	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellula region
0	Q12774	ARHG5_HUMAN	Homo sapiens	Rho guanine nucleotide exchange factor 5	25.924869	NaN	S184;S445;S450;S983;S1011;S1044;S1126	32574038	MEAEEAQRGASPPISAIEEFSIIPEAP MRSSQVSALGLEAQEDEDPSYKWR EEHRLSATQQSELRDVCDYAIETMP SFPKEGSADVEPNQESLVAEACDTP EHWEAVPQSLAGRQARTLAPPELW ACPIQSEHLDMAPFSSDLGSEEEEV EFWPGLTSLTLGSGQAEEEEETSSD NSGQTRYYSPCEEHPAETNQNEGSE SGTIRQGEELPPEELQESQGLLHPQ EVQVLEEQGQQEAGFRGEGTLREDV CADGLLGEEQMIEQVNDEKGEQKQ KQEQVQDVMLGRQGERMGLTGEPE GLNDGEWEQEDMERKAQGQGGPE QGEERKRELQVPEENRADSQDEKS QTFLGKSEEVTGKQEDHGIKEKGVP VSGQEAKEPESWDGGRLGAVGRAR SREEENEHHGPSMPALIAPEDSPHC DLFPGASYLMTQIPGTQTESRAEELS PAALSPSLEPIRCSHQPISLLGSFLTE ESPDKEIDQNSQQEESRLRKGTVSS QGTEVVFASASVTPPRTPDSAPPSPA EAYPITPASVSARPPVAFPRRETSCA ARAPETASAPLSMDDPSPCGTSEMC PAALYGFPSTGTSPPRPPANSTGTVQ HLRSDSFPGSHRTEQTPDLVGMLLS YSHSELPQRPPKPAIYSSVTPRRDRR SGRDYSTVSASPTALSTLKQDSQESI SNLERPSSPPSIQPWVSPHNPAFATE SPAYGSSPSFVSMEDVRIHEPLPPPP PQRRDTHPSVVETDGHARVVVPTLK QHSHPPPLALGSGLHAPHKGPLPQA SDPAVARQHRPLPSTPDSSHHAQAT PRWRYNKPLPPTPDLPQPHLPPISAP GSSRIYRPLPPLPIIDPPTEPPPLPPKS RGRSRSTRGGHMNSGGHAKTRPAC QDWTVPLPASAGRTSWPPATARSTE SFTSTSRSKSEVSPGMAFSNMTNFL CPSSPTTPWTPELQGPTSKDEAGVS EHPEAPAREPLRRTTPQQGASGPGR SPVGQARQPEKPSHLHLEKASSWPH RRDSGRPPGDSSGQAVAPSEGANKH KGWSRQGLRRPSILPEGSSDSRGPA VEKHPGPSDTVVFREKKPKEVMGGF SRRCSKLINSSQLLYQEYSDVVLNKE IQSQQRLESLSETPGPSSPRQPRKAL VSSESYLQRLSMASSGSLWQEIPVV RNSTVLLSMTHEDQKLQEVKFELIV SEASYLRSLNIAVDHFQLSTSLRATL SNQEHQWLFSRLQDVRDVSATFLS DLEENFENNIFSFQVCDVVLNHAPD FRRVYLPYVTNQTYQERTFQSLMNS NSNFREVLEKLESDPVCQRLSLKSF LILPFQRITRLKLLLQNILKRTQPGSS EEAEATKAHHALEQLIRDCNNNVQS MRRTEELIYLSQKIEFECKIFPLISQS RWLVKSGELTALEFSASPGLRRKLN TRPVHLHLFNDCLLLSRPREGSRFL VFDHAPFSSIRGEKCEMKLHGPHKN LFRLFLRQNTQGAQAEFLFRTETQS EKLRWISALAMPREELDLLECYNSP QVQCLRAYKPRENDELALEKADVVM VTQQSSDGWLEGVRLSDGERGWFP VQQVEFISNPEVRAQNLKEAHRVKT AKLQLVEQQA	True	False	4.443	5.0	0.857	0.544	0.524	4.259	0.889