| | UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochndrion | er re |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| **0** | P09951 | SYN1_RAT | Rattus norvegicus | Synapsin-1 | 53.033506 | S55;T56;T87;S96;S103;S261;S430;S516;T524;T562;S576 | S9;S39;S62;S67;T312;S425;S430;S432;T434;S436;S549;S551;S566;S566;S603;S662;S664;S682 | 12438562;29808487;38843836;20657584;18683930;24280219;10386995 | MNYLRRRLSDSNFMANLPNGYMTD<br>LQRPQPPPPPPSAASPGATPGSAAAS<br>AERASTAAPVASPAAPSPGSSGGGGF<br>FSSLSNAVKQTTAAAAATFSEQVGG<br>GSGGAGRGGAAARVLLVIDEPHTDW<br>AKYFKGKKIHGEIDIKVEQAEFSDLN<br>LVAHANGGFSVDMEVLRNGVKVVR<br>SLKPDFVLIRQHAFSMARNGDYRSL<br>VIGLQYAGIPSVNSLHSVYNFCDKP<br>WVFAQMVRLHKKLGTEEFPLIDQTF<br>YPNHKEMLSSTTYPVVVKMGHAHS<br>GMGKVKVDNQHDFQDIASVVALTK<br>TYATAEPFIDAKYDVRVQKIGQNYKA<br>YMRTSVSGNWKTNTGSAMLEQIAM<br>SDRYKLWVDTCSEIFGGLDICAVEAL<br>HGKDGRDHIIEVVGSSMPLIGDHQD<br>EDKQLIVELVVNKMTQALPRQRDAS<br>PGRGSHSQTPSPGALPLGRQTSQQP<br>AGPPAQQRPPPQGGPPQPGPGPQRQ<br>GPPLQQRPPPQGQQHLSGLGPPAGS<br>PLPQRLPSPTAAPQQSASQATPMTQ<br>GQGRQSRPVAGGPGAPPAARPPASP<br>SPQRQAGPPQATRQASISGPAPPKVS<br>GASPGGQQRQGPPQKPPGPAGPIRQ<br>ASQAGPGPRTGPPTTQQPRPSGPGP<br>AGRPTKPQLAQKPSQDVPPPIIAAAG<br>GPPHPQLNKSQSLTNAFNLPEPAPP<br>RPSLSQDEVKAETIRSLRKSFASLFS<br>D | True | False | 3.055 | 4.654 | 2.395 | 2.( |