	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	extracellular region
0	O75182	SIN3B_HUMAN	Homo sapiens	Paired amphipathic helix protein Sin3b	18.379595	NaN	S709;S712	30379171	MAHAGGGSGGSGAGGPAGRGLSGA RWGRSGSAGHEKLPVHVEDALTYLD QVKIRFGSDPATYNGFLEIMKEFKSQ SIDTPGVIRRVSQLFHEHPDLIVGFN AFLPLGYRIDIPKNGKLNIQSPLTSQE NSHNHGDGAEDFKQQVPYKEDKPQ VPLESDSVEFNNAISYVNKIKTRFLD HPEIYRSFLEILHTYQKEQLNTRGRP FRGMSEEEVFTEVANLFRGQEDLLS EFGQFLPEAKRSLFTGNGPCEMHSV QKNEHDKTPEHSRKRSRPSLLRPVS APAKKKMKLRGTKDLSIAAVGKYGT LQEFSFFDKVRRVLKSQEVYENFLR CIALFNQELVSGSELLQLVSPFLGKF PELFAQFKSFLGVKELSFAPPMSDR SGDGISREIDYASCKRIGSSYRALPKT YQQPKCSGRTAICKELDHWTLLQGS WTDDYCMSKFKNTCWIPGYSAGVL NDTWVSFPSWSEDSTFVSSKKTPYE EQLHRCEDERFELDVVLETNLATIR VLESVQKKLSRMAPEDQEKFRLDDS LGGTSEVIQRRAIYRIYGDKAPEIIESL KKNPVTAVPVVLKRLKAKEEEWREA QQGFNKIWREQYEKAYLKSLDHQAV NFKQNDTKALRSKSLLNEIESVYDE HQEQHSEGRSAPSSEPHLIFVYEDR QILEDAAALISYYVKRQPAIQKEDQG TIHQLLHQFVPSLFFSQQLDLGASEE SADEDRDSPQGQTTDPSERKKPAPG PHSSPPEEKGAFGDAPATEQPPLPPP APHKPLDDVYSLFFANNNWYFFLRL HQTLCSRLLKIYRQAQKQLLEYRTEK EREKLLCEGRREKGSDPAMELRLKQ PSEVELEEYYPAFLDMVRSLLEGSID PTQYEDTLREMFTIHAYVGFTMDKL VQNIARQLHHLVSDDVCLKVVELYL NEKKRGAAGGNLSSRCVRAARETSY QWKAERCMADENCFKVMFLQRKG QVIMTIELLDTEEAQTEDPVEVQHLA RYVEQYVGTEGASSSPTEGFLLKPVF LQRNLKKFRRRWQSEQARALRGEA RSSWKRLVGVESACDVDCRFKLSTH KMVFIVNSEDYMYRRGTLCRAKQVQ PLVLLRHHQHFEEWHSRWLEDNVT VEAASLVQDWLMGEEDEDMVPCKT LCETVHVHGLPVTRYRVQYSRRPAS P	None	None	None	None	None	None	None	None	None