	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	P49657	MNB_DROME	Drosophila melanogaster	Serine/threonine-protein kinase minibrain	25.496522	NaN	NaN	33925313	MYRLEDTNSGGVMDKNKQKLSAYG SSGGSVDAAQGSGSGGGRQRHAPLY GRFVDAEDLPATHRDVMHHHSSPS SSSEVRAMQARIPNHFREPASGPLR KLSVDLIKTYKHINEVYYAKKKRRAQ QTQGDDDSSNKKERKLYNDGYDDD NHDYIIKNGEKFLDRYEIDSLIGKGS FGQVVKAYDHEEQCHVAIKIIKNKKP FLNQAQIEVKLLEMMNRADAENKY YIVKLKRHFMWRNHLCLVFELLSYN LYDLLRNTNFRGVSLNLTRKFAQQL CTALLFLSTPELNIIHCDLKPENILLC NPKRSAIKIVDFGSSCQLGQRIYHYI QSRFYRSPEVLLGIQYDLAIDMWSL GCILVEMHTGEPLFSGCNEVDQMN KIVEVLGMPPKYLLDQAHKTRKFFD KIVADGSYVLKKNQNGRKYKPPGSR KLHDILGVETGGPGGRRLDEPGHSV SDYLKFKDLILRMLDFDPKTRVTPYY ALQHNFFKRTADEATNTSGAGATAN AGAGGSGSSGAGGSSGGGVGGGLG ASNSSSGAVSSSSAAAPTAATAAATA AGSSGSGSSVGGGSSAAQQQQAMP LPLPLPLPLPPLAGPGGASDGQCHGL LMHSVAANAMNNFSALSLQSNAHP PPSLANSHHSTNSLGSLNHISPGST GCHNNNSNSSNNNTRHSRLYGSN MVNMVGHHNSGSSNNHNSISYPH AMECDPPQMPPPPPNGHGRMRVPA IMQLQPNSYAPNSVPYYGNMSSSSV AAAAAAAAAAASHLMMTDSSVISAS AAGGGQGGGNPGQNPVTPSAAAFLF PSQPAGTLYGTALGSLSDLPLPMPLP MSVPLQLPPSSSSSVSSGSASVGSG GVGVGVVGQRRHITGPAAQVGISQS VGSGSSGSATGASSSDASSSSPMVG VCVQQNPVVIH	None	None	None	None	None	None	None	None	None