

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O70318	E41L2_MOUSE	Mus musculus	Band 4.1-like protein 2	24.308306	NaN	S7;S38;S86;S116;S201;S379;S395;S492;S543;S555;S561;S582;T606;S610;S630;S698;T745	34678516;34887587;22645316;28528544;25153642	MTTEVGSASEVKKGSDQAGADASKE KAKEVENEQTPVSEPEEEKGSQPGP PVERQSTPRLRKRKGDPSENRGISR FIPPWLKKQRSYNLVVAKDGGDKKE PTQADVEDQILGKEESLPEEESRAKG DAEEMAQRKHLEVQVEVREAKPAL KSSVETQPAEEVRKDKEETIQDTQE EKLEGGAAKRETKEVQTSELKAEVA SQKATKKTIVLAKVTLLDGTTEYSCD LEKRAKGQVLFDRVCEHLNLEKDY FGLLFQDHPEQKNWLDPAKEIKRQL KNLPWLFTFNVKFPDPSQLTEDIT RYFLCLQLRQDIASGRLPCSFVTHAL LGSYTLQAEHGDYDPEEYDSIDLGD FQFAPAHTKELEEKVSELHKTHRGL SPAQADSQFLENAKRLSMYGVDLH HAKDSEGVDIKLGCANGLLIYKDRL RINRFAWPKILKISYKRSNFYIKVRPA ELEQFESTIGFKLPNHRAAKRLWKV CVEHHTFYRLVSPEQPPKTKFRTLGS KFRYSGRTOAQTREASTLIDRPAPQF ERASSKRVSRLDGAPIGVVDQSPPG EGSVPGPGVISYTTIQDGRDSSKPT KATPLPAEGKKNTLRVDGDNIYVRH SNLMLEDLKAQEAILKHQASISEL KRNFMASSTPEPRPSEWEKRRVTPLP FQPQASSHETLNVVEEKKRAEVGKD ESVITEEMNGKEMSPGHGPGETRK VEPVAHKDSTLSSESSSSSESEED VGEYQPHHRVTEGTIREEQEEDDEE LEEEPGQAKVVEREAAPDAVPDR QAGASVLPVTEAQEHVVAQKLPGE KGAHGGTAEQDPREEAEDPHRVN GEVPHLDDLGLPEIICCSEPPVVKTE MVTISDASQRTEISTKEVPIVQTETKT ITYESPQIDGGAGGDSGVLLTAQTITS ESASTTTTTHTKTVKGGISETRIEKR IVITGDAALDHDQALAQAIAREAREQH PDMSVTRVVVHKETELAEEGEE