

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	g a
P98171	RHG04_HUMAN	Homo sapiens	Rho GTPase-activating protein 4	35.287989	S199;T202;T203;T204;T208;T686;S906;S913	S860;S901;S906	29351928;31492838;28657654;31637018;40914422;35254053;39302247;34019948;39531497	MAAHGKLRERRGLQAEYETQVKEMRWQLSEQLRCLLELQGLRRELLQELAEFMRRRAEVELEYSRGLKLAERFSSRGGRLGSSREHQSFKEPSLLSP LHCWAVLLQHTRRQSSRESAALSEVL AGPLAQRLSHIAEDVGRLVKKSRL EQQLQDELLEVVSELTAKKTYQAY HMEVNAEAKLREAERQEEKRAGR SVPTTTAGATEAGPLRKSSLKGGRL VEKRQAKFMEHKLKCTKARNEYLLS LASVNAAVSNYYLHDVLDLMDCCDT GFHLALGQVLRSYTAAESRTQASQV QGLGSLEEAVEALDPPGDKAKVLEV HATVFCPPPLRFDYHPPHDGDEVAEIC VEMELRDEILPRAQNIQSRLDRQTIE TEEVNKTLKATLQALLEVVASDDGD VLDSFQTSPTSELSKSTSSDPGSRQA GRRRGQQQETETTFYLTKLQEYLSGR SILAKLQAKHEKLEALQRGDKEEQ EVSWTQYTQRKFKSROPRPSSQYN QRLFGGDMKFIQSSGQPVPLVVES CIRFINLNGLOHEGIFRVSGAQLRVS EIRDAFERGEDPLVEGCTAHDLDV AGVLKLYFRSLEPPLFPDDLFGELLA SSELEATAERVEHVSRLWRLPAPV LVVLRYLFTFLNHLAQYSDENMMD PYNLAVCFGPTLLPVPAGQDPVALQ GRVNLVQTLIVQPDVFPPLTSLPG PVYEKCMAPPSASCLGDAQLESIGA DNEPELEAEMPAQEDDLEGVVEAV ACFAYTGRTAQELSFRRGDVLRRLHE RASSDWWRGEHNGMRGLIPHKYIT LPAGTEKQVVGAGLQTAGESGSSPE GLLASELVHRPEPCTSPPEAMGPSGH RRRCLVPASPEQHVEVDKAVAQNM DSVFKELGKTSVRQGLGPASTTSPS PGPRSPKAPPSSRLGRNKGFSRGP APASPSASHPQGLDITPKPH	True	False	4.692	4.756	0.906	False	0