

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q3TBD2	HMHA1_MOUSE	Mus musculus	Rho GTPase-activating protein 45	12.769218	S875	S23;S72;S92;S98;S568;S577;S591;S618;S946;S1017;S1020;S1022	34418053;29187734	MFSRKKRELMKTSPISKKNRAGSPN PQSSSGELPRKDWTEAPGLEPPATS LSTVAKGTGTLKRPTSLSRHASAAGF PLSGTATWTLVVRGYRSPLSAASPAEL PTEGAFPDGVEDISTLLADVARFAEG LEKLKECVLQDDLLEARRPLAHECL GEALRVMRQVISRYPLLNTVETLTAA GTLIAKVKAFHYECNNESDKREFEK ALETIAVSFSCTVSEFLLGEVDSSTLL AVPPGDPSQSMENLYGAGTEGPPH NVEECEEGCLPPEEVDMLLQRCEG GVDAALQYAKDMARYMKDLISYLEK RTTLEMFAKGLQKVHNCRQSVT HEPHMPLLSIYSLALEQDLEFGHGM VQAAGTLQTQTFMQPLTLRRLHER RRKEIKESWHRAQRKLQEAEANLRK AKQGYKQRCEDHDKARLQVAKAEE EQQGTGPGAGTAASKALDKRRRLEE EAKNKAEEAMATYRTCVA DAKTQKQ ELEDTKVTALRQIQEVIRQSDQTIKS ATISYYQLMHMQTAPLPVNFQMLCE SSKLYDPGQQYASHVRQLQRGEEP VRYDFEPYVSNSWSPIMRTRKGSF NPGDASGPEAAGSPPEEGGTSEAAP NKDHRGGRGHQVHKS WPISISDTEV GLDTSSGDLKKFDR TSSSGTMSSSE ELGDQEAGLVASAFDSADLNGMDP ELPVAMPSPFRHVLGSKAARTHRL RKL RTPAKCRECNSYVYFQGAECE CCLACHKKCLET LAIQCGHKKLQGR LQ LFGQDFSQAALSTPDGVPFIVKCC VCEIERRALHTKGIYRVNGVKTRVEK LCQAFENGKELVELSQASPHDISNV LKLYLRQLPEPLISFRFYHELVGLAK DSLKAEAEAKAASRGRQGGSESEAA TLAMVGRLRELMQDLPAENRATLLY LLKHLRRIVEME QDNKMTPGNLGIV FGPTLLRPRPTDATVSLSSLDYDPHQ ARVIETLIVHYGLVFEEPEEAPGSQ EGASTQCGQLESAEGIVFPLQEEAE DGSRESHAASNDSDSELEDASDPLS SSDASALHRLSFLEQTEAGLEEGPQ SHSGSEEQLEGEDGAPGPWLCHFN TNQSNNTSRAPLPTMRLRGGQITGG TSQERQPQFV