

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8CC35	SYNPO_MOUSE	Mus musculus	Synaptopodin	43.692727	S393;S397;S507;T551;T752;T753	S134;S202;S222;S258;S488;S512;T547;S567;S672;S689;S740;S744;S765;T769;S833;S854;S690;S789;S803;S819;S824;S852;S869;S892	33300544;26192747;34678516;22517741;35822049;22645316;16452088	MLGAHFPPPLGASEGRAAPCTFQIP DGSYRCLALEAEESSEDGLQGEVR LVDLEEEGTSQSRANHGTPPLSRAP AIIQPSSCHREARGGFQSRDRPSHD WDVVQARKVMTASGSSSPVPRVAQ KPALGRSTSFTENDLKEAKARSQQA AQLTTPSSNSRGVQLFNRRRQRVN EFTLESRGQRSPKLNQEALQTGRPL SPIGHAPGPSVKPTSPSKPGSPKHP PQSPSRGVAGHIMEGYSEEASLLRH LEKVASEEEEEVPLVVYLKENAALLTA NGLHLSQSRETQOSSPNPPDTEVPS PAADINQNPPSNATLTTLASSSHHS OPTADINQNPPAAITVPQNSSQAQ CSPNGTLDKPGTLCADDGQSPVPA EEVRSILLIDKVSAPPSAATSFSREA TPLSSSGPPAADLMSSSLIDMQPST LVAPAEQEVPGHVAVTTPTKVYSEV HLTLAKPASVVNRTARPFGIQSPGTS QIEQSPMMGRRQFGEKAWAPPASS MADRSPQQRHIMSRSPMVERRLL GQRSPVLERRPLGNFTPPPTYAETLS TAPVASRVRSPPSYSTLYPSSDPKPS HLKGQVAPANKTGILEESMARRGRS KSMFTFVEKPKVTPNPDLLDLVQTA DEKRRQRDHGEVGMEEEPFALGAE ASNFAQEPIARDRASPAAAEAVPE WASCLKSPRIQAKPKPKPNQNLSEA SGKGAELYARRQSRMEKYVIESSGH AELARCPSTMSLPSSWKYTTNAPG GFRVASLSPARTPPASLYHGYPENG VLRPEPTKQOPYQMRPSLYALSPVK EPAKASSRATSSRTPSRTVSPRAASP AKPSSLDLVPNLPRAGLPPSPALPRP SRSSPGLYTAPVQDSLQPTAVSPTYS SDISPVSPRAWSPRAKQAPRPSFST RNAGIEAQVWKPSFCFK

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