

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9WV92	E41L3_MOUSE	Mus musculus	Band 4.1-like protein 3	30.101893	S797	S96;S428;S451;S486;T495;T518;S543;T545;S547;T725;S802;S804;T923;S543;S543;S451;S525;S451	22517741;28528544;35822049;34887587;33300544;22645316;34678516	MTTESGSDSESKPDQEAEPQEAAGP QQQAGAQPGEPEAGGNGSLNGEKQ QPALEQFPEAAAHSSTPVKREIGDKD RDFAAAAAKQLEYQQFEDDKLSQRS SSSKLSRSPLKIVKRPKSMQCKVTLL DGSEYGCVDKRSRQVLFQDKVCE HLNLEKDYFGLTYRDAENQKNWL DPAKEIKKQIRSGAWHFSFNVKFYPP DPAQLSEDI TRY LCLQLRDDIVSGR LPCSFVTLALLGSYTVQSELGDYDPD ECGNDYISEFRFAPNHTKELEDKVIE LHKSHRGMTPAEAEMHFLENAKKL SMYGVDLHHAKDSEGVEIMLGVCA SGLLIYRDLRINRFAWPVKVLSISKYK RNNFYIKIRPGEFEQFESTIGFKLPN HRAAKRLWKVCVEHHTFFRLLLPEA PPKKFLTGLSKFRYSGRTQAQTRRAS ALIDRPAPYFERSSSKRYTMSRSLDG ASVSENHEIYMKDSVSAAEVGTGQY ATTKGISQTNLITTVTPEKKAEEERV EEEDRRKKAEEATPVTALRHEGKTD SERTDTAADGETSATESDQEEDAIEK AQDLDKTQDELMKHQTNISELKRTF LETSTETALTNEWKRLSTSPVRLAA RQEDAPMIEPLVPEETKQSSGEKLM DGSEILSLESARKPTEFIGGVSSTTQ SWVQKLETKTEPVEAEVESTPHQP LSTKVLQETILVEERHVMSVHASG DASHTARDEVDAEESTPTDRRHTGK GKEGSSVTEAAKEQRGEEVDQSAPE QEQPATVSHEEEQASTIRTSEGLEQ KSHFESSTVRVESTSVGSISPGGAKL EISTKEVPVVHTETKTITYESSQVDP GADLEPGVLMQAQTITSETTSTTTTT HITKTVKGGISETRIEKRIVITGDADI DHDQALAQAIKEAKEQHPDMSVTKV VVHKETEITPEDGED