

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q7Z7B0	FLIP1_HUMAN	Homo sapiens	Filamin-A-interacting protein 1	26.156049	S31;S33;S1104;T1130;T1135	S138;S979	30379171;30620550;28657654;29485866	MRSRNQGGESASDGHISCPKPSIIG NAGEKLSLEDKAKKKKSNRKEDDV MASGTVKRHLKTSGECKERKTKKSLE LSKEDLIQLLSIMEGELQAREDVIHM LKTEKTKPEVLEAHYGSAAEPEKVLRV LHRDAILAQEKSIGEDVYEKPISELD RLEEKQKETYRRMLEQLLAAEKCHR RTVYELENEKHKHTDYMNKSDDF NLLEQERERLKKLLEQEKAYQARKE KENAKRLNKLRLDELVKLKSFALMLV DERQMHIEQLGLQSQKVQDLTQKL REEEELKKAITSKSKEDRQKLLKLEV DFEHKASRFSQEHEEMNAKLANQE SHNRQLRLKLVGLTQRIEELEETNK NLQKAEFEELQELRDKIAKGECCGNSS LMAEVENLRKRVLMEGKDEEITKT ESQCRELRKKLQEEHHHSEKELRLEV EKLQKRMSELEKLEEAFAFSKSECT QLHLNLEKEKNLTKDLLNELEVVS RVKELECSERLEKAEKSLKDDLTKL KSFTVMLVDERKNMMEKIKQEERK VDGLNKNFKVEQGVMDVTEKLIIE SKKLLKSEMEEKVYNLTREDELI GKLKSEEEKSSELSCSVLLKRLD GIEEVEREITRGRSRKGSSELTCPEDN KIKELTLEIERLKKRLQLEVEGDL MKTEDEYDQLEQKFRTEQDKANFL SQQLEIKHQIAKNAIEKGEVVSQE AELRHRFRLEEAKSRDLKAEVQALK EKIHELMNKEDQLSQLQVDYSVLQQ RFMEEENKNKNMGQEVNLNLTKELE LSKRYSRALRPSVNGRRMVDVPTS TGVQTDVAVSGEAAEEETPAVFIKSF QEENHIMSINLRQVGLKKPVERSSVL DRYPPAANELTMRKSWIPWMRKRE NGPSITQEKGPRTNSSPGHPGEVVL SPKQGQPLHIRVTPDHENSTATLEIT SPTSEEFFSSTTVIPLGNQKPRITIP SPNVMPQKQKSGDTTLGPERAMSP VTITTFSSREKTPESGRGAFADRPTSPI QIMTVSTSAAPAEIAVSPESQEMPM GRTILKVTPEKQTVPTPVRKYNSNAN IITTEDNKIHIHLGSGQFKRSPGTSGE GVSPVITVRPVNVTAKEVSTGTVLR SPRNHLSSRPGASKVTSTITITPVTT SARGTQSVSGQDGSSQRPTPTIPM SKGMKAGKPVVAAPGAGNLTKEFPR AETQSMKIELKKAASSTTSLGGGK G	True	False	4.444	4.62	0.784	0.731	0.819