

UniprotKB ID	Entry name	organism	full name	oglcnaescore	oglcnaesites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	g	a
Q01813	PFKAP_HUMAN	Homo sapiens	ATP-dependent 6-phosphofructokinase, platelet type	24.772636	T538	S6;S12;S21;S142;S386;T651;S783	38665916;33214551;22923583;34019948;26853435;34725712;37217939;29351928	MDADDSRAPKGSRLRKFLEHLSGAG KAICVLTSGGDAQGMNAVRAVVR MGIYVGAKVYFIYEGYQGMVDGGSN IAEADWESVSSILQVGGTIGSARCQ AFRTREGRLKAACNLLQRGITNLCVI GGDGSALTGANLFRKEWSGLLEELAR NGQIDKEAVQKYAYLNVVGMVGSID NDFCGTDMTIGTDSALHRIEIVDAI MTTAQSHQRTFVLEV MGRHCGYLA LVSALACGADWVFLPESPPEEGWEE QMCVKLSENRAKRLNIIIVAEgai DTQNKPISEKIKELVVTQLGYDTRV TILGHVQRGGTPSAFDRILASRMGV EAVIALLEATPDTACVVSLNGNHAV RLPLMECVQMTQDVQKAMDERRFO DAVRLRGRSFAGNLNTYKRLAIKLP DDQIPKTNVAVINVGAPAAAGMNA AVRSAVRVGIADGHRMLAIYDGFDFG FAKGQIKEIGWTDVGGWTQGGGIL GTRVLPKYLEELATQMRTHSINAL LIIGGFEAYLGLLELSAAREKHEEFC VPMVMVPATVSNVPGSDFSIGADT ALNTITDTCDRIKQSASGTRRVFIE TMGGYCGLANMGGLAAGADAAYIF EPPFDIRDLQSNVEHLTEKMKTTIQ RGLVLRNESCSENYITDFIYQLYSEE GKGVFDCRKNVLGHMQGGAPSPF DRNFGTKISARAMEWITAKLKEARG RGKKFTDSSICVLGISKRNVIFQPV AELKKQTD FEHRIPKEQWWLKLRLPL MKILAKYKASYDVS DSGQLEHVQPW SV	True	True	4.867	4.571	2.473	1.591	1	