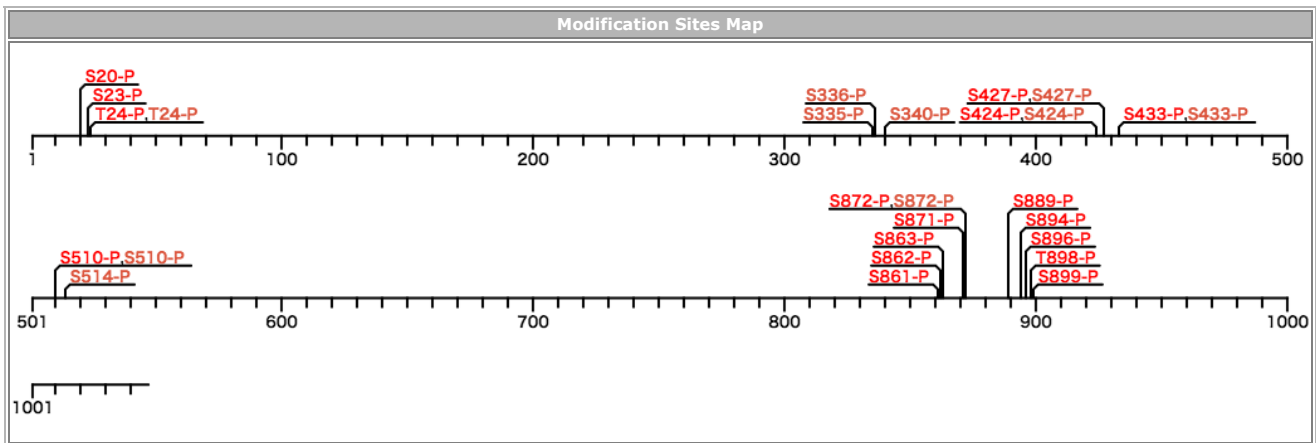


ID	Accession	GeneName	Chr.No.	Description
FARP1_HUMAN	Q9Y4F1	FARP1	13q32.2 98794816..99102027	FERM, RhoGEF and pleckstrin domain-containing protein 1



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
510	S	P	Lab	100628_akimura_pOVSAHO_1.mgf[F017460]
510	S	P	Lab	100628_akimura_pOVSAHO_2.mgf[F017461]
510	S	P	Lab	100628_akimura_pOVSAHO_3.mgf[F017462]
510	S	P	Lab	100520-GIST-IM1.mgf[F017509]
510	S	P	Lab	100520-GIST-R3.mgf[F017519]
510	S	P	Lab	100520-GIST-W2.mgf[F017522]
510	S	P	Lab	100520-GIST-W3.mgf[F017524]
510	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence	
MGEIEQRPTP GSRLGAPEN	S GIST LERGQK PPPTPSGKLV SIKIQMLDDT QEAFVFPQRA PGKVLDDAVC NHLNLVEGDY FGLEFPDHKK ITVWLDLLKP IVKQIRRPKH VVVK FVVKFF PPDHTQLQEE LTRYLFALQV KQDLAQGRLT CNDTSAALLI SHIVQSEIGD FDEALDREHL AKNKYIPQQD ALEDKIVEFH HNHIGQTPAE SDFQLLEIAR RLEMYGIRL H PAKDREGTKI NLAIVANTGIL VFQGFTKINA FNWAKVRKLS FKRKRFLIKL RPDANSAYQD TLEFLMASRD FCKSFWKICV EHHAFFRLFE EPKPKPKPVL FSRGSSFRFS GRT QKQVLDY VKEGGHKKVQ FERKHSKIHS IRSLASQPT E LNSEVLEQSQ QSTSLTFGEG AESPGGQSCR RGKEPKVSAG EPGSHSPAP RRS PAGNKQA DGAASAPTEE EEEV VKDRTQ QSKPQPPQPS TGSLTGSPHL SELSVNSQGG VAPANVTLSPL NLSPTDKQAS PLISPLLNDQ ACPRTDDEDE GRRKRFPDCK AYFIAKEVST TERTYLKDL E VITSWFQ STV SKEDAMPEAL KSLIFPNFEP LHKFHTNFKL EIEQRLALWE GRSNAQIRDY QRIGDVMLKN IQGMKHLAAH LWKHSEALEA LENGIKSSRR LENFCRDFEL QKVCYLPLNT FLLRPLHRLM HYKQVLERLC KHHPPSHADF RDCRAALAEI TEMVAQLHGT MIKMENFQKL HELKKDLIGI DNLVVPGREF IRLGSLSKLS GKGLQQRMMFF LFNDVLLYTS RGLT ASNQFK VHGQPLPYGM TIEESEDEWG VPHCLTLRGQ RQSIIVAASS RSEMEKWVED IQMAIDLAEK SSSPAPEFLA SSPDNKSPD EATAADQES E DDL SASRTSL ERQAPH RGNT MVHVCWHRNT SVSMVDFSIA VENQLSGNLL RKFKNNGWQ KLVVVFNTFC LFFYKSHQDN HPLASLPLLG YSLTIPSESE NIQKDYVFKL HFKSHVYFR AESEYTFE RW MEVIRSATSS ASRPHVLSHK ESLVY

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