	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plas mem
0	P19174	PLCG1_HUMAN	Homo sapiens	1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase gamma-1	26.040516	S63	T506;T771;T775;T783;T977;S1221;S1227;S1233;S1248;T1253;S1263;S1222	34725712;29351928	MAGAASPCANGCGPGAPSDAEVLH LCRSLEVGTVMTLFYSKKSQRPERK TFQVKLETRQITWSRGADKIEGAIDI REIKEIRPGKTSRDFDRYQEDPAFRP DQSHCFVILYGMEFRLKTLSLQATS EDEVNMWIKGLTWLMEDTLQAPTP LQIERWLRKQFYSVDRNREDRISAK DLKNMLSQVNYRVPNMRFLRERLT DLEQRSGDITYGQFAQLYRSLMYSA QKTMDLPFLEASTLRAGERPELCRV SLPEFQQFLLDYQGELWAVDRLQVQ EFMLSFLRDPLREIEEPYFFLDEFVT FLFSKENSVWNSQLDAVCPDTMNN PLSHYWISSSHNTYLTGDQFSSESSL EAYARCLRMGCRCIELDCWDGPDG MPVIYHGHTLTTKIKFSDVLHTIKEH AFVASEYPVILSIEDHCSIAQQRNMA QYFKKVLGDTLLTKPVEISADGLPSP NQLKRKILIKHKKLAEGSAYEEVPTS MMYSENDISNSIKNGILYLEDPVNH EWYPHYFVLTSSKIYYSEETSSDQGN EDEEEPKEVSSSTELHSNEKWFHG KLGAGRDGRHIAERLLTEYCIETGAP DGSFLVRESETFVGDYTLSFWRNGK VQHCRIHSRQDAGTPKFFLTDNLVF DSLYDLITHYQQVPLRCNEFEMRLS EPVPQTNAHESKEWYHASLTRAQAE HMLMRVPRDGAFLVRKRNEPNSYA ISFRAEGKIKHCRVQQEGQTVMLGN SEFDSLVDLISYYEKHPLYRKMKLRY PINEEALEKIGTAEPDYGALYEGRNP GFYVEANPMPTFKCAVKALFDYKAQ REDELTFIKSAIIQNVEKQEGGWWR GDYGGKKQLWFPSNYVEEMVNPVA LEPEREHLDENSPLGDLLRGVLDVP ACQIAIRPEGKNNRLFVFSISMASVA HWSLDVAADSQEELQDWVKKIREV AQTADARLTEGKIMERRKKIALELSE LVVYCRPVPFDEEKIGTERACYRDM SSFPETKAEKYVNKAKGKKFLQYNR LQLSRIYPKGQRLDSSNYDPLPMWI CGSQLVALNFQTPDKPMQMNQALF MTGRHCGYVLQPSTMRDEAFDPFD KSSLRGLEPCAISIEVLGARHLPKNG RGIVCPFVEIEVAGAEYDSTKQKTEF VVDNGLNPVWPAKPFHFQISNPEFA FLRFVVYEEDMFSDQNFLAQATFPV KGLKTGYRAVPLKNNYSEDLELASL LIKIDIFPAKENGDLSPFSGTSLRERG SDASGQLFHGRAREGSFESRYQQPF EDFRISQEHLADHFDSRERRAPRRT RVNGDNRL	True	True	4.898	5.0	2.151	2.448	1.769	5.0