

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	ext reg
Q6P9Q6	FKB15_MOUSE	Mus musculus	FK506-binding protein 15	35.493921	S306;S344	S14;S23;S306;S310;S342;S344;S617;S948;S1018;S1050;S1091;T1093;S1108;S1153;S1157;S1159;S1190;T1198	40885482;39627609	MFGAGDEDDTDFLSPSGGAKLASLF GLDQATMGHGNEFFOYTAPKQPKK GOGTAAGNQITAPKPAPATTGTSSVL FATAVHAYRYINGQYAKQKGFGAAV LGNHTSREYRILLYISQQQPVTVATI HLNFELMVRPNNYSTFYDDQRQNW SIMFESEKAAVSNKQVCVAKCNSI SSLDAVLCODLVAAEGPAVETGDSL EVAYTGWLLQNHVLGQVFDSTANK DKPLRLKLGSGKVVKGLDGLGGMK KGGKRLIITPSACAAGSEGVIGWTQP TDSILVFEVEVRRVKFARDSGSDGH SVSSRDSAAPSPIPASDLSADPVVT PLPLPLKPGEPGLRSKNSLSEQLTV NSNPDTVKAKLISRMAMKMGQPMPLPI LPPQLDSNDSETEDATVLRGAGQSL VTPSIQPSLQPAHPVLPQMASQAPQ PSGSGLQTPSAALMQAVSLDSHSAV SGNAQNFQPYAGVQAYAYPQTPSVT SQLQPVPRPLYPAPLSQAPHFQSGSD MMSFLMTEARQHNTAIRMAVNVKVA DKMDHLMTKVEELQKHSSGNSMLL PSMSVTMETSMIMSNIQRIQENER LKQELLEKSSRIEENDKISDLIERN QRYVEQSNLMMMEKRNNSLQTATEN TQARILHAEQEKAKVTEELAAATAQV SHLQLKMTAHQKKETELQLQTDN LKETDLLRGHVTRLQADSELREAS EQTQTKFKSEKQSRROLELKVTSLE EELTDLRAEKTSLEKNLSERKKKSA QERCQAEAEEMDEIRKSHQEELDRLR QLLKKARVSTDQAAAEQLTLAQAE QSQWEAKCEQLLASARDEHLQOYR EVCAQRDAHQQKLALLQDECLALQA QIAAFTEQKEHMQRLEKTKSQAPAG RAAADPSEKVKKIMNQVFQSLRGEF ELEESYDGGTILRTIMHTIKMVTLQL LNHQEEEEEEEEEEEEKKPLRPSL EQPGPATPGMPPAPPSGETQEAPPEV LPEQVVGETTPLPLQALPTPENGAT RKGEPAEAEVPSSEIKDSSLPPQAGI PAHRVLGPPTSIPPKPPGPVMDSES EEMLAADQRTVQPNGLLGEHRE VATDGLLQGNRRRLSLTPDPEKGE PALDPESQGGEAQPPECKQAEDVSS SGPRETLLDTELASAAAAGTSLRHNO DSQHCSSLGDEEDELFGKATLKVPR PTAQPEEEDDEVSMKGRPPPTPLF GDDDDDDDDIGWLG	False	False	2.209	2.85	0.835	0.812	0.935	0.857	0.5