

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracel region
Q8CIH5	PLCG2_MOUSE	Mus musculus	1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase gamma-2	25.685855	NaN	T753;T759;T1197;T1217;T1245	39627609	MTTMVNVDTLPPEYKESQIKRALELG TVMTVFNARKSTPERRIVQMIMETR QVAWSKTADKIEGFLDIMEIKEIRPG KNSKDFERAKAVRHKAECCTILYG TQFVLSLTLATDSKEDAVKWL SGL KILHQEAMSASTPTMIESWLRKQIYS VDQTRRNSISLRELKTLPLVNFKVS GIKFLKDKLVEIGAOKDELSFEQFHL FYKCLMFDQOKSILDEFKDDSSVFIL GNTDRPDASAVYLQDFQRFLHEQQ ELWAQDLNKVRRMTKFIDDTMRE TAEPLFVDFEFLTYLFSRENSIWDEK YDAVDMQDMNNPLSHYWISSHNT YLTGDQLRSESSTEAYIRCLRAGCRC IELDCWDGPDGKPIYHGWTRITTKIK FDDVVQAIRDHAFVTSSFPVILSIEE HCSVEQQRHMAKVFKEVLGDMMLT KPTASADQLPSPSQLREKIIKHKKL GPKGDVVDVNVEDKKDEHKPQGELY MWDSIDQKWTRHYCAIADAKLSFG DDIEQAVEEEPQDTPPTLHFGEK WFHKKVESRTSAEKLQEQYCAETGA KDGTFVRESETFPNDYTLFSWRSG RVQHCRIRSTMENGVMKYLLTDNL TFNSIYALIQHYREAHLRCAEFELRL TDPVNPNPHEKSPWYYDSLRSRGEA EDMLMRIPRDGAFILIRKREGTNSYAI TFRARGKVKHCRINRDGRHFVLGTS AYFESLVELVSYEKHALYRKMRLRY PVTPELLERYNMERDINSLYDVSRM YVDPSEINPSPMPQRTVKALYDYKAK RSDELTFCRGALIHNVSKPEGGW KGDYGTRIQQYFSPNYVEDISAGDAE EMEKQIIEDNPLGSLCKGILDNTYN VVKAPQKGNQKAFVFILEPKKQGD PVEFATDRVEELFEWFQSIREITWK MDTKENNMYWERNQSI AIELSDL VVYCKPTSKTKDHLENPDFREIRSFV ETKADSVRQKPVDLLRYNQKGLTR VYPKQQRVDSNYDPFRLWLCGSQ MVALNFQTPDKYMQMNHAFSLN GRTGYVLQPESMRSEKYDMPLESQ RKILMTLTVKVLGARHLPKLGRSIAC PFVEVEICGAEYDSNKFKTTVNDN GLSPVWAPTQEKVTFEYDPNLAFLR FVVEEDMFSDPNFLAHATYPIKGIK SGFRSVPLKNGYSEDIELASLLVFCE MRPVLESEELYSSCRLRRRQEEL NNQLFLYDTHQNLRGANRDALVKE FNVNENQLQLYQEKCNRRRLREKRV SNSRFYS	True	True	4.779	3.247	1.94	2.12	1.294	4.559	2.372