

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golg. appa
Q80TZ3	AUXI_MOUSE	Mus musculus	Auxilin	52.100753	T588;T590;S591;T605;S623;S628	S137;S478;S481;S595	33300544;22517741;22645316;40885482;35822049;40021952	MTNPKSGVAESAGLACSRRAAGEN RMKDSEKNGASSPDMEPSYGGGLF DMVKGGAGRLFNSLNKDNLKDITLKD TSSRVIQSVSSYTKGDLDFYVTSRII VMSFPVDSVDIGFRNQVDDIRSFLLD SRRLDHYTVYNLSPKSYRTAKFHRSR VSECSWPIRQAPSLHNLFAVCRNMY NWLLOPKNVVHCLDGRAASSIL VGAMFIFCNLYSTPGPAVRLLYAKRP GIGLSPSHRRYLGYMCDLLADKPYR PHFKPLTIKAITVSPVPPFNKQRNGC RPYCDVLIGETKIYSTCTDFERMKEY RVQDGKIFIPLNITVQGDVIVSMYHL RSTIGSRLQAKVTNTQIFQLQFHSGF IPLDITVLKFTKPELDACDVPEKYPQ LFQVTLDIEVQPQDKVIDLTPPWEHY CTKDVNPSILFSSQHEQDITLALGG QAPADLPPDHPRNLGQGGFFASLC WQDQKSEKSRCEEDHAALVNQESE QSDELLTLSSPHGNAEGDKPHGAK KPGKKQPEAAPPPEVDLLGLEG SDVSTNFSSLAAPPSNELLSDLFGG VGATGPAQAGQAGVEDVFHPSGPVS AQSTPRRTATSASASPTLRVGEATF DPFGAPAKPPGQDLLGSFLNTSSAS SDPFLQPTRSPSPVHASSTPAVNIQ PDIAGGWDWHTKPGGFGMGSKSAA TSPTGSSHGTPHQSKPQTLDFAD LGTGSSSFASKPTTPTGLGGFPPL SSPQKASPQPMGGGWQFPAGYNW QQTQSKPQSSMPHSSPQNRPNYNV SFSAMPAGQSERGKGSNTLEGKQK AADFEDLLSSQGFNAHKDKKGPRTI AEMRKEEMAKEMDPEKLEWIE GKERNIRALLSTMHTVLWAGETKW KPVGMADLVTPEQVKKVYRRAVLVV HPDKATGQPYEQYAKMIFMELNDA WSEFENQGGKPLY	False	False	2.626	2.726	1.735	1.551	1.911