

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	p
P52480	KPYM_MOUSE	Mus musculus	Pyruvate kinase PKM	44.724957	NaN	S37;T41;S97;S100;T105;S127;T148;T175;T195	33300544;22645316;34678516;28528544;34887587;37453647;24788674;21540332;36288343;39627609;36064721;37507081	MPKPHSEAGTAFIQQLHAAMADT FLEHMCRLDIDSAPITARNTGICTIG PASRSVEMLKEMIKSGMNVARLNF SHGTHEYHAETIKNVREATESFASD PILYRPVAVALDTKGPEIRTGLIKGSG TAEVELKKGATLKITLDNAYMEKCD ENILWLDYKNICKVVEVGSKIYVDDG LISLQVKEKGADFLVTEVENGGSLGS KKGVNLPGAAVDLPVSEKDIQDLK FGVEQDVMVFASFIRKAADVHEVR KVLGEKGKNIKIISKIENHEGVRFFD EILEASDGIMVARGDLGIEIPAEKVFL AQKMMIGRCNRAGKPVICATQMLE SMIKKPRPTRAEGSDVANAVLDGAD CIMLSGETAKGDYPLEAVRMQHLLIA REAEAAIYHLQLFEELRRRAPITSDPT EAAAVGAVEASFCCSGAIIVLTKSG RSAHQVARYRPRAPIIAVTRNPQTAR QAHLYRGIFPVLCDAVLNWAEDV DLRVNLAMDVVGKARGFFKKGDDVIV LTGWRPESGFTNTMRVVPV	True	True	5.0	4.62	4.668	5.0	1.784	2