

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	extra regic
Q9Y678	COPG1_HUMAN	Homo sapiens	Coatomer subunit gamma-1	14.54405	S725	T594	38665916;35083852;33214551;23301498;30397120;34019948;34725712;32119511;30459710;37217939;29351928	MLKKFDKKDEESGGSNPFQHLEK SAVLQEARVFNETPINPRKCAHILTK ILYLINQGEHLGTTEATEAFFAMTKL FQSNDP TLRMCYLTIKEMSCIAED VIIVTSSLTKDMTGKEDNYRGPVAVRA LCQITDSTMLQAIERYMKQAIVDKVP SVSSSALVSSLHLLKCSFDVVKRWV NEAQEAASSDNIMVQYHALGLLYHV RKNDRLAVNKMISKVTRHGLKSPFA YCM MIRVASKQLEEDGSRDSPFLD FIESCLR NKHEMVVYEAASAVNLP CSAKELAPAVSVLQLFCSSPKAALRY AAVRTLNKVAMKHPSAVTACNLDLE NLVTD SNRSIATLAIITLLKTGSESSI DRLMKQISSFMSEISDEFKVVVVQAI SALCQKYPRKHAVLMNFLTMLREE GGFEYKRAIVDCIISHEENSEKETG LSHLCEFIEDCEFTVLATRILHLLGQ EGPKTTNPSKYIRFIYNRVLEHEEV RAGAVSALAKFGAQNEEMLPSILVL LKRCVMDDDNEVRDRATFYLVLE QKQKALNAGYILNGLTVSIPGLERAL QYTLPESEKPFDLKSVPLATAPMA EQRTESTPITAVKQPEKVAATRQEIF QEQLAAVPEFRGLGPLFKSSPEPVAL TESETEYVIRCTKHTFTNHMFQFD CTNTLNDQ TLENVTVQMEPT EAYEV LCYVPARSLPYNQPGTCYTLVALPKE DPTAVACTFSCMMKFTVKDCDPTTG ETDDEGYEDEYVLEDELEVTVADHIQ KVMKLNFEAAWDEVGDEFEEKEET TLSTIKTLEEAVGNIVKFLGMHPCER SDKVDPDNKNTHTLLLAGVFRGGHDI LVRSRLLLLDTVTMQVTARSLEELPV DIILASVG	True	True	4.623	2.355	1.06	4.323	4.451	1.449	1.203