

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 regio
P63010	AP2B1_HUMAN	Homo sapiens	AP-2 complex subunit beta	7.372219	S671;S672	S4;T737;T928	38665916;29351928;33214551;23301498;35138101;34019948;32119511;34725712;37217939;28510447	MTDSKYFTTNKKGEIFELKAELNNE KKEKRKEAVKKVIAAMTVGKDVSSL FPDVVNCMQTDNLELKKLVLYLML NYAKSQPDMAIMAVNSFVKDCEDP NPLIRALAVRTMGCIRVDKITEYLCE PLRKCLKDEDPYVRKTAAVCVAKLH DINAQMVEDQGFLDSLRLDIADSNP MNVANAVAALSEISESHPNSNLLDL NPQINIKLLTALNECTEWGQIFILD CLSNYNPKDDREAQSICERVTPRLS HANSAVVLSAVKVLKFLLELLPKDS DYNNMLLKKLAPPLVTLSSGPEVQ YVALRNINLIVQKRPEILKQEIKVFFV KYNDPIYVKLEKLDIMIRLASQANIA QVLAELKEYATEVDVDFVRKAVRAIG RCAIKVEQSAERCVSTLLDLIQTKVN YVVQEAIVIRDIFRKYPNKYESIATL CENLDSLDEPDARAAMIWIVGEYAE RIDNADELLESFLEGFHDESTQVQL LLTAIVKFLKKPSETQELVQVLS LATQSDNPDRLDRGYIYWRLLSTD PVTAKEVVLSEKPLISEETDLIEPTLL DELICHIGSLASVYHKPPNAFVEGSH GIHRKHLPIHHGSTDAGDSPVGT ATNLEQPQVIPSQGDLLGDLNLDL GPPVNVQVSSMQMGAVDLLGGGL DSLVGQSFIPSSVPATFAPSPTPAVVS SGLNDLFELSTGIGMAPGGYVAPKA VWLPVAVKAKGLEISGTFTHRQGHY MEMNFTNKALQHMTDFAIQFNKNS FGVIPSTPLAIHTPLMPNQSIDVSLPL NTLGPVMKMEPLNQLQVAVKNNID VFYFSLIPLNVLFVEDGKMERQVF LATWKDIPNENELQFQIKECHLNAD TVSSKLQNNVYTIAKRNVEGDML YQSKLKTNGIWLAELRIQGNPNYT LSLKCRAPEVSYQIYQVYDSILKN	True	True	4.66	1.749	1.359	1.358	1.808	5.0	1.57