

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q6PAJ1	BCR_MOUSE	Mus musculus	Breakpoint cluster region protein	47.286703	T149;S150;S402	S122;S139;T178;S203;S216;S237;T247;S358;S379;S384;T387;S461;S465;S475;S487;T553;T640;T643;T692;S1263	22517741;33300544;40885482;36852467;35822049	MVDSVGF AEAWRAQFPDSEPPRME LRSVGDIEQE LERCKASIRRLLEQEVN QERFRMIYLQTL LAKEKKS YDRQRW GFRRAAQPPDGA AEPRASAPRPPPA PADGADPAPVEESEARP DGEGSPSK GRSASARRPAAAASADRDRGPPPTS VAALRSNF EKIRKGPAPG SADA EK PFYVNV E FHH ERGLVKVNDKEVSD RISSLGSQAMQMERKKSQQSAGQG LGEAPRPHYGRSSESSCGLDGDYE DAELNPRFLKDNLINANGGNRPPW PPLEYQPYQSIYVGGMMVEGEGKSP LLRSQSTSEQEKRLTWPRRSYSPRS FEDSGGGYTPDCSSNENLTSSEEDF SSGQSSRVSPSTTYRMRDKSRSP SQNSQQSFDSSPPTPQCQKRHRQC QVVVSEATIVGVRK TQGIWPSDGDST FQGEADSSFGT PPGYGCAADQAE EQ RRHQDGLPYIDDS PSSHLSKGR GSLASGALDPTKVSELDLEKGLEMR KVVLSGLASEETYL SHLEALLPMK PLKAAATTSQPVLT SQIETIFFKVPE LYEIHKEFYDGLFPRVQQW SHQQRV GDLFQKLASQLGVYRAFVDNYGVA METAEKCCQANAQFAEISENLRARS NKDVKDS TTKN SLETL LYPVDRVT RSTLVLDLLKHTPSSHDPHSLLDQ ALRISQNFLSSINEEITPRRQSM TVK KGEHRQLLKDSFMVELVEGARKLR HIFLFTDLLLCTKLKQSGGKTQQYD CKWYIPLTDL SFQMVDELEALNIPL VPDEELDALKIKISQIKSDIQREKRAN KGSKVMERLRK KLESEQLLLMSP SMAFRVHSRNGKSYTFLISSDYERA EWRESIREQQKCFKSFSLTSVELQ MLTNSCVKLQTVHHIPLTINKEDDE SPGLYGFLHVIVHSATGFKQSSNLYC TLEVDSFGYFVNKAKTRVYRDTTEP NWN EEF EIELEGSQTLRILCYEKCY NKMKMTKEDGESADKLMGKGQVQ LDPQTLQDRD WQRTVIDMNGIEVKL SVKFTSREFSLKRMP SRKQTGVFGV KIAVVTKRERSKVPYIVRQCVEEIERR GMEEVGIYRVSGVATDIQALKAAPDV NNKDVSMVMSEMDVNAIAGTLKLY FRELPEPLFTDEFYPNFAEGIALSDP VAKESCMLNLLLSLPEANLLTFLFLL DHLKRVAEKETV NKMSLHNLATVF GPTLLRPSEKESKLPANPSQPITMTD SWSLEVMSQVQLLYFLQLEAIPAP DSKRQSILFSTEV	True	True	5.0	3.003	2.421	2.238	1.91