	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	ex re
0	Q9UQ26	RIMS2_HUMAN	Homo sapiens	Regulating synaptic membrane exocytosis protein 2	25.946761	S1409	S400;T689;S791;S794;S1148;S1396;S1399	28657654;37217939	MSAPVGPRGRLAPIPAASQPPLQPE MPDLSHLTEEERKIILAVMDRQKKK VKEEHKPQLTQWFPFSGITELVNNV LQPQQKQQNEKEPQTKLHQQFEMY KEQVKKMGEESQQQQEQKGDAPTC GICHKTKFADGCGHNCSYCQTKFCA RCGGRVSLRSNKVMWVCNLCRKQQ EILTKSGAWFYNSGSNTPQQPDQKV LRGLRNEEAPQEKKPKLHEQTQFQG PSGDLSVPAVEKSRSHGLTRQHSIK NGSGVKHHIASDIASDRKRSPSVSR DQNRRYDQREEREEYSQYATSDTA MPRSPSDYADRRSQHEPQFYEDSD HLSYRDSNRRSHRHSKEYIVDDEDV ESRDEYERQRREEEYQSRYRSDPNL ARYPVKPQPYEEQMRIHAEVSRARH ERRHSDVSLANADLEDSRISMLRM DRPSRQRSISERRAAMENQRSYSME RTREAQGPSSYAQRTTNHSPPTPRR SPLPIDRPDLRRTDSLRKQHHLDPSS AVRKTKREKMETMLRNDSLSSDQS ESVRPPPPKPHKSKKGGKMRQISLS SSEEELASTPEYTSCDDVEIESESVS EKGDSQKGKRKTSEQAVLSDSNTRS ERQKEMMYFGGHSLEEDLEWSEPQ IKDSGVDTCSSTTLNEEHSHSDKHP VTWQPSKDGDRLIGRILLNKRLKDG SVPRDSGAMLGLKVVGGKMTESGR LCAFITKVKKGSLADTVGHLRPGDEV LEWNGRLLQGATFEEVYNIILESKPE PQVELVVSRPIGDIPRIPDSTHAQLES SSSSFESQKMDRPSISVTSPMSPGM LRDVPQFLSGQLSIKLWFDKVGHQL IVTILGAKDLPSREDGRPRNPYVKIYF LPDRSDKNKRRTKTVKKTLEPKWN QTFIYSPVHRREFRERMLEITLWDQ ARVREEESEFLGEILIELETALLDDEP HWYKLQTHDVSSLPLPHPSPYMPRR QLHGESPTRRLQRSKRISDSEVSDY DCDDGIGVVSDYRHDGRDLQSSTLS VPEQVMSSNHCSPSGSPHRVDVIGR TRSWSPSVPPPQSRNVEQGLRGTRT MTGHYNTISRMDRHRVMDDHYSPD RDRDCEAADRQPYHRSRSTEQRPLL ERTTTRSRSTERPDTNLMRSMPSLM TGRSAPPSPALSRSHPRTGSVQTSPS STPVAGRRGRQLPQLPPKGTLDRKA GGKKLRSTVQRSTETGLAVEMRNW MTRQASRESTDGSMNSYSSEGNLIF PGVRLASDSQFSDFLDGLGPAQLVG RQTLATPAMGDIQVGMMDKKGQLE VEIIRARGLVVKPGSKTLPAPYVKVYL LDNGVCIAKKKTKVARKTLEPLYQQL LSFEESPQGKVLQIIVWGDYGRMDH KSFMGVAQILLDELELSNMVIGWFK LFPPSSLVDPTLAPLTRRASQSSLES STGPSYSRS	None	None	None	None	None	None	None	None	N