

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	extracellular region
B5DEL3	TTC17_RAT	Rattus norvegicus	Tetratricopeptide repeat protein 17	23.439218	NaN	NaN	38843836	MAAAIGVGRGFELLSCSGPGWLISLS ALLSVVARGALATTHWVVTEDEGKIQ QQVDSPMNLKHPHDLVILMRQETT VNYLKELEKQLVAQKIHIENEDRD TGLEQRHNKEDPDCIKAKVPLGDLD LYDGYITLESKDVRPEDFIDTESVVP PDPEQPDCTKILELPYSIHAFQHLLRG VQERVNLSAPLLPKEDIPTYLSKRL GRSIDDIGHLIHEGLQKNSSSWVLY NLASFYWRKNEPYQVVECTMRALH FSSRHNKDIALVNLANVLHRAHFS DAAVVHAALDDSAFFTSYTLGNIY AMLGEYNHSLVLCYDHALQAKPGFE QAIKRKHAVLCQOKLEQKLEAQHRS LQRTLNELKEYKQKQHDHYLRQOEIL EKHKLIQEEQILRNIIHETQMAKEAQ LGNHQICRLVQHQHSLHCQWDQPV RYHRGDI FENVVYVQFGDDSSSTSSM MSVNFVPTNQSDVSESVRSPVAH SVLWVWGRDS DAYRDKQHILWPKR ADCTDSYPRVPLGGELPTYFLPPENK GLRIHELTSDDYSSEEAQTPDCSIT DFRKSHTLSYLVELEVRMDLKAKIP DDHARKILLSRIKNYTIPEEEIGSFLF HAINKPNAPVWLILNEAGLYWRAVG NSTFAIACLQRALNLPVQYQDIPLV NLANLLIHYGLHLDATKLLQALAV NSSEPLTFLSLGNAYLALKNVSGALE AFRQALKLTTRCPECESSLKLIRCMQ FYPFLYNITSSVCSGHCHEKTLDNS HDKQKYFAKPSLDAEEEPSRHGA DEDPVLSVENAGRSDALRELESTVV EESNGSDEVEKSEETKMSEEILALV DEFQAWPLEGLGGTLEMKGRRLD LQGIRVLKKGPDGVAKSSCYGDCR SEDDATEWITFQVKRVKPKGDHK KPPGKKVEASQAENGQRYQANLEIT GPKVASPGPQEKRDYQSMGWPS DECLKLRWVELTAIVSTWLAVSSKNI DITEHIDFATPIQQPAMEPLCNGNLP TSMHTLDHLHGVSNRASLHYTGES QLTEVLQNLGKDOYPOOSLEQIGTRI AKVLEKNQTSWVLSMAALYWRVK GQGKAIDCLRQALHYAPHQMKDV PLISLANILHNAKLWNDVAVVATMA VEIAPHFAVNHFTLGNVYVAMEEFE KALVWYESTLKLQPEFVPAKNRIQTI QCHMLLKKGRSP	False	False	2.74	0.281	0.422	0.375	False	4.188	0.375