	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparat
0	Q5H9R7	PP6R3_HUMAN	Homo sapiens	Serine/threonine-protein phosphatase 6 regulatory subunit 3	15.308014	NaN	S315;S579;S617;T631;S634;S722;S853	30379171;23301498;28510447	MFWKFDLHSSSHIDTLLEREDVTLK ELMDEEDVLQECKAQNRKLIEFLLK AECLEDLVSFIIEEPPQDMDEKIRYK YPNISCELLTSDVSQMNDRLGEDES LLMKLYSFLLNDSPLNPLLASFFSKV LSILISRKPEQIVDFLKKKHDFVDLII KHIGTSAIMDLLLRLLTCIEPPQPRQ DVLNWLNEEKIIQRLVEIVHPSQEED RHSNASQSLCEIVRLSRDQMLQIQN STEPDPLLATLEKQEIIEQLLSNIFHK EKNESAIVSAIQILLTLLETRRPTFEG HIEICPPGMSHSACSVNKSVLEAIRG RLGSFHELLLEPPKKSVMKTTWGVL DPPVGNTRLNVIRLISSLLQTNTSSI NGDLMELNSIGVILNMFFKYTWNN FLHTQVEICIALILASPFENTENATIT DQDSTGDNLLLKHLFQKCQLIERILE AWEMNEKKQAEGGRRHGYMGHLT RIANCIVHSTDKGPNSALVQQLIKDL PDEVRERWETFCTSSLGETNKRNTV DLVTTCHIHSSSDDEIDFKETGFSQD SSLQQAFSDYQMQQMTSNFIDQFG FNDEKFADQDDIGNVSFDRVSDINF TLNTNESGNIALFEACCKERIQQFD DGGSDEEDIWEEKHIAFTPESQRRS SSGSTDSEESTDSEEEDGAKQDLFE PSSANTEDKMEVDLSEPPNWSANF DVPMETTHGAPLDSVGSDVWSTEEP MPTKETGWASFSEFTSSLSTKDSLR SNSPVEMETSTEPMDPLTPSAAALA VQPEAAGSVAMEASSDGEEDAESTD KVTETVMNGGMKETLSLTVDAKTET AVFKSEEGKLSTSQDAACKDAEECP ETAEAKCAAPRPPSSSPEQRTGQPSA PGDTSVNGPV	None	None	None	None	None	None	None