

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q52LW3	RHG29_HUMAN	Homo sapiens	Rho GTPase-activating protein 29	6.445675	NaN	S171;S176;S179;S190;S499;S519;S552;S913;S949;S1019;S1144;S1146	28411811;34019948;28510447	MIAHKQKKTKKKRAWASGQLSTDIT TSEMGLKSLSSNSIFDPDYIKELVND IRKFSHMLLYLKEAIFSDCFKEVIHIR LEELLRVLKSIMNKHQNLNSVDLQN AAEMLTAKVKAVNFTEVNEENKND LFQEVFSSIETLAFTFGNLTNFLMG DVGNDSSLRLPVSRETKSFENVSVE SVDSSSEKGNFSPLELDNVLLKNTD SIELALSYAKTWSKYTKNIVSWVEKK LNLELESTRNMVLAEATRTNIGIQE FMPLQSLFTNALLNDIESSHLLQOTI AALQANKFVQPLLGRKNEMEKQRK EIKELWKQEQNKMLEAENALKKAK LLCMQRQDEYEKAKSSMFRAEEEH LSSSGGLAKNLNKQLEKKRRLEEEA LQKVEEANELYKVCVTNVEERRNDL ENTKREILAQLRTLVFQCDLTLKAVT VNLFHMQLQAASLADSLQSLCDS AKLYDPGQYSEFVKATNSTEEEKV DGNVNKHLNSSQPSGFGPANSLD VVRLPDSSNKIEEDRCSNSADITGPS FIRSWTFGMFSDSESTGGSSSESRL DSESISPGDFHRKLRPTPSSGTMSSA DDLDEREPPSPSETGPNLSTGTFKKT MSKAALTHKFRKLRSPKCRDCEGI VVFQVECEECLLVCHRKCLLENLVII CGHQKLPKIHFLFGAFTQVAKKEP DGIPFILKICASEIENRALCLQGIYRV CGNKIKTEKLCQALENGMHLVDISE FSSHDICDVLKLYLRQLPEPFILFRLY KEFIDLAKIQLHVNEEQETKNSLE DKKWPNMCIENRILLKSKDLLRQLP ASNFNSLHFLIVHLKRVVDHAEENK MNSKNLGVIFGPSLIRPRPTTAPITIS SLAEYSNQARLVEFLITYSQKIFDGS LQPQDVMCSIGVVDQGCFFPKPLLS EERDIERSMKSLFFSSKEDIHTSESE SKIFERATSFEESEKQNALGKCA CLSDKAQLLLDQEAESASQKIEDGK TPKPLSLKSDRSTNNVERHTPRTKIR PVSLPVDRLLLASPPNERNRNMG NVNLDKFCNPAFEGVNRKDAATT VCSKFNFGFDQQTLOKIQDKQYEQNS LTAKTTMIMPALQEKGVTTSLQISG DHSINATQPSKPYAEPVRSVREASER RSSDSYPLAPVRAPRTLQPHWTTF YKPHAPIISIRGNEEKPASPSAAVPPG TDHDPHGLVVKSMPPDPKASACPG QATGQPKEDSEELGLPDVNPQCQR PRLKRMQQFEDLEGEIPQFV