	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrac region
0	Q91YE8	SYNP2_MOUSE	Mus musculus	Synaptopodin-2	26.088944	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	MGTGDFICISMTGGAPWGFRLQGG KEEQQPLQVAKIRSQSKASGSGLRE GDEVVSINGNPCADLTYPEVIKLME GITDSLHLLVKRPSSGTSETLDSESE TTNHQHLTHEGPMESTTLQIQQATE TQSEDFFLAPVQTKVPLTEDQSNAW GYAECPKEEQAPPMLGSQEGHLVEE VILRQKAEAGQPGHVVELQLSLSKE RHQCTSGPIVTLQGNDKSTSPDPDW SSQLERTVHINSIPAPEKADTSLTSST SSGRELRVIQGRDPGGAGLPQVEVIL DCSDRLKAEECRLQTGRGCVASPVE GGRSEAPPSLVSFAVSSEGTEHGED QRSGKDQSRPHKHRARHARLRRSE SLSEKQVKEAKSKCKSIALLLTDAPN PNSKGVLMFKKRRRRARKYTLVSYG TGELEREEEEEEDQEAGDKDEISEV AFLGTSESEVDEELLSDVDDNTQVV NFDWDSGLVDIEKRLNRGDKMEML PDTTGKGALMFAKRRERMEQFTAQ NEEEKTGGMAGGGPDALQTDGLRT MTSYQRKEESVRMQSSVSESSFQM GRSLASVPQQNGFSGVSETAGAQR MFPMNRTAKPFLGSMNQPAAPFSP TRSVTSPISDFPAPPPYSAVSPPPEAF SRGVSSPVAGPAQPPPWPQPAPWSQ PAFYDSSEQIASRDERIAVPAKRTGIL QEAKRRGTTKPMFTFKETKVSPNPE LLSLLQNAEGKRGTGGDSGPEEDYL SLGAEACNFMQSSAKQKTPPPVAPK PAVKSPSSSQPVAPVSPVWSPGVAPA QRPAFSTSNPPNPPQVTAVSSIKIAQ PAAPPARPASALNLAGPFKGPQAVV VSHNYTPKPSAPTPLVNAAPAGAGG PSNELPGMSGKGAQLFAKRQSRME KYVVDSDTVQAHTVRAQSPTPSLPAS WKYSSNVRAPPPVAYNPIHSPSYPLA AIKSQPPGAQASKTSKKKGKKPLNTL DVMKHQPYQLNASLFTFQPPDSKD GLPQKSTVKVSSAPAMKQALPPRQA NVGSPTNAQASSVYSVPAYTSQPNF FAAEATSPVSASPVPVSVPTSPKQES TSTSYFVAPRPKFSAKKSGVTVQVW KPSVVEE	None	None	None	None	None	None	None	None	None