

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	extra regic
Q15393	SF3B3_HUMAN	Homo sapiens	Splicing factor 3B subunit 3	21.922984	S133;T237	S156;T1200	23301498;37217939;33214551;16408927;22121020;40596516;32522876;38665916;34725712;32119511;40914422;30620550;35083852;34019948;28510447;29351928	MFLYNLTLQRATGISFAIHGNFSGTK QOEIVVSRGKILELLRPDPNTGKVHT LLTVEVFGVIRSLMAFRLTGGTKDYI VVGSDSGRIVILEYQPSKNMFEKIHQ ETFGKSGCRRIVPGQFLAVDPKGRA VMISAIEKQKLVYILNRDAAARLTISS PLEAHKANTLVYHVVGVDVGFENP MFACLEMDYEEADNDPTGEEAAANT QQTLTIFYELDLGLNHVVRKYSEPLE EHGNFLITVPGGSDGPGSVLICSENY ITYKNFGDQPDIRCIPIRRRNDLDDP ERGMIFVCSATHKTKSMFFFLAQTE QGDIFKITLETDEDMVTEIRLKYFDT VPVAAAMCVLKTGFLFVASEFGNHY LYQIAHLGDDDEEPEFSSAMPLEEG DTFFFQPRPLKNLVLVDELDSLSPIL FCQIADLANEDTPQLYVACGRGPRS SLRVLRHGLEVSEMAVSELPGNPNA VWTVRRHIEDEFDAYIIVSFVNATLV LSIGETVEEVTDSGFLGTTPTLSCSLL GDDALVQVYPDGIRHIRADKRVNEW KTPGKKTIVKCAVNQRQVVIALTGGE LVYFEMDPSGQLNEYTERKEMSAD VVCMSLANVPPGEQRSRFLAVGLVD NTVRIISLDPSDCLQPLSMQALPAQP ESLCIVEMGGTEKQDELGERGSIGF LYLNIGLQNGVLLRTVLDPVTGDLS TRTRYLGSRPVLFVRVMQGEAVL AMSSRSWLSYSYQSRFHLLTPLYET LEFASGFASEQCPEGIVAISTNTRLRIL ALEKLGAVFNQVAFPLQYTPRKFSVIH PESNNLIHIEDHNAYTEATKAQRKQ QMAEEMVEAAGEDERELAAEMAAA FLNENLPESIFGAPKAGNGQWASVI RVMNPIQGNLTLDLVQLEQNEAFSV AVCRFSNTGEDWYVLVGVAKDLILN PRSVAGGFVYTYKLVNNGEKLEFLH KTPVEEVPAAIAPFQGRVLIGVGKLL RVYDLGKKLLRKCCENKHIANYSIGI QTIGHRVIVSDVQESFIWVRYKRNE NQLIIFADDTYPRWVTTASLLDYDTV AGADKFGNICVVRLPPNTNDEVDED PTGNKALWDRLLNGASQKAEVIM NYHVGETVLSLQKTTLIPGGSSESLVY TTLGGGIGILVPFTSHEDHDFQHVE MHLRSEHPPLCGRDHLSFRSYFFPV KNVIDGDLCEQFNSMEPNKQKNVS EELDRTPPEVSKKLEDIRTRYAF	True	False	1.419	5.0	1.364	1.268	1.018	3.23	1.334