

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
G3X8Y3	G3X8Y3_MOUSE	Mus musculus	NaN	28.757601	NaN	NaN	25153642;28528544	MPAVSLPPKENALFKRILRCYEHKO YRNLKFKCKQILSNPKFAEHGETLA MKGLTLNCLGKKEEAYELVRRGLRN DLKSHVCWHVYGLLQRSDDKKYDEAI KCYRNALKWDKDNLQILRDLSSLQI QMRDLEGYRETRYQLLQLRPAQRAS WIGYAIAYHLLLEDYEMAAKILEEFRK TQQTSPDKVDYEYSELLYQNVLR EAGLYREALEHLCTYEKQICKLAVE ETKGELLQLCRLEDAADVYRGLQE RNPENWAYYKGLEKALKPANMLER LKIYEEAWTKYPRGLVPRRLPLNFLS GEKFKECLDRFLRMNFSKGCPPVF NTLRSLYRDKEKVAIVEELVVGYES LKSCLFNPNDGKEEPTTLLWVQ YYLAQHVDKIGQPSIALEYINTAIEST PTLIELFLVKAKIYKHAGNIKEAARW MDEAQLDTADRFINSKCAKYMLKA NLIKEAEEMCSKFTREGTSAVENLN EMQCMWFQTECAQAYKAMNKFGE ALKKCHEIERHFIEITDDQDFHTYC MRKITLRSYVDLLKLEDVLRQHPFYF KAARIAIEIYLKLDNPLTDENKEHE ADTANMSDKELKLRNKORRAQKK AQIEEEKNAEKEKQQRNQRKQKDD DDDEEIGGPKEELIPEKLAKVETPLE EAIKFLTPLKNLVKNKIETHLFAFEIY FRKEKFLMLQSVKRAFAIDSSHPW LHECMIRLFHVSCEKDLPETVRTV LKQEMNRLFGATNPKNFNETFLKR NSDSLPHRLSAAKMVYYLDSSSQKR AIELATTLDGSLTNRNLQTCMEVLE ALCDGSLGDCKEAAEAYRASCHKLF PYALAFMPPGYEEDMKITVNGDSSA ETEELANEI	False	False	3.332	3.859	0.932	0.751	1.596	1.412	0.614