

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membr
Q7TSI3	PP6R1_MOUSE	Mus musculus	Serine/threonine-protein phosphatase 6 regulatory subunit 1	28.201711	NaN	S232;T524;S529;S530;S531;S633;S636;S698;S739;S826	39627609;33300544	MFWKFDLHTSSHLDTLLEKEDLSLP ELLDEEDVLQECKVVNRKLLDFLLQ PSHLQAMVAWVTQEPASGEERLRY KYPVACEILTSQVINDALGADES LLNRLYGFLQSGDSLNPLLASFFSKV MGILINRKTQDLVSLRKKDDFDL LLRHIGTSAIMDLLLRLLTCVERPQL RODFVFNWLNEEKIVQRLIEQIHPK DDNQHSNASQSLCDIIRLSREQMIQ GDSPPEPDQLLATLEKQETIEQLLSN MFEGEQCQSVIVSGIQVLLTLEPRR PRSDSVTMNNFFSSVDGQLELLAQQ ALDDALSSMGALHALRPRDRFHQL LLEPPKLEPLQMTWGS LAPLGNT LHVVKLLASALSANAAALTQELLVLD VPNTLLDLFFHYVFNNFLHAQVEVC VSAMLSGGPPDSSSETPVNPVIVKH LLQHCRVLVERILASWEENDRVQSGG GPRKGYMGHLTRVANAVVQNAEQG PNAEQLGQLLKELPEEQQRWEAF VSGPLAETNKKNTVDLVNTHLHSS SDEDDRLEKFNPEEAVLQOAFM DFQMQRMTSAFIDHFGFNDEEFGE QEEVSNAPFDKTANITFSLNADDE PNANLLEICYKDRIQQFDDEEEEEE EGQGSAESDGEYGAWQGSQPVRAS QASQPPGVRS GGSTDEEEDEEED EEDEGAEQAACGRTPSSFPSPSTQ PPGPSWTATFDTPMDAPTGPVSK EADMSSIQIPSSPPAHGSPQLRSQDP THPSAPQEVTDSSKVAEPLAPCQALV SVADVQATLHGMRSAPSSLDSATRD PSTSVPDFKAHQSPQTMEGKRSP LGLPQSQSALEMPNGSTPGGPISSG SQ	False	False	3.429	3.144	0.523	False	0.629	0.894