

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	extrac region
Q6NZC7	S231P_MOUSE	Mus musculus	SEC23-interacting protein	35.309632	NaN	S600;S735;S748;S924	33300544;35822049;22645316	MADRKANGGGASASSSGTNLLFSSS ATEFSFNVPFIPVTQAAASPALLLP GEDSTDVGEEDSFLGQTSHTSTPQ TFSYFSQVSSSDPPGNIGQSPLTTS AMSAGQSALPKPPATLAFITGSQDA LSAFPSPVSKAPGATPPSQMGTSTYS PSQPSSLPPNFGSPPOGIPQQGHNP YRHTPVSSRANPYITPPQLQQQMP GHSSYPSPGPPVQTYQMPGPLPPL PTAMQSPAQQVVPARPAGPLVQGPS PFVLQNYEPVQPHWFYCKEVEYKQ LWMPFSVLDLNLLEIYNSVQDPDE SVVLGTDGGRYDVLYDRMRKSVYW EEEPTEVRRCTWFKGDTDSRFIPYT EEFSEKLEAEYKKAVSTNOWHRRLE FPSGETIVMHNPKVIVQFQSSVPDE WGTTQDGGQTRPVVVKRGIDDSLDEI PDGEMPQVDHLVFMVHGIGPVCDEL RFRSIECVDDFRVVSLLKLLQTHFKK SVDEGKVS RVEFLPVHWH SALGGH ATGVDNRNKKITLPSIGRFRHFTNET LLDVL FYN SPTYCQAIVEKVEVEINR LHSLFMSRNP NFKGKVS VAGHSLG SLILFDILSNQRDMCVSKSPGSAVS NGVIKHSQFQEKLI SEESKLMSDES CDLDVEDEEPLTLHGTLAALSFDYI STFEKEKIDMESLLMCTVDDLKEMG IPLGPRKKIANFVKLKA AKLEQKAAA AEKKAALALTKGQDESAPKTKEMA SPSSESNE SKRKL SVGAYVSSVRVDY ESFEVGTGQVSVAYSSLD FEPEIFFA LGSPIGMLLTIRGVARIDEKYRLPTC KGFNFIYHPLDPVAYRLEPMIAPDL LKAVLVPHHKGRKRLHLELKE SLSR MGSDLKQGFISLKS AWQTLNEFAR AHTSTTQLQE ELEKVANQIKEEE EK QVVEAKKKKESPELSKDEDYLGKVG MLNGGRRIDYVLQEKPIESFNEYLFA LQSHLCYWGSEDTALLLKEIYRTM NISPEQPQH	None	None	None	None	None	None	None		

0