

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|-------------|--------------|----------------------------------|--------------|---------------|-----------------------|-------------------|--|
| Q7Z6I6 | RHG30_HUMAN | Homo sapiens | Rho GTPase-activating protein 30 | 16.686362 | S632;S678 | S576;S996 | 35132862;29351928 | MKSRQKGKKKGSAKERVFGCDLQE HLQHSGQEVPPQVLKSCAEFVEEYGV VDGIYRLSGVSSNIQKLRQEFESERK PDLRRDVYLDIHCVSSLCKAYFREL PDPLLYRLYDKFAEAVGVQLEPERL VKILEVLRELPVPNYRTLEFLMRHLV HMASFSAQTNMHARNLAIWAPNL LRSKDIEASGFNGTAAFMEVRVQSIV VEFILTHVDQLFGGAALSGGEVESG WRSLPGTRASGSPEDLMRPLPYHL PSILQAGDGPPQMRPYHTIIEIAEHK RKGSLKVRKWRSIFNLGRSGHETKR KLPRGAEDREDKSNKGLRPAKSM DLSAAAGASDEPEGLVGPSSPRPSP LLPESENDSIEAAEGEQEPEAEALG GTNSEPGTPRAGRSAIRAGGSSRAE RCAGVHISDPYVNLPLHITSILSVPP NIISNVSLARLTRGLECPALQHRPSP ASGPGPGPLGPGPPDEKLEASPASS PLADSGPDDLAPALEDSLQEVQDS FSFLEDSSSSEPEWVGAEDGEVAQA EAAGAAFSPGEDDPGMGYLEELLGV GPQVEEFSVEPPLDDLSDLDEAQFVL APSCCSLDSAGPRPEVEEENGEEVF LSAYDDLSPLLGPKPIWKSGSLEG EAAGCGRQALGQGGEQACWEVGE DKQAEPGGRLDIREEAEGSPETKVE AGKASEDRGEAGGSQETKVRLREGS REETEAKKEKSKGQKKADSMKAG VEEPGGDEYTDEKEKEIEREEDQR EEAQVEAGRDLQGAQEDQVAEEK WEVVQKQEAEGVREDEDKGQREKG YHEARKDQGDGDSRSPEAATEGGA GEVSKERESGDGEAEGDQRAGGYL EEDTLSESGVASLEVDCAKEGNPH SSEMEEVAPQPPQPEEMEPEGQSP DGCLCPCSLGLGGVGMRLASTLVQV QQVRSVPVPPKQFAKMPSAMCSK IHVAPANPCPRPGRDGTGERAWG SRASRSSWRNGGSLSFDAVALARD RQRTEAQGVRRQTCTEGGDYCLIP RTSPCSMISAHSRPLSCLLPSEGA EGSGRSRSLPPEPQVPDPLLSSQ RRSYAFETQANPGKGEGL |