

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9C091	GRB1L_HUMAN	Homo sapiens	GREB1-like protein	6.287639	NaN	NaN	34019948	<p>MGNSYAGQLKSARFEEALHNSIEAS LRCSSVVPPIFSQLYLDPDQHPFSS ADVKPKVEDLDKDLVNRYTQNGSLD FSNNLTVNEMEDDEDDEEMSDSNS PPIPYSQKPAPEGSCTTDGFCQAGKD LRLVSLCMEQIDIPAGFLLVGAKSPN LPEHILVCAVDKRFLPDDHGKNALL GFSGNCIGCGERGFYFTEFSNHIN LKLTTQPKKQKHLKYYLVRSSQGVL SKGPLICWKECRSRQSSASCHSIKPS SSVSSTVTPENGTNGYKSGFTQTD AANGNSSHGKGSASSSTPAHTGN YLSLSPRPSYASGDQATMFISGPPKKR HRGWYPGSPLPQPGLVVPVPTVRPL SRTEPLLSAPVPQTPLTGILQPRPIPA GETVIVPENLLSNSGVRPVILIGYGT PYFYGNVGDIVVSPLLVNCYKIPQLE NKDLEKLGLTGSQFLSVENMILLTIQ YLVRLGPDQVPLREEFEQIMLKAMQ EFTLRERALQIGAQCVPVSPGQLPW LARLIASVSQDLVHVVTQNSLAEGI SETLRTLSEMRHYQRLPDYVVVICAS KIRGNEFCVVVLGQHQSRAEAESML TTSEFLKEISYELITGKVSFLASHFKT TSLGDDLDKLEKMQRRGDSVVT FDGDLNECVSPQEAAMIPTQNLDL DNETFHIIYQPQLTVARKLLSQVCAIA DSGSQSLDLGHFSKVDIIIIVRSEVL VQQTQRIRQSGVLDLGLENGTA HQRAEKYVVRDNEIQTKFEVFMRR VKQNPYTLFVLVHDNSHVELTSVIS GSLHSEPSHGLADRVINCREVLEA FNLLVLQVSSFPYTLQTQQRISSEN EVHWIQLDTGEDVGCEEKLYFGLSE YSKSLQWGITSPLLRCDETFEKMVN TLLERYPRLHSMVVRCYLLIQYSEA LMALTTMASLRDHSTPETLSIMDDL ISSPGKNKSGRGHMLIIRVPSVQLA MLAKERLQEVKDKLGLQYRFEIILG NPATELSVATHFVARLKSWRGNEPE EWIPRTYQDLGLPCIVILTGKDPLG ETFPRSLKYCDLRLIDSSYLTRTALE QEVGLACCYVSKEVIRGPTVALDLSG KEQERAASVENDSDELLIDLERPQS NSSAVTGTSGSIMENGVSSSSTADK SQKQSLTPSFQSPATSLGLDEGVAS SAGAGAGETLKQECDSLGPQMASST TSKPSSSSSGPRTLPPWPGQPIRGCRG PQAALPPVILSKAAYSLLSQKSGK</p>

LPSSSSLLPHADVAWVSSLRPLLK
DMSSEEQSLYRQWTLARQHHADY
SNQLDPASGTRNFHPRLLLLTGPPQ
VGKTGSYLQFLRILFRMLIRLLEVDV
YDEEINTDHNESSEVSQSEGEWP
DIESFSKMPFDVSVHDPKYSLMSLV
YTEKLAGVKQEVIKESKVEEPRKRET
VSIMLTKYAAYNTFHHCEQCROYM
DFTSASQMSDSTLHAFTFSSSMLGE
EVQLYFIIPKSKESHFVFSKQGKHLE
SMRLPLVSDKNLNAVKSPIFTPSSGR
HEHGLLNLFHAMEGISHLHLLVVKE
YEMPLYRKYWPNHIMLVLPGMFNN
AGVGAARFLIKELSYHNLELERNRL
EELGIKRQCVWPFIVMMDDSCVLW
NIHSVQEPSSQPMVEGVSSKNVSLK
TVLQHIEATPKIVHYAILGIQKWSSKL
TSQSLKAPFSRCHVHDFILLNTDLTQ
NVQYDFNRYFCEDADFNLRTNSSGL
LICRFNNFSLMKKHVQVGGQRDFII
KPKIMVSESLAPILPLQYICAPDSEHT
LLAAPAQFLLEKFLQHASYKLFPKAI
HNFRSPVLAIDCYLNIGPEVAICYISS
RPHSSNVNCEGVFFSGLLLYLCDSF
VGADLKKFKFLKGATLCVICQDRSSL
RQTIVRLELEDEWQFRLRDEFQTAN
SSDDKPLYFLTGRHV