

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane
Q6NZC7	S23IP_MOUSE	Mus musculus	SEC23-interacting protein	45.142925	S600;S607	S600;S735;S748;S924	22645316;33300544;40885482;35822049;39627609	MADRKANGGGASASSGTNLLFSSS ATEFSFNVPFIPVTQAAASPASLLLP GEDSTDVGEEDSFLGQTSHTSTPQ TFSYFSQVSSSDPFGNIGQSPLTTS AMSAGQSALPKPPATLAFITGSQDA LSAFPPSVSKAPGATPPSQMGSTYS PSQPSLPPNFGSPQGIQOGHNP YRHTPVSSRANPYITPPQLQCCOMP GHSSYPPSPGPVQTYQMPPLPLPL PTAMQSPAQQQVPARPAGPLVQGPS PFVLQNYEPVQPHWFYCKEVEYKQ LWMPFVLDLNLLEEIYNSVQDPDE SVVLGTDGGRYDVLYDRMRKSVYW EEEPTEVRRCTWFYKGDTSRFIPYT EEFSEKLEAEYKAVSTNQWHRRL FPSGETIVMHNPKVIVQFQPSSVPDE WGTTQDGGQTRPRVVKRGIDSLDEI PDGEMPVQVDHLVFMVHGIGPVCDL RFRSIECVDDFRVSLKLLQTHFKK SVDEGKVSVEFLPVHWSALGGH ATGVDRNIKITLPSIGFRFRHFTNET LLDVLFYNSPTYCQAIKVEVEINR LHSLFMSRNPNFKGKVSVAGHSLG SLILFDILSNQRDMCVSKSPGSAVVS NGVIKHSQFQEKLISEESKLSMDES CDLDVEDEEPLTLHGTLLEALSFDYI STFEKEKIDMESLLMCTVDDLKEMG IPLGPRKKIANFVKLKAALKLEQKAA AEKKAALAALTKGQDESAPKTKEMA SPSSESNEKRKLSVGAYVSSVRVDY ESFEVGTGQVSVAYSSLDPEPEIFFA LGSPIGMLLTIRGVARIDEKYRLPTC KGFNHYHPLDPVAYRLEPMIAPDL LKAVLVPHHKGRKRLHLELKESLSR MGSDLKQGFISLKSAAWQTLNEFAR AHTSTTQLQEELEKVANQIKKEEEK QVVEAKKKKESPELSKDEDYLGKVG MLNGRRIDYVLQEKPIESFNEYLF LQSHLCYWGS EDTALLLKEIYRTM NISPEQPQH	False	True	2.194	2.831	1.116	5.0	5.0	1.045