	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrac region
0	Q6P2K6	P4R3A_MOUSE	Mus musculus	Serine/threonine-protein phosphatase 4 regulatory subunit 3A	25.019509	T241	S117;S127;S685;S728;S755;S758;S761;S764;S767	34418053	MTDTRRRVKVYTLNEDRQWDDRGT GHVSSGYVERLKGMSLLVRAESDGS LLLESKINPNTAYQKQQDTLIVWSEA ENYDLALSFQEKAGCDEIWEKICQV QGKDPSVDITQDLVDESEEERFDDM SSPGLELPSCELSRLEEIAELVASSLP SPLRREKLALALENEGYIKKLLELFH VCEDLENIEGLHHLYEIIKGIFLLNRT ALFEVMFSEECIMDVIGCLEYDPTLS QPRKHREFLTKTAKFKEVIPISDPEL KQKIHQTYRVQYIQDMVLPTPSVFEE NMLSTLHSFIFFNKVEIVGMLQEDE KFLTDLFAQLTDEATDEEKRQELVN FLKEFCAFSQTLQPQNRDAFFKTLS NMGILPALEVILGMDDTQVRSAATDI FSYLVEYNPSMVREFVMQEAQQND DDILLINLIIEHMICDTDPELGGAVQL MGLLRTLVDPENMLATANKTEKTEF LGFFYKHCMHVLTAPLLANTTEDKP SKDDFQTAQLLALVLELLTFCVEHH TYHIKNYIINKDILRRVLVLMASKHA FLALCALRFKRKIIGLKDEFYNRYIM KSFLFEPVVKAFLNNGSRYNLMNSA IIEMFEFIRVEDIKSLTAHVIENYWKA LEDVDYVQTFKGLKLRFEQQRERQD NPKLDSMRSILRNHRYRRDARTLED EEEMWFNTDEDDMEDGEAVVSPSD KTKNDDDIMDPISKFMERKKLKESE EKEVLLKTNLSGRQSPSFKLSLSSGT KTNLTSQSSATSLPGSPGSPGSPGSP GSPGSVPKSTSQTAAITTKGGLVGLV DYPDDDEDDDEDEDKEDTLPVSKKA KFES	None	None	None	None	None	None	None	None	None