

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q640N3-2	RHG30_MOUSE	Mus musculus	Isoform 2 of Rho GTPase-activating protein 30	14.32282	NaN	S578;S875;S996	36084651	MKSRQKGKKKGSSKERVFGCDLRE HLQHSGQEVPPQVLRSCAEFVQEQYGV VDGIYRLSGVSSNIQKLRQEFETERK PDLRRDVYLDIHCVSSLCKAYFREL PDPLLTYRLYDKFAEAVAVQLEPERL VKILEVLQELPIQNYRTLEFLMRHLV HMASFSAQTNMHARNLAIWVAPNL LRSKDIEASGFNGTAAFMEVRVQSIV VEFILTHVDQLFRGDSLSAGVDLESG WKSPLGARASGSSEDLMPTSLPYHL PSILQAGDGPPQIRPYHTIIEIAEHKR KGSLKVRKWRISIFNLGRSGHETKRK LPLRVEDREEKSSKGLRPAKSMDS LSAAAGASDEPEGLVGSSSSQPSL MPESLESNSMEGEQEPEAEAPGSA NSEPGTPRAGRS AVRALGSSRAERC AGVHISDPYNVNPLPHITSILSVPPNI ISNVSLVRLTRGLECPALQPRPSPAV GPLASDEKSEARSVPGPLDDSSPAA MTPALEDSLSQEVQDSFSFLEDLSS SEPEWVGVEEREVAKAEAAGAAGAA AFSLGEDDPGMGYLELLRVGPQVE EFSVEPPLDDLSDDTQYVLAPNCC SLDSAVSTPDVEEDYGEEVFLSAYD DLSPLLGPKPINWEGVGSLEEEAAG CGKQPPTQDEEEQACSETRQEKEAK PRSTSDNREEAEATPETEMEAGKAD AEGGEAERSQKVMDSFKEGSREELE AKEENSEGREVESIKETKDVEKIIGE PGKDEEREIGREEGAEGDDTPVDS DMDPEHVFQEDLVLEESWEVVHKKH EAEKGRESETKELRRKSDLKSREDQ GHSEDSGSPEEGDDRKEGVFSKEQ KSIDVETEVMRGVGDHLEEGALSEG PGVELLRVDSTEEINEQTSEMKAQAP LQPSEPEGMEAEGQLNPETCDLYSC PCGSAGGVGMRLASTLVQVRQVRSV PVVPPKPQFAKMP SAMCSKIHVAPA SPCPRPGRLDGTPGEKAWGSRASW RNGGSLSFDAVALARERQRTESQG VLRTQTCTGGGDYLSRTPPCSMIL AHSSRPLSCLERPPEGTEGSEPRSL SLPPRELHPVPLVAPQRQTYAFETQ TNHGKDEGV