

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8BWB1	SYP2L_MOUSE	Mus musculus	Synaptopodin 2-like protein	15.565522	S89	S105;S108;T138;S140;S163;S175;S177;S342;S347;S371;S378;S381;S667;S675;T702;T710;S787;S789;T791;S890;T891;T897	36288343;36852467	MGAEEEVQVTLAGGAPWGFRLQGG TEQRKPLQIRRRSQAGRAGLRERDQ LLAINGVSCTNFNSHASAMTLIDASGR QLVLTVRRVTDEGSVRSPSPGELQV LSPLSPLSPEPPGAPVSQALQPTSLR SPPDSEAYYGETDSVDGDPATQEKP RRTRRRGPARPSLPGAPPDEVYLSDS PAEPAPVKTGSPSQGDSRVSSPSWE EGAALQPPPAEALLPHGPLRPGPH LIPMVGVPVPHVAEDLTTTTYTQKAKQ AKLQRAESLQEKSVKEARTKCRITIAS LLTAAPNPHSKGVLMFKRRRQRAKK YTLVSFGAAAGTGTEEDGIPPTSES ELDEETFSDARSLTNQSDWDSPLYD MELARAGLGTAESQNSGLGGQLSEV SGRGVQLFEQQRQRVASSSQELAQV GPAAMLNGQSLQSPRAQSAPPEAA VLPLSPLSVPVAVSPTPFFPDGGAPIPA PSIFNRSARPFTPLGQQRSGTTSVI FRPLAPKKVNEGLGSTSPAPSPFAAP PQGPTPLPSFTTVVPSHTPVSGASS TQRSSGPVTATSSLYIPAPSRPVTPG GAPEPPTPPSAAAMTSTASIFLSTPL RNSARPEAPGPAVPEPASVREQRISV PAARTGILQEARRRGTRKQMFPRGK EETKNSPPELLSLVQNLDEKPRAG GAESGPEEDALSLGAEACNFMQPLG GRSYKTLQVSPKTPPPMAPKTPPPT TPKTPPPVAPKPGSRGLLDGLVNGST PMVGIPEPPRLQGRGGELFAKRQSR ADRYVVEATSGSSLNPGLRPRSPSPT PSLPPSWKYSNIRAPPPIAYNPLLSP FFPQAARTLPNKAQSQGPRVTPKQG IKALDFMRHQPYQLKTAMFCFDEGS STPGPTSGPPKTARVQEIRRFSTPAP QPTAEPLAPTVLVPRAATTLDEPIWR AELASTPVPNPDHQESLRSFAAAPSS CGFQVARPRFSATRTGLQAHVWRPG AGHQ