

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|-------------|--------------|----------------------------------|--------------|---------------|---|----------|---|
| A7KAX9 | RHG32_HUMAN | Homo sapiens | Rho GTPase-activating protein 32 | 12.453809 | NaN | S706;S709;S732;S738;S852;S856;S892;S952;S1203;S1585 | 35132862 | METESESSTLGDDSVFWLESEVIIQV TDCEEEEREKFRMKSSVHSEED DFVPELHRNVHPRERPDWEETLSA MARGADVPEIPGDLTLKTCGSTASM KVKHVKKLPFTKGHFPKMAECAHF HYENVEFGSIQLSLSEEQNEVMKNG CESKELVYLVQIACQGKSWIVKRSYE DFRVLDKHLHLCIYDRRFSQLSELPR SDTLKDSPESVTQMLMAYLSRLSAIA GNKINCGPALTWMEIDNKGHLLV HEESSINTPAVGAAHVIKRYTARAPD ELTLEVGDIVSVIDMPPKVLSTWWR GKHGFQVGLFPGHCVELINQKVPQS VTNSVPKPVSKKHGKLITFLRTFMKS RPTKQKQKQKRGILKERVFGCDLGEH LLNSGFVQPVLQSCAFIERYGIVD GIYRLSGVASNIQRLRHEFDSEHVPD LTKEPYVQDIHSVGLCKLYFRELPN PLLYQLYEKFSDAVSAATDEERLIKI HDVIQQLPPPHYRTLEFLMRHLSLL ADYCSITNMHAKNLAIVWAPNLLRS KQIESACFSGTAAFMEVRIQSVVVEF ILNHVDVLFSGRISMAMQEGAASLS RPKSLVSSPSTKLLTLEEAQARTQA QVNSPIVTENKYIEVGEGPAALQKGF HTIIEFPLERKRPQNKMKKSPVGSW RSFFNLGKSSSVSKRKLQRNESEPS EMKAMALKGGRAEGTLRSKSEESL TSLHAVDGDGSKLFRPRRPRSSDAL SASFNGEMLGNRCNSYDNLPHDNE SEEGGLLHIPALMSPHSAEDVDLS PPDIGVASLDFDPMSFQCSPPKAESE CLESGASFLDSPGYSKDKPSANKKD AETGSSQCQTPGSTASSEPVSPQEK LSPFFTLDSPTEDKSSKPSSTFKV VYAFSPKIGRKLKSPSMSISEPISVT LPPRVSEVIGTVSNTTAQNASSSTW DKCVEERDATNRSPTQIVKMKTNET VAQEAYESEVQPLDQVAEEVELPG KEDQSVSSSQSKAVASGQTQTGAVT HDPPQDSVPVSSVSLIPPPPPKNVA RMLALALAESAQCASTQSLKRPGTS QAGYTNYGDIAVATTEDNLSSSYSAV ALDKAYFQTRPAEQFHLQNNAPG NCDHPLPETTATGDPHTSNTTESGE QHHQVDLTGNQPHQAYLSGDPEKA RITSVPLDSEKSDDHVSFPEDQSGK NSMPTVSFLDQDQSPRFYSGDQPP SYLGASVDKLLHHPLEFADKSPTPPN |

LPSDKIYPPSGSPEENTSTATMTYMT
TTPATAQMSTKEASWDVAEQPTTAD
FAAATLQRTHRTNRPLPPPSQRSA
EQPPVVGQVQAATNIGLNNSHKVQG
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RSVPISRTEVPPDDEPAYCPRPLYQY
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VHYRYSYSSSSSSSYSPDGALCDVD
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PEGEDRFYRRHPEAEMDRAHHHGG
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MGCSLPEHRAHQEASHRQFCESKN
GPPYPQGAGQLDYGSKGIPDTSEPV S
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SKEMERPWVRQPSAPEKHSRDCYK
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QNVERDPSVLYQYQPHGKRQSSVTV
VSQYDNLEDYHSLPQHQRGVFGGG
GMGTYVPPGFPHQSRTYATALGQG
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