

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
F1LVL5	F1LVL5_RAT	Rattus norvegicus	NaN	23.439218	NaN	NaN	38843836	MENRPGSFQYVPVQLQGGAPWGFT LKGGLEHCEPLTVSKIEDGGKAALS QKMRTGDELVNINGTPLYGSRQEAL ILIKGSFRILKLVRRRNTVPSRPHSW HVAKLLEGCPDVATTMHFPSEAFSL SWHSGCNTSDVSVQWCPLSRHCST EKSSSIGSMESLEQPGQATYEGHLL PIEQNMYPQRSAYSFSASSNAS DCALSLKPEEPPSTDVCMMSGPGTK VTDDQANVSETSGSSHSTSGGHMT STSHASSCSEEGHHSGPAKMVRGPP EPPVRRDSLQASRAQLLNGEQHRAS EPVDSLQKQEKPALETMLPPRSSNQ FCCISGQDQVTGEDHQNCESLKPSE SSQDDCDHLLIEGSSKALDSPKAHD KGSNKEFGLLKEASADLANTLSFGAI PHLRGSIHRHSAPELLASHLQQM HLDNRGNKGMELPTGQDGHQWTM SPLHNNPKGKKSPLPTGGTQDQTS KERRIKPTDDKLMVSANQSQSDALL GEVDGHPSEPTYLDSNRASRASSDL TGQQPSATCSPVQOTRDFFSANKVV GHTEATEEGDNEPKEGGRLGRRS GGPRGRSIQNRKSERFATNLRNEI QRRKAQLQKSKGPLSQLCDTKEDVE ETQEPPESPPLPASNSLLPSYKNVP SHGDQLFNKSMILRARSSECLSQAS ESSKARVGLGRMSPGQRSGQSSLA LNTWWKASDSSSTDTEKANAHHG CRGHWWRWSPHNAQPOVALSTEGP SNPDDNKELKIPTQAGDEAILMPF ADRRKFFEESSKSLTSHLPGLTTH NKPFQIRPKPIDQNFQSVYRELKCH PMDQSYHSADQPYHAADQSYHSL PLQSEAPTYPDCFAPKGVDNSLYCK PVHHGDCDYHRTCSHPCSAQGTVM HDPCCICSGEICPALLKRNLKPKCH NCRCHHHQCIRCTGCCSHPOHGAH EDSSMASGNAWKSRKAAIQEFPVVK WKPIGNRKTSHSGREMAHSKANL SLSAPFRPHIENSALDLSNYRAVSSL DLLGDFKHASNTPEESSVYEDESSV ASMPRPLRSRAFSESHISLEPQNTRA WGKQQRRESFSGKSETQPDTLGARK KAFPPRPPPPNWEKYRLFRAAQQQ QQQQQQQQQQQQQQQQQCEKEEE EEKEQEEKGEKEDLPPQYFSSETT GSCAPITEEQPQSLKMGHQEASRQS SQLQEQEAYAVHPSNFVPAVRSYP VTQPEKAQPPCYGGAHGLWRITTEQE ATVTPKQEFQHFSPPTGASGIPTSYS AYYNISVAKAELLNKLKQPMAEAE GLGEEGVYELAQKKIQLIESISRKLS VLREAQRGLLDDINANAALGEEVEV NLKAVCKSNELEKYHLFIGDLKVV NLLLSLGRLARVENALNSIDAESN QEKLVLRKQQLTNQLADAKELKE HVDREKLVFSMVSRYLPQDQLQDY QHFVKMKSALIIEQRELEEKIKLGEE QLKCLKESLHLGPSNF	False	False	2.77	3.896	False	False	4.0	False	