

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochondrion | endoplasmic reticulum | golgi apparatus |
|--------------|-------------|--------------|----------------------------------|--------------|---------------|--|---|--|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|
| Q68EM7 | RHG17_HUMAN | Homo sapiens | Rho GTPase-activating protein 17 | 14.378567 | S820 | S484;S575;T679;T682;S702;S704;T753;T757;T759;S762;T763 | 30379171;31492838;35254053;29485866;37217939;38665916 | MKKQFNRMKQLANQTVGRAEKTEV LSEDLQIERRLDTVRSICHHSHKRL VACFQGHQGTDAERRHKKLPLTALA QNMQEASTQLEDSLLGKMLETCGD AENQLALELSQHEVFVKEIVDPLY GIAEVEIPNIQKQRKQLARLVLDWDS VRARWNQAHKSSGTNFQGLPSKIDT LKEEMDEAGNKVEQCKDQLAADMV NFMMAKEGEYGKFFVTLLAEQADYHR KALAVLEKTLPEMRAHQDKWAEKP AFGTPLEEHLKRSGREIALPIEACVM LLETGMKEEGLFRIGAGASKLKKL KAALDCSTSHLDEFYSDPHAVAGAL KSYLRELPEPLMTFNLYEWTQVAS VQDQDKKLQDLWRTCOKLPQNFV NFRYLIKFLAKLAQTSVDVNMKTPSNI AIVLGNLLWARNEGTLAEMAAAATS VHVVAVIEPIIQHADWFFFPEVEFNV SEAFVPLTTPSSNHSFHTGNDSDSG TLERKRPAVMMEGDLVKESFGV KLMDFQAHRRGGTLNRKHISPAFQP PLPTDGVTVVPAPEPPPPQSSRAES SSGGGTVPSSAGILEQGSPGDGSP KPKDPVSAAVPAPGRNNSQIASGQN QPQAAAGSHQLSMGQPHNAAGPSP HTLRRAVKKPAPAPPKGNPPPGHP GGQSSSGTSQHPPSLSPKPPTRSPSP PTQHTGQPPGQPSAPSQLSAPRRYS SSLSPIQAPNHPPQPPTQATPLMHT KPNSQGPNNMMPALPSEHGLEQPSH TPPQTPTPPSTPPLGKQNPSPAPQT LAGGNPETAQPHAGTLPRPRVVPKP RNRPSVPPPPQPPGVHSAGDSSLTN TAPTASKIVTDSNSRVSEPHRSIFPE MHSDSASKDVPGRILLDIDNDTEST AL | True | False | 4.71 | 3.837 | 0.964 | 0.792 | 1.16 |