	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatu
0	Q922D4	PP6R3_MOUSE	Mus musculus	Serine/threonine-protein phosphatase 6 regulatory subunit 3	29.536434	NaN	S315;S550;S588;T602;S605;S693;T820;S824;S525	28135057;34887587;28528544	MFWKFDLHSSSHIDTLLEREDVTLK ELMDEEDVLQECKAQNRKLIEFLLK AECLEDLVSFIIEEPPQDMDEKIRYK YPNISCELLTSDVSQMNDRLGEDES LLMKLYSFLLNESPLNPLLASFFSKV LSILISRKPEQIVDFLKKKRDFVDLIIK HIGTSAIMDLLLRLLTCIEPPQPRQD VLNWLNEERIIQRLVEIVHPSQEEDR HSNASQSLCEIVRLSRDQMLQVQNS TEPDPLLATLEKQEIIEQLLSNIFHKE KNESAIVSAIQILLTLLETRRPTFEGH IEICPPGMSHSACSVNKSVLEAIRGR LGSFHELLLEPPKKSVMKTTWGILD PPVGNTRLNVIRLISSLLQTNTSSIN GDLMELNSIGVILDMFFKYTWNNFL HTQVEICIALILASPFENAENGTITDQ DSTGDNLLLKHLFQKCQLIERILEA WDTNEKKQAEGGRRHGYMGHLTRI ANCIVHSTDKGPNSALVQQLIKDLP DEVRERWETFCTNSLGETNKRNTV DLAFSDYQMQQMTSNFIDQFGFND EKFADQDDIGNVSFDRVSDINFTLN TNESGNIALFEACCKERIQQFDDGG SDEEDIWEEKHIAFTPESQRRSSSGS TDSEESTDSEEEDGAKQDLFESSSA NTEDKMEVDLNEPPTWSANFDVPM ETTHGAPLDSVGSDVWSTEEPMPTK ETGWASFSEFTSSLSTKESLRSNSPV EMETSTESVDPLTPGAAALATQPEAP GSMAMEASSDGEEDAESTDKVTET VMNGGMKETLSLTVDAKTETAVFKS EEEKLSTSQDAACKDAEETPEPAEA KCTAPLTPSSSPEQRTDQPSMPSDP SVNGPV	None	None	None	None	None	None	None