	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	O55029	COPB2_MOUSE	Mus musculus	Coatomer subunit beta'	23.627151	NaN	S859	37453647	MPLRLDIKRKLTARSDRVKSVDLHP TEPWMLASLYNGSVCVWNHETQTL VKTFEVCDLPVRAAKFVARKNWVVT GADDMQIRVFNYNTLERVHMFEAH SDYIRCIAVHPTQPFILTSSDDMLIKL WDWDKKWSCSQVFEGHTHYVMQI VINPKDNNQFASASLDRTIKVWQLG SSSPNFTLEGHEKGVNCIDYYSGGD KPYLISGADDRLVKIWDYQNKTCVQ TLEGHAQNVSCASFHPELPIIITGSE DGTVRIWHSSTYRLESTLNYGMERV WCVASLRGSNNVALGYDEGSIIVKL GREEPAMSMDANGKIIWAKHSEVQ QANLKAMGDTEIKDGERLPLAVKD MGSCEIYPQTIQHNPNGRFVVVCGD GEYIIYTAMALRNKSFGSAQEFAWA HDSSEYAIRESNSIVKIFKNFKEKKS FKPDFGAESIYGGFLLGVRSVNGLAF YDWENTELIRRIEIQPKHIFWSDSGE LVCIATEESFFILKYLSEKVLAAQETH EGVTEDGIEDAFEVLGEIQEIVKTGL WVGDCFIYTSSVNRLNYYVGGEIVTI AHLDRTMYLLGYIPKDNRLYLGDKE LNIVSYSLLVSVLEYQTAVMRRDFS MADKVLPTIPKEQRTRVAHFLEKQG FKQQALTVSTDPEHRFELALQLGEL KIAYQLAVEAESEQKWKQLAELAISK CQFSLAQECLHHAQDYGGLLLLATA SGNASMVNKLAEGAERDGKNNVAF MSYFLQGKLDACLELLIRTGRLPEAA FLARTYLPSQVSRVVKLWRENLSKV NQKAAESLADPTEYENLFPGLKEAF VVEEWVKETHADLWPAKQYPLVTP NEERNVMEEAKGFQPSRPTAQQEP DGKPASSPVIMASQTTHKEEKSLLEL EVDLDNLELEDIDTTDINLDEDILDD	True	True	4.001	2.451	1.526	1.839	5.0	1.574	1.26