

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8N3V7	SYNPO_HUMAN	Homo sapiens	Synaptopodin	26.066898	T406;T409;T410;T411;S412;T413;S415;S448;T560;T564;T566;S570	S140;S207;S263;S501;S525;T560;S580;S685;S702;T746;S754;S758;S779;T783;S833;S854;S738;S784;S804;S812;S826;S854;S871;S894	30379171;20068230;30059200;31492838;28657654;29237092;30620550	MLGPHLPPPPLAPSEGRPTPCAFQIP DGSYRCLALEAEESSGEEGLQGEVG PTDLEEDGVSRSRSGDDSAACRVTQGT PQLPKALGIQPPSCSREEQGASQHD DRASQDWDVVKAGQMMTASPSGP GPRVAQKPALGRSTSLTEKDLKEAK ARSQQIAAQLTTPSSNSRGVQLFN RRRQRVNEFTLESHGQRGQKPSQE SLRVLPSLPGHAPGLSLSSTSLPEP GPPRHPSQSPDRGVPGHSMEGYSE EASLLRHLEKVAEEEEVPLVVYLKE NAALLTANGLHLSQNREAQQSSPAP PPAEVHSPAADVQNQLASPSATLTT PTSNSSHNPATDVNQNPPATVVVQ SLPLSSIQQNSSEALPSNGTGPASK PSTLCADGQPQAPAEVRCSTLLIDK VSTPATTSTFSREATLIPSRPPASD FMSSLLIDIQNTLVVSADQEMSG RAAATTPTKVYSEVHFTLAKPPSVVN RTARPFGIQAPGGTSMERSPMLER RHFGEKAPAPQPPSLPDRSPRQRHI MSRSPMVERRMMGQRSPASERRPL GNFTAPPTYTETLSTAPLASWVRSP SYSVLYPSSDPKSSHLLKQAVPASKT GILEESMARRGSRKSMFTFVEKPKV TPNPDLLDLVQTADKRRQRDQGEV GVVEEPPFALGAEASNFQQEPAPRDR ASPAAAEVVPWASCLKSPRIQAKP KPKPNQNLSEASGKGAELYARRQSR MEKYVIESSHTPELARCPSPMSLP SSWKYPTNAPGAFRVASRSPARTPP ASLYHGYPENGVLRPEPTKQPPYQL RPSLFVLSPIKEPAKVSFRAASPAKPS SLDLVNLPGKALPPSPALPRPSRSS PGLYTSFGQDSLQPTAVSPPYGGDIS PVSPSRAWSPRAKQAPRPSFSTRNA GIEAQVWKPSFCFK