

UniprotKB ID	Entry name	organism	full name	oglcnaescore	oglcnaesites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P30835	PFKAL_RAT	Rattus norvegicus	ATP-dependent 6-phosphofructokinase, liver type	23.439218	NaN	S377;T640;S775	38843836	MATVDLEKLRMSGAGKAIGVLTSGG DAQGMNAAVRAVTRMGIYVGAKVF LIYEGYEGLVEGGENIKPANWLSVS NIIQLGGTIIGSARCKAFTTREGRLAA AYNLLQHGITNLCVIGGDGSLTGANI FRNEWGSLLEELVKEGKISESTAQN YAHLSIAGLVGSIDNDFCGTDMTIGT DSALHRIMEVIDAITTTAQSHQRTFV LEVMGRRHCGYLALVSALASGADWLF IPEAPPEDGWENFMCERLGETRSRG SRLNIIIIAEGADRHGKPISSSYVKDL VVQRLGFDTRVTVLGHVQRGGTPSA FDRVLSKMGMEAVMALLEATPDT PACVVSLSGNQSVRLPLMECVQVK DVQKAMDEKRFDEAIQLRGRSFEN NWKIYKLLAHQKVSKEKSNFSLAIL NVGAPAAGMNAAVRSVRTGISEGH TVYVVHDGFEGLAGKQVQEVGWHHD VAGWLGRRGSM LGTKR TL PKPHLE AIVENLRTYNIHALLVIGGFAYEGV LQLVEARGRYEELCIVMCPATISN NVPGTDFSLGSDTAVNAAMESCDRI KQSASGTRRRVFIVETMGGYCGYLA TVTGI AVGADAAYVFEDPFNIHDLKA NVEHMT EKMTDIQRGLVLRNEKC HEHYTTEFLYNLYSSEGRGVFDCRT NVLGHLQGGAPTFDRNYGTKLGV KAMLWMSEKLRDVRKGRVFANAP DSACVIGLRKKVAFSSVTELKKETD FEHRMPREQWWNLRLMLKMLAH YRISMADYVSGELEHVTRR TL SIDKG F	True	False	4.742	2.667	2.142	1.268	False	1.564	1.699