

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membra
Q91YE8	SYNP2_MOUSE	Mus musculus	Synaptopodin-2	33.354824	S990;S991	S300;S319;S320;T323;S540;S541;S543;S546;S555;S596;T602;S603;T614;S618;S697;S719;T744;S767;S771;S895;S899;S903;S1008;S1050	30016717;39927985	MGTGDFICISMTGGAPWGFRLQGG KEEQQLQVAKIRSQSKASGSLRE GDEVVSINGNPCADLTYPEVIKLE GITDSLHLLVKRPSSGTSETLDSESE TTNHQHLLTHEGPMESTTLQIQQATE TQSEDFFLAPVQTKVPLTEDQSNAW GYAECPKEEQAPPMLGSOEGHLVEE VILRQKAEAGQPGHVVELQLSLSKE RHQCTSGPIVTLQGNKSTSPDPDW SSQLERTVHINSIPAPEKADTSLTSST SSGRELRVIQGRDPGGAGLPQVEVIL DCSDRLKAEECRLQTGRGCVASPVE GGRSEAPPSPVSVFAVSSEGTEHGED QRSKQSRPHKHRARHARLRSE SLSEKQVKEAKSKCKSIALLLTDAPN PNSKGVLMFKRRRRRARKYTLVSYG TGELEEEEEEDQEAGDKDEISEV AFLGTSESEVDEELLSDVDDNTQVV NFDWDSGLVDIEKRLNRGDKMEML PDTTGKALMFARRRERMEQFTAQ NEEEKTGGMAGGGPDALQTDGLRT MTSYQRKEESVRMQSSVSESSFQM GRSLASVPQQNGFSGVSETAGAQR MFPMNRTAKPFLGSMNQPAAPFSP TRSVTSPISDFPAPPYSAVSPPEAF SRGVSSPVAGPAQPPWPQAPWSQ PAFYDSSEQIASRDERIAVPAKRTGIL QEAARRGTTKPMFTFKETKVSNPPE LLSLLQNAEGKRGTTGGDSGPEEDYL SLGAEACNFMQSSAKQKTPPPVAPK PAVKSPSSQPVAVSPVWSPGVAPA QRPAFSTSNPPNPQVTAVSSIKIAQ PAAPPARPASALNLAGPFGKQAVV VSHNYTPKPSAPTPLVNAAPAGAGG PSNELPGMSGKGAQLFAKRQSRME KYVVDSDTVQAHTVRAQSPTPSLPAS WKYSSNVRAPPVAYNPIHSPSYPLA AIKSQPPGAQASKTSKKKGGKPLNLT DVMKHQPYQLNASLFTFQPPDSKD GLPKSTVKVSSAPAMKQALPPROA NVGSPTNAQASSVYVPAYTSQPNF FAAEATSPVSAASPVPVSVPTSPKQES TTSYFVAPRPKFSAKKSGVTVQVW KPSVVEE	True	False	3.601	5.0	0.938	False	False	False