

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q9Y3P9	RBGP1_HUMAN	Homo sapiens	Rab GTPase-activating protein 1	24.348488	NaN	S42;S360;T996	34725712;23301498	MDDKASVGGKISVSSDSVSTLNSEDF VLVSRQGDDETPSTNNGSDEKTKGLK IVGNGSEQQLQKELADVLMDPPMD DQPGKEKELVKRSQLDGEKDGPLSN QLSASSTINPVVLVGLQKPEMSLFPVK PGQGDSEASSPFTPVADEDSVVFVK LTYLGCASVNA PRSEVEALRMMMSIL RSQCQISLDVTLVSPNVSEGIVRLLD PQINTEIANYPYKILFCVRGHDGTP ESDCFAFTESHYNAELFRHVFRCIEI QEA VSRILYSFATAFRRS AKQTPLSA TAAFQTPDSDIFTFVSVLEIKEDDGK GYFSAVPKDKDRQCFKLRQIDKKIV IYVQQTNNKELAIERCFGLLLSPGKD VRNSDMHLLDLESMGKSSDGKSYVI TGSWNPKSPHFQVNEETPKDKVLF MTTAVDLVITEVQEPVRFLLLETKVRV CSPNERLFWPFSKRSTTENFFLKLK QIKQRERKNNTDTLYEVVCLSESE RERRKTTASPSVRLPQSGSQSSVIPS PPEDDEEEDNDEPLLSGSGDVSKEC AEKILETWGELLSKWHLNLRVFPKQ LSSLVRNGVPEALRGEVWQLLAGCH NNDHLVEKYRILITKESPDQSAITRDI NRTFPAHDYFKDTGGDGGDSLYKIC KAYSVDDEIIGYCQGGQFLAAVLLH MPEEQAFSVLVKIMFDYGLRELFLKQ NFEDLHCKFYQLERLMOEYIPDLYN HFLDISLEAHMYASQWFLTLFTAKF PLYMVFHIDLLCEGISVIFNVALGL LKTSKDDLLTDFEGALKFFRVQLPK RYRSEENAKKLMELACNMKISQKKL KKYEKEYHTMREQQAQEDPIERFE RENRRLQEANMRLEQENDDLAHEL VTSKIALRKDLDNAEEKADALNKEL LMTKQKLIDAEKRRLEESAQLK EMCRRELDKAESEIKNNSIIGDYKQ ICSQLSERLEKQQTANKVIEIKIRQK VDDCERCREFFNKEGRVKGISSTKE VLDEDTDEEKETLKNQLREMELELA QTKLQVVEACKIQDLEHHLGLALN EVQAAKKTWFNRTLSSIKTATGVQG KETC	True	False	4.582	1.892	0.718	1.424	1.419	1.247	0.906

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