

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q24546	SYN_DROME	Drosophila melanogaster	Synapsin	25.496522	NaN	S539	33925313	MKRGFSSGDLSEVDDVDPNSLPPA ARPIQDQPTKPPVAGGPPNMPPPPA PGQPAGAAPELSLSFGAGKTPATAAP APPRGVSAPTSPAKSRESLLQRVQSL TGAARDQGASILGAAVQSATQRAPA FSKDKYFTLLVLDQNTDWSKYFRG RRLHGDFDIRVEQAEFRDITVSSAD TGPVVTMAAYRSCTRVARSRPFDV LIRQPPRDGSSDYRSTILGLKYGGVP SINSLHSIQFQDKPWVFSHLLQLQ RRLGRDGFPLIEQTFFPNRDLFQFT KFPSVLKAGHCHGGVATARLENQSA LQDAAGLVSGAGNDSHCYCTIEPYID AKFSVHIQKIGNNYKAFMRKSITGN WKTNQGSAMLEQITLTKYKSWVD EISELFGGMEVCGLSVVAKDGREYI ISACDSTFALIGDTQEEDRRQIADLV SGRMQNVCRPSMAQTGPGKLPSSRS SVSSRAESPTDEGVAPTPPLPAGPRP APMGGPPPPIPERTSPAVGSIGRLSSR SSISEVPEEPSGSPSTVGGVRRDSQ TSQSSTISSSVSRAGQRPPQTQNSVV EDAEDTMKNLRKTFAGIFGDMXEIA NKKRGRTASETSSGSGPGSVPSAG PGSGFSSFLGKQFSFAGKGEVIST QPTQRPSEPPAIPTTASSAVRPES VSVSDSRNTDILTERAGAGYQPVTN YEQQERVNPFDKPSKSGSAASIHT SSSSSSSSSSSRINRNGNAIQSPPP PAGPPPPPTNVTAVGSNANSSSGY RNSFSSSLSKDKTSYGNYSSTTSVET ITRMDTNTTNGATATEAGEASGVTA ITNISNSDGIVAPTTGTTTTSVTTNDW RSAIGMRSASVYSAPAAVTTVLPGDT SGYDSNSIASQGEGLNPNPSDLPSYT RPSYSRSESNAKHSDDLDFVFGDSKT TPASYGNGKYTRAAGSISDADMIFGG PPSNYKTRDFGASKSMSMTSGGVGS GNGSGSLGGYKIYDSIQNAAFSDFS DSGSMSSISGSHTKRWSASKEEDEL DLK	None	None	None	None	None	None	None		

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