

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
Q86X10	RLGPB_HUMAN	Homo sapiens	Ral GTPase-activating protein subunit beta	29.413267	S343;T476	S359;T363;T379;S421;S720;T734;S1285	28657654;38253038	<p>MYSEWRSLLHLVIQNDQGHSTVLHS YVESVGREVANAVRPLGQVLGTPS VAGSENLLKTDKEVKWTVMEVICYGL TLPLDGETVKYCVDVYTDWIMALVL PKDSIPLPVIKEPNQYVQTIKHLQNL LFVPRQEQGSSQIRLCLQVLRAIQKL ARESSLMARETWVLLLLFLLQINDIL LAPPTVOGGIAENLAEKLGVLFEVW LLACTRCFPTPPYWKTAKEVMANW RHHPAVVEQWSKVICALTSRLLRFT YGPSFPAFKVPDEASLIPPEMDNE CVAQTWFRFLHMLSNPVDLSNPAII SSTPKFOEQFLNVS GMPQELNOYPC LKHLPOIFFRAMRGISCLVDAFLGIS RPRSASAPPTPVNRLSMPQSAAVST TPPHNRRHRAVTVNKATMKTSTVST AHASKVQHTSSTSPSSPNTSSE PRPLPAPRRPKVNSILNLFGSWLFDA AFVHCKLHNGINRDSMTAITTQAS MEFRRKGSQMSTDTMVSNNPMFDAS EFPDNYEAGRAEACGTLCRIFCSKKT GEEILPAYLSRFYMLLIQQLINDYV CHPVLASVILNSPPLFCCDLKIDV VPYFISALETILPDRELSKFKSYVNP ELRRSSINILLSLLPLPHHFGTVKSE VVLEGFNSDDSSSYDKPITFLSLKL RLVNILGALQTEETDPNNTQMILGA MLNIVQDSALLEAIGCOMEMGGGE NNLKSHSRTNSGISSASGSTEPTTP DSERPAQALLRDYALNTDSAAGLLIR SIHLVTQRLNSQWRQDMSISLALE LLSGLAKVKVMVDSGDRKRAISSVC TYIVYQCSRPAPLHSRDLHSMIVAFA QCLCVWLTEHPDMLDEKDKLKEVL EIVELGISGSKSKNNEQEVKYKGD EPNPASMRVKDAEATLTCIMQLL AFPSPSGPASPCSLVNETTLIKYSRLP TINKHSFRYFVLDNSVILAMLEQPLG NEQNDFFPSVTVLVRGMSGRLAWA QQLCLLPRGAKANQKLFVPEPRPVP KNDVGFKYVKHRPFPEEVDKIPFV KADLSIPDLHEIVTEELERHEKLR GMAQQIAYEIHLEQQSEELQKRSF PDPVTDCKPPPAQEFQTARLFLSHF GFLSLEALKEPANSRLPPHIALDSTI PGFFDDIGYLDLLPCRPFDTVFIFYM KPGQKTNQEILKNVSSRTVQPHFL EFLSLGWSVDVGRHPGWTGHVST SWSINCCDDGEGSQQVEISSDIGA SIFNGQKVVLYADALTEIAFVVPSP VESLTDLSLENSISDQSDSNMDLMP GILKQPSLTLLEFPNHTDNLNSSQR LSPSSRMKLPQGRPVVPLGPETRV SVVVERYDDIENFPLSELMTEISTG VETTANSSTSLRSTTLEKEVPVIFHP LNTGLFRIQIGATGKFNMVPLVDG MIVSRRALGFLVRQTVINICRRKLE SDSYSPPHVRRKQKITDIVNKYRNKQ LEPEFYTSLFQEVGLKNCSS</p>	False	False	3.096	2.558	1.285	0.672	0.95	1.255