

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
F1MAP4	F1MAP4_RAT	Rattus norvegicus	NaN	23.439218	NaN	NaN	38843836	MAISSSSCLGLICSLCHWLRAASSL NLEDPNVCSHWESYSVTVQESYPHP FDQIYTSCTDILNWFKCTRHRVSYR TAYRHGEKTMYYRKSQCCPGFYESR DVCVPHCADKCVHGRCIAPNTCQCE PGWGGTNCSSACDGDHWGPHCSS RCQCKNRALCNPITGACHCAAGYRG WRCEDRCEQGTYGNDCHQRCQCQ NGATCDHITGECRCSPGYTGAFCE LCPPGKHGQCEQRCPCQNGGVCH HVTGECSCPSGWMGTVCGQPCPEG RFGKNCSEQCQHNGGACDAATGO CHCSPGYTGERCQDECPVGTGVRC AETCRVNGGKCYHVSGLCEAGF SGEFCEARLCPEGLYGIKDKRCP HLDNTHSCHPMSGECGCKPGWSGL YCNETCSPGFYGEACQIQCSQNGA DCDSVSGRCACAPGFKGTDCSTPCP LGRYGINCSSRCGCKNDAVCSPVDG SCICKAGWHGVDCSISCPSTWGFG CNLTCQCLNGGACNTLDGTCTCAP GWRGEKCEFPQDGTYGLNCAERC DCSHADGCHPTTGHCRCLPGWSGV HCDSVCAEGRWGPNCSLPCYCKNG ASCSRDDGICECAPFRGTTQQRICS PGFYGHRCSTQCPQCVHSSGPCHHI TGLCDCLPGFTGALCNEVCPSGRFG KNCAGVCTTNGTGNPIDRSCQCY PGWIGSDCSQPCPPAHWGPNCIHTC NCHNGAFCSAYDGECKCTPGWTGL YCTQRCLPGFYGKDCALVCQCNNGA DCDHISGQCTCRTGFMGRHCEQKC PAGTYGYGCRQICDCLNNSTCDHIT GTCYSPGWKGARCDQAGVIVGNL NSLSRTSTALPADSYQIGAIAGIVLL LVVFLFLLFVYRHKQKRKESMPA VYTPAMRVINADYTIAETLPHSNGG NANSHYFTNPSYHTLSQCAASPHVN NRDRMTIAKSKNNQLFVNKKNVNP GKRGALVDCTGTLPADWKQGGYLN ELGAFGLDRSYMGSKSLDGLKNSY NSSTCSLSSSENPYATIKDPPALLPK SSECGYVEMKSPARRDSPYAEINNS TSANRNVYEVEPTVSVVQGVFSNNG HVTQDPYDLPKNSHIPCHYDLLPVR DSSSSPKREDGGSSSTTTSSNSN SSE	False	True	1.067	1.461	1.323	1.068	0.575	4.727	2.691