

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q96T37	RBM15_HUMAN	Homo sapiens	RNA-binding protein 15	25.019945	T401;S692;S757	S109;S179;S208;S210;S253;S257;S259;T266;S292;S294;S365;T568;S622;S656;S670;S674;S700;S741;S765;S767;S781;S935	25367160;37217939;40596516;40914422;30620550;34019948	MRTAGRDPVPRRSFWRRAVPLCET SAGRRVTQLRGDDLRRPATMKGKE RSPVKAKRSGGEDSTSRGERSKKL GGSGGNSGSSGKTDSGGGSRRL HLDKSSSRGGSREYDTGGGSSSSRL HSYSSPSTKNSSGGGESRSSSRGGG GESRSSGAASSAPGGGDGAEYKTLKI SELGSQLSDEAVEDGLFHEFKRFGD VSVKISHLSGSGGDERVAFVNFRR PEDARAAKHARGRLVLYDRPLKIEAV YVSRRRRSRPLDKDTPPSASVVGAS VGGHRHPPGGGGQRSLSPGGAAAL GYRDYRLQQLALGRLPPPPPLPRD LERERDYPFYERVRPAYSLEPRVGAG AGAAPFREVD EISPEDDQRANRTLF LGNLDITVTESDLRRAFDRFGVITEV DIKRPSRGQTSYGFLENDLMSH RAKLAMSGKIIIRNPIKIGYGKATPTT RLWVGGGLGPVVPLAALAREDFRFGT IRTIDYRKGDSSWAYIQYESLDAAHAA WTHMRGFPLGGPDRRLRVDFADTE HRYQQYLQPLPLTHYELVTDAFGH RAPDPLRGARDRTPPLLYRDRDRDL YPDSDWVPPPPVRESTRTAATSVP AYEPLDSLDRRRDGSWLDLDRDRGDR DLPSSRDQPRKRLPEESGGRHLDR SPESDRPRKRHCAPSPDRSPELSSS RDRYNSDNDRSRLLLLERPSPIRDR RGSLEKSQGDKRDRKNSASAERDR KHRTTAPTEGKSPKKEDRSDGSAP STSTASSKLKSPSQKQDGGTAPVASA SPKLCCLAWQGMILLKNSNFPSNMH LLQGDLOVASSLLVEGSTGGKVAQL KITQRLRLDQPKLDEVTRRIKVAGPN GYAILLAVPGSSDSRSSSSAASDTA TSTQRPLRNLVSYLKQKQAAGVISLP VGGNKDKENTGVLFHAFPPCEFSQQ FLDSPAKALAKSEEDYLVMIIVRFGF FQIGVRYENKKRENALATLL	True	False	2.745	5.0	1.879	1.315	0.96