

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	extracel region
Q64700	RBL2_MOUSE	Mus musculus	Retinoblastoma-like protein 2	25.590209	NaN	S410;T414;S636;T639;S659;S669;S684;S942;S946;S960;S965;S967;T968;S975;S976;T980;S1031;S1064;S1076;S1108	35822049	MASGGNQSPPPPPAAAASSEEEEEED GDAADRAQAGSPSHQIQRFEEELC SRLNMDEAARAEAWSSYRSMSESY TLEGNDLHWLACALYVACRKSVPV SKGTAEAGNYVSLTRILRCSEQSLIEF FNKMKKWEDMANLPPHFRERTERL ERNFTVSAVIFKYEPIFQDIFKYPOE EQPRQQRGRKQRRQFCITSEIFHFC WVLFYAKGNFPMISDDLVSNSYHLL LCALDLVYGNALQCSNRKELVNPNF KGLSEDCPKDSKASSDPPCVIEKLC SLHDGLVLEAKGIKEHFVWPKYIRKLF EKLLKKGKEENLTGFLEPGNFGESF KAVNKAYEEYVLAAGNLDERVFLGE DAEEEVGTLSRCLSAASGTEAERT QMRDILQOHLDKSKALRVCTPLTGV RYVQENSPCVTPVSTAAHSLSRLHT MLSGLRNAPSEKLERILRSCSRDPT QAIADRLKEMYEYVQFQDENFNS NCAKEIANKHFRFAEMLYYKLVESVI EQEQKRLGDMDLSGVLEHDAFHRS LLACCLEVVAFSHKPPGNFPFIAEIF DVPHYHFYKVEVFIRAEDGLCREVV KHLNQIEEQILDHLAWKTKSPLWDR IRDNENRVPTCEEVMPQNLERTDE IYIAGSPLTPRRVGEVRADAGGLGRSI TSPTTLYDRYSSPTVSTTRRRLFEND SPSEGSTSGRIPPQLVNAVVPQNVV GETVSVTPVPGQTLVMTATVTAN NGQVTIPVQGANENGGITFFPVQV NVGGQAQAVAGSIQPLSAQALAGSL SSQQVTGTTLQVPGVVAIQISPGGQ QQNPGQPLTSSSIRPKRTSSLALFFR KVYYLAGVRLRDLCKIKLDSDELKKI WTCFEFSIIQCTELMMDRHLQDL MCAIYVMAKVTKEDRSFQNMRCYR TQPARSQVYRSVLIKGRNRSGSS ESRSHQNSPTELNDRASRDSSPVM RSNSTLPVPQSSAPPTPTRLTGASS DVEEEERGLIQFYNNIYRKQIQFA MKYSQANAQTDTPPLSPYPFVRTGS PRRVQLSQSHPIYISPHNNEAMPSP REKIFYYSNSPSKRLREINSMIRTG ETPTKRGILLDDGSESPAKRICPEN HSALLRRLQDVANDRGSQ	True	False	3.704	5.0	2.136	1.6	1.038	1.878	1.961

0