

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum
Q9HBD1	RC3H2_HUMAN	Homo sapiens	Roquin-2	33.827608	T468;T470;T471;T472;S478;S492;S497;S556;T561;S562;S566;S593;S645;S714;S716;S725;S726;S731;T755;T756;T769;S770;T801;T843;T883;T887;S897;S903;S909;S927;S930;S952;T959;S960	S549;S808;S983;S1119	40136647;35132862;35289036;32574038;30059200;41350704;39302247;28657654;40307207;38665916;35254053;33214551;34019948;31492838;29237092;30620550;39534244;37340703;31637018;35138101;39531497	MPVQAAQWTEFLSCPICYNEFDENVHKPISLGCSTVCKTCLNKLHRKACPFQDTAINTDIDVLPVNFALLQLVGAQVPDHDQSIKLSNLGENKHYEVAKKVEDLALYLKPLSGGKGVASLNGSALSRPMPQRKLVTLVNCQLVEEGRVRA MRAARSLGERTVTELILQHONPQQL SANLWAAVRRARGCQFLGPAMQEEA LKLVLLALEDGSALS RKVLLVLFVVQR LEPRFPQASKTSIGHVVQLLYRASCF KVTKRDESSLMQLKEEFRSYEALR REHDAQIVHIAMEAGLRISPEQWSS LLYGDLAHKSHMQSIHDKLQSPESFA KSVQELTIVLQRTGDPANLNRRLRPH LELLANIDPNPDVSPWTEQLENAM VAVKTVVHGLVDFIQNYSRKGHETP QPQNSKYKTSMCRDLRQGGCPR GTNCTFAHSQEELEKYRLRNKKINA TVRTFPLLNKVGVNNTVTTAGNVIS VIGSTETTGVKIVPSTNGISNAENSVS QLISRSTDSTLRALETVKKVGKVGAN GQNAAGPSADSVTENKIGSPKTPV SNVAATSAGPSNVGTELNVPQKSS PFLTRVPVYPHSENIQYFQDPRTQI PFEVPOYPQTGYPPPTVPAGVAPC VPRFVRSNNVPESSLPPASMPYADH YSTFSPRDRMNSSPYQPPPPOPYGP VPPVPSGMYAPVYDSRRIRWPPMYQ RDDIIRSNSLPPMDVMHSSVYQTL RERYNSLDGYYSVACQPPSEPRTTVP LPREPCGHLKTSCEEQIRRPDQWA QYHTQKAPLVSSTLPVATQSPPTPPSP LFSVDFRADFSESVSGTKFEEDHLS HYSWPWCGTIGSCINAIDSEPKDVIA NSNAVMDLDSGDVKRRVHLFETQ RRTKEEDPIIPFSDGPIISKWGAISRS SRTGYHTTDPVQATASQGSATKPISV SDYVPYVNAVDSRWSSYGNEATSSA HYVERDRFIVTDLSGHRKHSSTGDL LSLELQQAQKSNLQLQREANALAMQ QKWNLSLDEGRHLTLNLLSKEIELRN GELQSDYTEDATDTKPRDIELELSA LDTDEPDGQSEPIEILDIQLGISSQN DQLLNGMAVENGHPVQQHQKEPPK QKKQSLGEDHVILEEQKTLIPVTSFCF SQPLPVSISNASCLPITTSVSGNLIL KTHVMSEDKNDFLKPVANGKMNVS	False	False	2.29	3.956	0.783	1.14