

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q80TH2	ERBIN_MOUSE	Mus musculus	Erbin	19.869727	NaN	S440;S444;T483;T485;S595;S599;S600;S617;S712;S849;S854;S869;T914;T917;S928;T970;T1097;S1150;S1171;S1276;S1231;S1234	22645316;28528544	MTTKRSLFVRLVPCRCLRGEETVT TLDYSHCSLEQVPKEIFTFEKTLEEL YLDANQIEELPKQLFNCQSLHKLSL PDNDLTTLPASIANLINLRELDVSKN GIQEFPENIKNCKVLTIVEASVNPISK LPDGFSQLLNLTQLYLNDAFLFLPA NFGRLTKLQILELRENQLKMLPKTM NRLTQLERLDLGSNEFTEVPEVLEQ LSGLREFWMDGNRLTFIPGFISLR QLTYLDVSKNNIEMVEEGISTCENL QDFLLSSNSLQQLPETIGSLKNVTTL KIDENQLMYLPDSIGGLRSIEELDCS FNEIEALPSSIGQLTNMRTFAADHN YLQQLPPEIGNWKNITVFLHCNKL ETLPEEMGDMQKLKVINLSDNRLK NLPFSFTKLQQLTAMWLSDNQSKPL IPLQKETDTETQKMVLTYNMFPPQP RTEDVMFISDNESFNPALWEEQRK QRAQVAFECDEKDEREAPPREGNL KRYPTYPDELKNMVKTIVQTIHRL KDEETNEESGRDLKQHEDQQVVNK DKCVKTSESTTTKSKLDEREKYMNS VQKMSEPEAETNGGNLPTASMKL SGNLKHIVNHDDVFEESEELSSDEE MKMAEMRPPLIESSINQPKVVALSN NKKDDAKDADSLSDVTHNSNQNN SNCSSPSRMSDSVSLNTDSSQDTSL CSPVKQTPVDSNSKVRQEDENFNSL LQNGVNLNNSPEEKFKINDKKDFKL PEYDLNIEEQVLVIEKDIDSKATSDD SRQLDHINMNINKLVTNNIFQPEVM ERSKMQDIVLGTGFLSIHPKNEAEHI ENGAKFPNLESINKVNGLCDTAPS PGRVEPQKASSADVGISKSTEDLSP QRSGPTGAVVKSHSITNMETGGLKI YDILGDDGPQPPSAAVKIASAVDGKN IVRSKATLLYDQPLQVFTAASSSSE LLSGTKAVFKFDSNHNPEEPDIIRAA TVSGPQSTPHLYGPPQYNVQYSGSA TVKDTLWHPKQNPQIDPVSFPPQRL PRSESAENHSYAKHSANMNFNSHN NVRANTGYHLQQLAPARHGEMWA ISPNDRLVPAVTRTTIQRQSSVSSTA SVNLGDPTRRTEGDYLSYRELHSMG RTPVMSGSRPLSARAYSIDGPNTS RPQSARPSINEIPERTMSVSDFNYSR TSPSKRPNTRVGSEHSLDPPGKSK VPHDWREQVLRHIEAKKLEKHPQTS SPGECCQDDRFMSEEQNHPGALS

