

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	extracellular region
Q7TSH2	KPBB_MOUSE	Mus musculus	Phosphorylase b kinase regulatory subunit beta	30.924281	NaN	S10;S19;S693	39927985;33300544	MANSPDAAFSSPALLRSGSVYEPLK SINLPRPDNETLWDKLDHYYRIVKST MLMYQSPTTGLFPKTGCGEEKSKV HESLYCAAGAWALALAYRRIDDDKG RTHELEHSAIKCMRGILYCYMRQAD KVQQFKQDPRPTTCLHSVFSVHTGD ELLSYEEYGHLLQINAVSLFLLYLVEV ISSGLQIYNTDEVSFIONLVFCVERV YRVPDFGVWERGSKYNNGSTELHS SSVGLAKAALEAINGFNLFNGQGC WSVIFVDLDAHNRNRQTLCSLLPRE SRSHNTDAALLPCISYPAFALDDEAL FSQTLDKVIRKLGKYGFKRFLRDGY RTPLEDPNRRYYKPAEIKLFDGIECE FPIFFLYMMIDGVFRNGLEQVKEYQ DLLTPLLHQTTTEGYVVPKYYVVPAD FVECEKRNPQSGKRFPSNCGRDGKL FLWGOALYIIAKLLADELISPKDIDPV QRFVPLQNQRNVSMRYSNQGPLEN DLVVHVALVAESQRLQVFLNTYGIQ TQTPQQVEPIQIWPQQELVKAYPHL GINEKLGSLGRPDRPIGCLGTSKIYRI LGKTVVCPYPIIFDLSDFYMSQDVLLLI DDIKNALQFIKQYWKMHGRPLFLVL IREDNIRGSRFNPILDMLAAFKKGIIG GVKVHVDRLQTLISGAVVEQLDFLRI SDTEKLPEFKSFEELEFPKHSKVKR QSSADAPEAQHEPGITITIEWKNKST HEILQKLNDCGCLAGQTILLGILLKR EGPNFITMEGTVSDHIERYVRRAGS KKLWSVVRRAASLLNKVVDSLAPSIT NVLVQKQVTLGAFGHEEEVISNPL SPRVIKNIIYKCNTHDEREAVIQQE LVHIGWIISNSPELFSGMLKIRIGWII HAMEYELQVRGGDKPAVDLYQLSPS EVKQLLLDILQPOQSGRCWLNRRQI DGS LNRTPEFYDRVWQILERTPNGI VVAGKHLPOQPTLSDMTMYEMNFS LLVEDMLGNIDQPKYRQIIVELLMVV SIVLERNPELEFQDKVDLDRLVKEAF HEFQKDESRLKEIEKQDDMTSFYNT PPLGKRGTCSYLTKVMNSLLEGEV KPSNEDSCLVS	False	False	2.016	1.219	0.234	0.234	False	3.112	False