

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
015034	RIMB2_HUMAN	Homo sapiens	RIMS-binding protein 2	27.086674	S665;T666;T667	S704;S712;S832;S839;T841	28657654	MREAAERRQQQLQLEHDQALAVLSA KQQEIDLLQKSKVRELEEKRCRTQSE QFNLLSRDLEKFRQHAGKIDLLGGS AVAPLDISTAPSKPFPQFMNGLATSL GKGQESAIGGSSAIGEYIRPLPQPGD RPEPLSAKPTFLSRSGSARCRSESD MENERNSENTSKQRYSGKVHLCVAR YSYNPFDDGPNENPEAELPLTAGKYL YVYGDMDDEDGFYEGELLDGQRGLV PSNFVDFVQDNESRLASTLGNEDQ QNFINHSGIGLEGEHILDHSPTHID AGITDNSAGTLDVNIDDIGEDIVPYP RKITLIKQLAKSVIVGWEPVAVPPGW GTVSSYNVLVDKETRMNLTGSRK ALIEKLNMAACTYRISVQCVTSRGSS DELQCTLLVGKDVVAPSHLRVDNI TQISAQLSWLPTNSNYSHVIFLNREE FDIVKAARYKYQFFNLRPNMAYKVK VLAKPHQMPWQPLEQREKKEAFV EFSTLPAGPPAPPQDVTVQAGVTPAT IRVSWRPPVLTPTGLSNGANVTGYG VYAKGQ RVAEVIFFPTADSTAVELVRL RSLEAKGVTVRTLSAQGESVDSAVA AVPPELLVPTPHPRPAPQSKPLASS GVPETKDEHLGPHARMDEAWEQSR APGPVHGHMLEPPVGPGRRSPPSPSR ILPQPQGTVPVSTTVAKAMAREAAQR VAESSRLEKRSVFLERSAGQYAAAD EEDAYDSPDFKRRGASVDDFLKGSE LGKQPHCCHGDEYHTESSRGSDLS DIMEEDEEELYSEMQLLEDGRRRPS GTSHNALKILGNPASAGRVDHMGR RFRPGSAGPQRSRPVTVPSIDYGRD RLSPDFYEESETDPGAEELPARIFVA LFDYDPLTMSNPDAEEELPFKEG QIIKYYGDKDADGFYRGETCARLGLI PCNMVSEIQADDEEMMDQLLRQGF LPLNTPVEKIERRRSGRRHSVSTRR MVALYDYPRESSPNVDVEAELTFC TGDITVFGIIDEEDGFYYGELNGOKG LVPSNFLEEVDDVEVYLSDAPSHYS QDTPMRSAKRKRSVHFTF	False	False	1.192	1.269	0.818	0.735	1.94	3.558	0.929