

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
P59729	RIN3_MOUSE	Mus musculus	Ras and Rab interactor 3	25.875327	T862	NaN	30059200	MRRAEAPSSAHPAGPIPDAGKGEGE EDEEKDGTRELGLSTTPRNCIPRRGIS VLEKLVKTCPVWLQGLGQAEAAKI LQQEMAGMFLVCRDNNLKQLVLCV HFPSLKGSSAEVLEYPIKEEKAILYLE GSVLFVEDIFRLIAFYCVSRDLLPFTL RLPQAILEASSFLELETISNLGLGFW DSSLNSRGSAEPLRSPAGTPASSSL RPTTHYANCSCIEELSVGNDRLWVF NPIFIEDCILPADPPPLPTGSYPPRPT PATPDATSPTS KGSPPRRPPPPPLPT VPPTGPARPLAPPVPPAGPLPNSPLT PTSHLAPHAPGPPGHSNQPPMTACE SLPRPAVGLGPFGEEMKPGTTPNP LHQAPPPPLPKKALPAAPRRRISE RVSLESQNVGTSTDRDHSGISRTAS LNLPPQSTVSSLGDRPRTTEQSQD TEAKASHADSIPVPPGKAKQPPVPPP RKKRVSRQLASTLLSPLEPIQEASS EKQATGASWEGLSVPRQAGMOHLQ VQSSSCPQSSPEFKGSQASLSDSLG VPASAADQDSYSTSSAEELFSSPN VKKKPSMILDKARHRLSFVVFASVF HAFLLSSDRKLYKKVELAQDKSSYF GSLVQDYKVYSLEMMARQTSSTEM LQEIRTMMTQLKSYLLQSTELKALV EPTLHSEEELEAIVESALYKCVLKPL KEAINSSLEIHSRDGSLQQLKENQL VVLATTTTDLGVTTTSVPEVAVMEKIL QKLTSMHKAYSPGKKISILLKTCKLIY DSMALGNPGKPYGADDFLPVLMYVL ARSNLTEMLLNVEYMMELMDPALQ LGECSYLLTTYGALEHIKNYDKITV TRQLSVEVQDSIHRWERRTLNKKAR ASRSSVQDFICVSYLKPEQQSRTLAS RADTAAQALCAQCAEKFEVSQPQDY RLFVLVDGRCFQLADEALPHRIKGYL LRSEPKRDFHFVYRPQDSGKDASSQ PCIIVREPNFL	True	False	4.289	3.161	1.187	0.929	1.371	1.524	1.328