

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q6NXL1	Q6NXL1_MOUSE	Mus musculus	NaN	42.188058	NaN	NaN	28135057;40997131;28528544	MSQQGYVATPPYSQSQPGMGISPPH YGHYGDPSHASSPPGVMKPLGPSAA PSGMLPPGFLPPGPPQFGPNGAHTP GHPPQRFPGPPVNSVAPSYASGQT LPQSSYPGPGSTSSVTQLGSQFSSM QINSYGVGATPQSQGPPGPQAGAF QGPPQPAQPSILQPGHQVPPPPPTAL NGPGASPMSPPTHRODGLPGPAPLN AQYQPPPPGQTLGPGYPPQATNY GPMGGAQMSYGGFPGGPAQMA GPAPQLQRKLDPSIPSPQVIENDR ATRGGQVYTTNTRGQVPLVTTECVI QDQGHSSPRYIRCTTYCFPCTSDMA KQAQIPLAAVIKPFADIPPNETPLYLV NHGESGPVRCNRCKAYMCPFMQFI EGGRRYQCGFCSCVNEVPPFFYQHL DHIGRRLDHYEKPELSLGSYEYVATL DYCRKNKPPSPAFIFMIDVSYSNIK NGLVKLICEELKTALKRKPKEEHEET SAIRVGFITYNKVLHFFNVKSNLAQP QMMVVTDVGEVFPVLLDGLFVNYE ESQSVIHNLDDQIPEMFADSNENET VFAPVIQAGMEALKAECPGKLFIFH SSLPTAEAPGKLNRRDDKLVNTDK EKILFQPQTAVYESLAKDCVANSCSV TLFLFPQFVDSASLGLVPLLTGGTL YKYNVFIHSDSQRFITDLRNDIEKK IGFDAIMRVRTSTGFRATDFGGIFM NNTTDEVMAAIDCDKAVTVEFKHD DKLSEVGVGALIQCAVLYTTISGQRRL RIHNLALNCSTQLADLYKSCETDALI NFFAKSAFKAVLNQPLKAIREILVNQ TAHMLACYRKHCAASPSAASQILIPD SMKVLVPVYMNSLLKNCVLLSRSEIS PDERAYQRLVMTMGVADSQLFFY PLLLPIHTLDVKSAAALPPAVRCSESR LSEEGIFLLANGLNMFLWFGVGSPP ELIQGIFNVPSFAHINTDMTSLPEVG SPHSQQLRMIMNNIQKKPYSMKLI VVKQREQREMAFRQFLVEDKGLYG GSSYVDFLCCVHKEICQLLN	True	False	4.211	2.3	1.146	3.881	2.08	1.555	1.451

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