

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q04736	YES_MOUSE	Mus musculus	Tyrosine-protein kinase Yes	35.498311	NaN	T32;T334;T343;T424;T535	34678516;22645316	MGCIKSKENKSPAIKYTPENLTPVSP SASHYGV EHA TVAPTSSSTKGASVN FNSLSMTPFGGSSGVT PFGGASSSF SVVSSSYPTGLTGGVTIFVALYDYE A RTTEDLSFKKGERFQIINNT EGDWW EARS IATGKSGYIPSNYVVPADSIQAE EWYFGKMGRKDAERLLNPGNQRG IFLVRESETTKGAYSL SIRDWDEV RG DNVKHYKIRKLDNNGYYITTRAQFD TLQKLVKHYTEHADGLCHKLTTVCP TVKPTQGLAKDAWEIPRESLRLEV KLGQGC FGEVWMTWNGTTKVAIK TLKPGTMMPEAFLQEAQIMKLRH DKLVPLYAVVSEEPYIVTEFMSKGS LLDFLKEGDGKYLKLPQLVDMAAQI ADGMAYIERMNYIHRDLRAANILVG ENLICKIADFGLARLIEDNEYTARQG AKFPIKWTAPEAALYGRFTIKSDVWS FGILQTELVTKGRVPYPGMVNREVL EQVERGYRMPCPQGPCPESLHELMN LCWKKDPDERPTFEYIQSFLEDYFTA TEPQQPGENL	True	True	4.657	2.512	2.067	1.24	2.737	5.0	1.545