

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
Q9UPN7	PP6R1_HUMAN	Homo sapiens	Serine/threonine-protein phosphatase 6 regulatory subunit 1	27.216596	S800	S232;T524;S529;S530;S531;S635;S638;S702;S726;S759;S846	35132862;23301498	MFWKFIDLHTSSHLDTLEREDLSLP ELLDEEDVLQECKVNRKLLDFLLQ PPHLQAMVAWVTQEPDSDGEERLR YKYPVACEILTSQVQINDALGADE SLLNRLYGFLQSTGSLNPLASFFSK VMGILINRKTQDLVSLRKKDDFVD LLLQHGTSAIMDLLRLLTCVERPQ LRQDVVNWLNEEKIVORLIEQIHPS KDENQHSNASQSLCDIIRLSREQMI QVQDSEPDQLLATLEKQETIEQLLS NMFEGEQSQSVIVSGIQVLLTLEPR RPRSESVTVNSFFSSVDGQLELLAQ GALESTVSSVGALHALRPRLSCFHQ LLEPPKLEPLQMTWGMLAPPLGNT RLHVVKLLASALSANDAALTHELLA LDVPNTMLDLFFHYVFNNFLHAQV EGCVSTMLSLGPPDSSPETPIQNPV VKHLLQQCRLVERILTSWEENDRVQ CAGGPRKGYMGLHTRVAGALVQNT EKGPNAEQLRQLLKELPSEQEQEW EAFVSGPLAETNKKNMVDLVNTHH LHSSDDEDDLKFEFNFPEEAVLQ AFMDFQMQRMTSAFIDHFGFNDEE FGEQEEVSNAPFDKTANITFSLNAD DENPNANLLEICYKDRIQQFDDDEE EEDEEEAQSAGESDGEDGAWQGSQ LARGARLGPPGVRSGGSTDSEDEE EEDEEEDEEGIGCAARGGATPLS YSPGPQPPGWSWTATFDVPTDAPT SPRVSGEEELHTGPPAPQGPLSVQ GLPTQSLASPPARDALQRSQDPTPP SAPQEATEGSKVTEPSAPCQALVSI DLQATFHGIRSAPESSDSATRDPSTS VPASGAHQPPQTTEGEKSPEPLGLP QSQAQALTPPIPNGSAPEGPASPG SQ	True	False	4.746	4.288	1.053	0.687	0.802	1.186