

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
Q921K2	Q921K2_MOUSE	Mus musculus	Poly [ADP-ribose] polymerase	28.854855	NaN	NaN	34887587;28528544	MAEASERLYRVEYAKSGRASCCKCS ESIPKDSLRLMAIMVQSPMFDGKVP WYHFSCFWKVGHSIRQPDVEVDGF SELRWDDQQKVKKTAEGGVAGKG QDGGGKAEKTLGDFLAEYAKSNRS MCKGCKLEKIEKGMRLSKKMVDPE KPQLGMIDRWYHPTCFVKKRDELGF RPEYSASQLKGFSLLSAEDKEALKK QLPAIKNEGKRRKGDEVDTDEVAKK KSKKGGKDKDSSKLEKALKAQNELIW NIKDELKKACSTNDLKELLIFNQQQ VPSGESAILDRVADGMAFGALLPCK ECSGQLVFKSDAYYCTGDVTAWTKC MVKTQNPSPRKEWVTPKEFREISYLK KLKVKKQDRIFPPESSAPAPLALPLS VTSAPTAVNSSAPADKPLSNMKILTL GKLSQNKDEAKAVIEKLGKLTGSA NKASLCISTKKEVEKMSKKMEEVKA ANVRVVCEDFLQDVSASTKSLQELL SAHSLSSWGAEVKAEPGEVVAPKGG SAAPSKKSKGAVKEEGVNVKSEKRMK LTLKGGAAVDPDSGLEHSAHVLEKG GKVFSAATLGLVDIVKGTNSYYKLQLL EDDKESRYWIFRSWGRVGTVIGSNK LEQMPKEDAVEHFMKLYEEKTGN AWHSKNFTKYPKKFYPLEIDYQDE EAVKCLTVKPGTKSKLKPQVQELVG MIFDVEISMKKALVEYIDLQKMPLG KLSRRQIQAAAYSILSEVQOAVSQGSS ESQILDLSNRFYTLIPHDFGMKKPPL LNNADSVQAKVEMLDNLLDIEVAYS LLRGGSDSSKDPIDVNYEKLKTDIK VVDRDSEEADEVIRKYVKNTHATTHN AYDLEVIDIFKIEREGESQRYKPFRL HNRRLWLHGSRTTNFAGILSOGLRI APPEAPVTGYMFGKGIYFADMVSKS ANYCHTSQGDPIGLILLGEVALGNM YELKHASHISKLPGKHSVKGLGKT TPDPSASITLEGVEVPLGTGIPSGVN DTCLLYNEYIVYDIAQVNLKYLKLLK FNFKTSW	True	False	4.622	5.0	4.678	2.765	2.016	2.88	2.826