	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	Q9VTU0	KRI1_DROME	Drosophila melanogaster	Protein KRI1 homolog	28.06426	T119	S95;S97;S98;S137;S138;S179	40245499	MNNNLFEGSDDDGEDLQLSTNKDY AKTYNILRKKELLQKYKDRGLDVSES EFDSDSSSSEEDEVDPKFDQDFFKT LSSLKSKDPCIYDKGTKFFSESSGDE DDKDGEAPKKKKKAKPVTLKDYERK VILEHNGKFESSDEEQQEKEHEELQ RAQSPSAVEEERRLKAEFRKVMNKE DDSEDEEFGGIFKKRSKTKEQTAAE EADFAKWLAGKQAEIQETDKKQLEP LKQYWSSNKLTQGESFLRDYILNKG YANTDESAIPTYDEIVGEAAPLSEDE QELEKQAEFEHKYNFRFEEPDADFI KRYPRTIEQSLRRTDDKRKEKRKELK ERKDQEKQQKMKELELVKEMKRKE IDEKIRKLKAVTGNDELGFRDEELEE DFDPAAHDRRMQELFDDEYYNVDE GEEKPECPSDIDELVLEDWDNYDPR QHANGGGEDYEGHCEDDDFNMDC DYDPSTAKEQLQQELIENTRGRKGR KGRRNRFMEMIQAEKPAFNPEDEK TYSEYIDEYYQMDCEDIIGDQPCRFK YVETTPNDFGLTIEEILLAKNKELNQ WASLKKAVQNRPEHVEKKEQRLYK MKAKNEDLKRKIFKSLYGEGSDDEE QPAEEKPEVTPAEATAPAENGQVST EGLSKSKRKRLKRKAAAAAASAPKV LKEESDSKDPKEADGSTEDVQAESS KKKVDTPSKKGKDDANQETKNSPQS TEKTKNNNALKNNKKEPKNVQNGF QKPQNQANKSAKTKSNQPFKTTESA PAKAEKSNGNNPFNKPQSKSQQRQ ELPPIHKNQGGNKKGPRNANGTNP FKKSNQKPSAPFPAKKTNNFKAKNK QNNNSGITDDRLKAYGINPRKFHKR EKYGKKDN	True	False	1.594	4.233	False	False	False	0.613	False