

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	extra regic
P47857	PFKAM_MOUSE	Mus musculus	ATP-dependent 6-phosphofructokinase, muscle type	24.048659	NaN	S133;S377;S667;S775	33300544;37507081	MTHEEHHAAKTLGIGKAIAVLTSGG DAQGMNAAVRAVVRVGIPTGARVFF VHEGYQGLVDGGEHIREATWESVS MMLQLGGTVIGSARCKDFREREGRL RAAHNLVKRGITNLICVIGDGLTGT ADTFRSEWSDLLNDLQKDKGKITAEE ATKSSYLNIVGLVGSIDNDFCGTDM TIGTDSALHRIVEIVDAITTTAQSHQR TFVLEVMGRHCGYLALVTSLSGAD WVFIPECPDDDDWEEHLCLRLSETR TRGSRLNIIIVAEGAIDKNGKPTSED IKNLVVKRLGYDTRVTVLGHVQGGG TPSAFDRILGSRMGVEAVMALLEGT PDTACVVVLSGNQAVRLPLMECVQ VTKDVTKAMDEKRFDEAIKLRGRSF MNNWEVYKLLAHVRRPPVSKGGLHT VAVMNVGAPAAGMNAAVRSTVIRGL IQGNRVLVVDHGFGLAKGQIEEAG WSYVGGWTGQGGSKLGTKRTLPKK NLEQISANITKFNIQGLVIHGGFEAYT GGLELMGRKQFDELCPFVVIPATV SNNVPGSDFSIGADTALNICTTCDR IKQSAAGTKRRVFIHETMGGYCGYLA TMAGLAAGADAAYIFEEPFTIRDLQV NVEHLVQKMKTTVKRGLVLRNEKC NENYTTDFIFNLYSEEGKGFIDSRKN VLGHMQQGSPTPFDRNFATKMGA KAMNWMMSGKIKESYRNGRIFANTP DSGCVLGMRKRALVFQVTELDQDT DFEHRIPKEQWWLKLRLPILKILAKYE IDLDTSDAHLEHISRKRSGEAAV	True	False	5.0	3.549	2.317	1.55	0.968	3.124	2.19