

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
Q9CQ73	PKP2_MOUSE	Mus musculus	Plakophilin-2	22.380232	NaN	S44;S82;S132;S135;S151;S154;S155;S172;S188;S232;S265;S287	37507081	MAVPGSLAECGYIRTVLGOQILGHL DSSSLALPSEARLRLAGSSGRGDPAAR RSQRIQEQVQOTLARRGRSSAVSGN LHRTSSVPEYVYKLVVENDFVGRQ SPVTRDYDMLKAGMTATYGSRWGR AAAQYSSQKSVEERSWRQPLRRLEI SPDSSPERAHYGHSEYQYAWRSHV PGGRLTLPRYARSEILGLRQAGTARR PPGCGSFSDAVFDNGPLKPTMPHTP PGTSHSAGSLEETTIVRVSQARLQST QSRTARSSWPRSSVRSSLREPGRML TTAGQAAVGSQDAHGDRSVFADAQL GNADIEMTLERAVNMLDADHVPVS KISAAATFIQHEFSQKSEARKRVNQL RGIPKLLQLLKLQNEVDVQRAACGAL RNLVFEEDNDNKLEVAELNGVPRLLQ VLKQTRDLETKKQITGLLWNLSSSD KCLKHLMITEALLTLTESVIIPFSGWP EGDYPKANGLLDFDIFYNVVTGCLR MSSAGPDGRKMMRRCDGLIDSLVH YVRGTIADYQPD DKATENCVCILHNL SYQLEAELPEKYSQSIYMQNRNIQT NSNKSIGCFGSRSRKLKEQYQDLQM PEERSNPHGIEWLWHSIVIRMYLSLI AKSTRNTQEASLGALQNLTAGGGP IPTLVARMVVQKENGLOHTRKMLH VGDPSVKKTAVSLLRNLNRNLSLQ EIAKETLPDLVSIIPDTVPSTDLLIETT ASACYTLNLMQNSYQNARDLLNT GGLQKIMTISIGEGYAPNKASKAASV LLYSLWAHTELHHAYKKAQFKKTD VNSRTAKAYHSLKD	True	True	3.473	4.558	1.757	2.107	0.879	4.431	1.669