

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
Q921M3	SF3B3_MOUSE	Mus musculus	Splicing factor 3B subunit 3	25.161187	NaN	S156;T1200	34887587	MFLYNLTLQRATGISFAIHGNFSGTK QQEIVVSRGKILELLRPDPNTGKVHT LLTVEVFGVIRSLMAFRLTGGTKDYI VVGSDSGRIVILEYQPSKNMFEKIHQ ETFGKSGCRRIVPGQFLAVDPKGRA VMISAIEKQKLVYILNRDAAARLTISS PLEAHKANTLVYHVVGVDVGFENP MFACLEMDYEADNDPTGEEAANT QQTLTFFYELDLGLNHVVRKYSEPLE EHGNFLITVPGGSDGPGSLICSENY ITYKNFGDQPDIRCPPIRRRNDLDDP ERGMIFVCSATHKTKSMFFFLAQTE QGDIFKITLETDEDMVTEIRLKYFDT VPVAAAMCVLKTGFLVASEFGNHY LYQIAHLGDDDEEPEFSSAMPLEEG DTFFFQPRPLKNLVLVDELDSLSPIL FCQIADLANEDTPQLYVACGRGPRS SLRVLRHGLEVSEMAVSELPGNPNA VWTVRRHIEDEFDAYIVSVFNATLV LSIGETVEEVTDSGFLGTTPTLSCSLL GDDALVQVYPDGIRHIRADKRVNEW KTPGKKTIVKCAVNQRQVIALTGGE LVYFEMDPSGQLNEYTERKEMSAD VVCMSLANVPPGEQRSRFLAVGLVD NTVRIISLDPSDCLQPLSMQALPAQP ESLCIVEMGGTEKQDELGERGSIGF LYLNIGLQNGVLLRVTVDPTGDLSD TRTRYLGSRPVKLFRVRMQGQEAVL AMSSRSWLSYSYQSRFHLTPLSYET LEFASGFASEQCPEGIVAISTNTRLRIL ALEKLGAVFNQVAFPLQYTPRKFSVIH PESNNLIIIEDHNAYTEATKAQRKQ QMAEEMVEAAGEDERELAAEMAAA FLNENLPESIFGAPKAGNGQWASVI RVMNPIQGNLTLVQLEQNEAAFSV AVCRFSNTGEDWYVLVGVAKDLILS PRSVAGGFVYTYKLVNNGEKLEFLH KTPVEEVPAAIAPFQGRVLIGVGKLL RVYDLGKKLLRKCENKHIANYSISGI QTIGHRVIVSDVQESFIWVRYKRNE NQLIIFADDTYPRWVTASLLDYDTV AGADKFGNICVVRLPPNTNDEVED PTGNKALWDRLLNGASQKAEVIM NYHVGETVLSLQKTTLIPGGSESLVY TTLSGGIGILVPFTSHEDHDFFOHVE MHLRSEHPPLCGRDHLSFRSYFPV KNVIDGDLCEQFNSMEPNKQKNVS EELDRTPPEVSKKLEDIRTRYAF	True	False	1.456	4.456	1.313	1.118	0.805	3.211	1.296

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