDY DSMEREI AEP LFDLKVGM EQ LVQ NATFR CI LATLLAVGNF LNSQS S GFE LSYLEKVSEV KDTVRRQSL L HHLCSLV LQT RPESDLYSE IPALTRCAKV DFEQLTENLG QLERRSRAAE E SLRSLAKHE LAPALRARLT HFLDQCARRV AMLRIVHRRV CNRFHAFLLY LGYTPQAARE VRIMQFCHTL REFALRYTC RERV LQQQK QATYRERNKT RGRMITETEK FSGV AGEAPS NPSVPVAVSS GPGRGDADSH ASMKSLT SR PEDTTHNRRS RGMVQSSSPI MPTVGPSTAS PEEPPGSSLP SDTSDEIMDL LVQSVTKSSP RALAARERKR SRGNR KSLRR T L K S GLGDDL VQALGLSKGP GLEV

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