

UniprotKB ID	Entry name	organism	full name	oglcnaescore	oglcnaesites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrace region
Q9EQ32	BCAP_MOUSE	Mus musculus	Phosphoinositide 3-kinase adapter protein 1	25.685855	NaN	T264;T420;T445;T460;T513;T553;T570;T594;S642;T694;S718	39627609	MAASGWGRGCDILIFYSFDAEEWCQ YLQDLFVSCRQVRSOKTQTYRLVDP ASFSAQDLWVFRDARCVLVLLSAGL VGCFGQPGLLPMLQRACHPPQRVV RLLCGVQPGDEDFQAFPPDWAHWQ EMTCDDPEPETYLA AVRKAISED SGC DSVTDTEPEDERELPF SKQTNLPPEI SPGNLMVVQPD RIRCGAETT VYIIVR CKLDEKVSTEA EFSPE D SPSIRVEGT LENEYTVSVKAPDLSSGNVSLKVYS GDLVVCETT VSYTDM EIGNLLSSA ANPVFEMCQAFKIVPYNTETLDKLL TESLKNNIPASGLHLFGINQLEEDD MMTNQRDEELPTLLHFAAKYGLKN LTALLLTCPGALQAYSVANKHGHP NTIAEKHGFRDLRQFIDEYVETVDM LKTHIKEELMQGEEADVYESMAHL STDLLMKCSLNP GCDD ELYESMAAF APAATEDLYVEMLQASAGNPVSGES FSRPTKDSMIRKFLGNSVKPASWE REQHPYGEELYHIVDEDETF SVDL ANRPPVPVPRPEASAPGPPPPDNEP YISKVFAEKSQERLGNFYVSESIRK EPLVRPWRDRPPSSIYDPFAGMKT GQRQLITLQEQVKLGIVNVDEAVLH FKEWQLNQKRSSEFRFQENLKR LRESITRRRKEKPKSGKHTDLEITVPI RHSQHLPEKVEFGVYESGPRKSVLP ARTELRRGDWKTDSMSSTASSTSNR SSTRSLLSVSSGMEGDNEDNEIPEIT RSRGPPTQVDGAPVVTGTVPVGTLE RPPRVPPRAASQRPLTRESFHPPPPV PPRGR	True	True	5.0	2.322	1.099	0.915	False	5.0	1.304