

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane
Q9WUB3	PYGM_MOUSE	Mus musculus	Glycogen phosphorylase, muscle form	26.740217	NaN	S15;T204;T227;S430;T473;S514;S747;S748	33300544;36288343	MSRPLSDQDKRKQISVRGLAGVENV SELKKNFNRLHLFTLVKDRNVATPR DYYFALAHTVRDHLVGRWIRTQQHY YEKDPKRIYYLSLEFYMGRTLQNTM VNLALENACDEATYQLGLDMEELEE IEEDAGLGNGLGRLAACFLDSMAT LGLAAYGYGIRYFYGIFNQKICGGWQ MEEADDWLRYPGNPWEKARPEFTLP VHFYGRVEHTSQGAKWVDTQVLA MPYDTPVPGYRNNVVNTMRLWSAK APNDFNLKDFNVGGYIQAVALDRNLA ENISRVLYPNDFNFEKELRLKQEY FVVAATLQDIIRRFKSSKFGSRDPVR TNFDAFPDKVAIQLNDTHPSLAPEL MRILVDLERLDWDKAWDVTVKTCA YTNHTVLPALERWPVHLMETLLPR HLQIYEINQRFLNRVAAAFPGDVDR LRRMSLVEEGAVKRINMAHLCIAGS HAVNGVARIHSEILKKTIFKDFYELE PHKFQNKTNGITPRRWLVLCNPGLA EVIAERIGEDYISDLQLRKLLSYVD DEAFIRDVAKVKQENKLFSAYLEY EYKVHINPNSLFDVQVKRIHEYKRO LLNCLHITLYNRIKREPFRFVMPRT IMIGKAAPGYHMAKMIKLIAGD VVNHDPAVGDRLRVIFLENYRVSLA EKVIPAADLSEQISTAGTEASGTGNM KFMLNGALTIGTMDGANVEMAEAA GEENFFIFGMRVEDVERLDQRGYNA QEYYDRIPELRQIEQLSSGFFSPKQP DLFKDIVNMLMHHDRFKVFADYEE YKCKQKVSSELYKNPREWTRMVIRN IATSGKFSDDRTIAQYAREIWGVEPS RQLRPAPDEKI	False	True	3.147	2.751	3.098	4.427	1.244	2.102