

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
M0R715	M0R715_RAT	Rattus norvegicus	NaN	23.439218	NaN	NaN	38843836	MGTPSHELNTTSSGSEVIRKALEEG LRRICVAQPVPFVQVLGVMIGAGV AVLVTAVLILLVRRRLRVOKTPAPEG PRYFRKRDKVLFYGRKIMRKVSQS TSSLVDTSVSTTSRPRMKKLKMMLN IAKKILRIQKETPTLQRKEPPPSVLEA DLTEGDLANSHLPSEVLYMLKNVRV LGHFEKPLFLELCRHMVFQRLGQG DYVFRPGQPDASIYVVDGLLELCLP GPDGKECVVKEVVPGDSVNSLLSIL DVITGHQHPQRTVSARAARDSTVLR LPVEAFSAVFTKYPESLVRVQIIMV RLQRVTFLLALHNYLGLTNEFSHEI QPLRLFPSPGLPTRTSPVRGSKRVVS TSGTEDTPKETSGRPLDPTGAPLPGP AGDPVKPTSLEAPSAPLLSRCISMPV DISGLQGGPRSDFDMAYERGRISVS LQEEASGGPQTASPRTPTQELREQP AGACEYSFCEDESATGGCPFGFYQG RQTSSIFEAAKRELAKLMRIEDPSLL NSRVLLHHAKAGTHARQGDQDVSL HFVLWGCLHVYQRMIDKAEVECLFV AQPGEVLVGLAVLTGEPLIFTLRAQR DCTFLRISKSDFYEMRAQPSVLSA AHTVAARMSPFVRQMDFAIDWTAV EAGRALYRQGDRSDCTYIVLNGRLR SVIQRGSKKELVGEYGRGDLIGVVE ALTRQPRATTVHAVRDTLAKLPEG TLGHIKRRYPQVVTRLIHLLSQKILG NLQQLQGGPFGSGLSVPHSELTNP ASNLSTVAILPVCAEVPMMMAFTLEL QHALQAIGPTLLLNSDIIRARLGASA LDSIQEFRLSGWLAQQEDAHRIVLY QTDTSLTPWTVRCLRQADCILIVGLG DQEPTVGQLEQMLENTAVRALKQLV LLHREEGPGPRTVWLNMRSWCS GHLHLRCPRLFSRRSPAKLHELVE KVFSRRADRHSDFSRLARVLTGNTI ALVLGGGGARGCSHIGVLKALEEAG VPVDLVGGTSGSFIGALYAEERSAS RTKQRAREWAKSMTSVLEPVLDLTY PVTSMFTGSAFNRSIHRVFDKQIE DLWLPYFNVTITDITASAMRVHKDGS LWRYVRASMTLSGYLPLCDPKDGH LLMDGGYINNLPAIARSMGARTVI AIDVGSQDETDLSTYGDLSGWWLL WKRLNPWADKVKVPDMAEIQSRLA YVSCVRQLEVVKSSSYCEYLPSIDC FKTMDFGKFDQIYDVGYQYKAVFG GWTRGEVIEKMLTDRRSTDLNESRR ADILAFPSSGFTDLAEIVSRIEPPTSY VSDGCADGEESDCLTEYEEDAGPDC SRDEGGSPEGASPSTASEVEEEKSTL RQRFLPQETGAQ	False	False	2.491	0.823	0.919	3.967	0.678	1.667	1.321