

UniprotKB ID	Entry name	organism	full name	oglcnaicscore	oglcnaicsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q62074	KPCI_MOUSE	Mus musculus	Protein kinase C iota type	27.03061	NaN	T3;S7;S8;T9;T264;T279;T333;T411;T563	25153642	MPTQRDSSTMSHTVACGGGGDHS QVRVKAYYRGDIMITHFEPSSIFEGL CSEVRDMCSFDNEQPFTMKWIDEE GDPCTVSSQLEEEAFRLYELNKDS ELLIHVFPVPERPGMPCPGEDKSIY RRGARRWRKLYCANGHTFQAKRFN RRAHCAICTDRIWGLGRQGYKCINC KLLVHKKCHKLVITIECGRHSLPPEP MMPMDQTMHPDHTQTVIPYNPSSH ESLDQVGEEKEAMNTRESGKASSSL GLQDFDLRVIGRGSYAKVLLVRLKK TDRIYAMKVVKELVNDDEDIDWVQ TEKHVFEQASNHPFLVGLHSCFQTE SRLFFVIEYVNGGDLMFHMQRQRK LPEEHARFYSAEISLALNYLHERGIY RDLKLDNVLLDSEGHILTDYGMCK EGLRPGDITSTFCGTPNYIAPEILRG EDYGFSDWWALGVLMFEMMAGR SPFDIVGSSDNDQNTEDYLFQVILE KQIRIPRSLSVKAASVLKSFLNKDPK ERLGCHPQTGFADIQGHPPFRNVD WDMMEQKQVPPFKPNISGEFGLD NFDSQFTNEPVQLTPDDDDIVRKID QSEFEGFEYINPLLSAEECV	True	True	4.462	5.0	1.745	1.494	2.86	5.0	3.206

0