

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q8JZQ9	EIF3B_MOUSE	Mus musculus	Eukaryotic translation initiation factor 3 subunit B	35.528647	NaN	T31;S37;S40;S68;T72;T74;S75;S79;S84;S88;S90;S111;S120;S123;S141;S143;S153;S228	34887587;37453647;21540332;36288343;39627609;37507081	MQDAENVAVPEAAEERAEPARQQP ASESPPTDEAAGSGGSEVGTEDAE EDAEAGPEPEVRAPAAQSEETAT SPAASPTPQSAERSPSQEPSAPGKAE AVGEQARGHPSAGAEIEGGSDGSAA EAEPRALENGEADEPSFSDPEDFVD DVSEELLGDVLDKDRPQEAADGIDSVI VVDNVPOVGFDRLEKLNVIHKIFS KFGKIINDYYPEEDGKTKGYIFLEYAS PAHAVDAVKNADGYKLDKQHTFRV NLFTDFDKYMTISDEWDIPEKQPFK DLGNLRYWLEEAECRDQYSVIFESG DRTSIFWNDVKDPVSIEERARWTET YVRWSPKGTYLATFHQRGIALWGGD KFKQIQRFHQGVQLIDFSPCERYLV TFSPMLDTQDDPQAIHWDLTGHHK RGFHCESSAHWPIFKWSDHGKFFA RMTLDTLSIYETPSMGLLDKKSLSKIS GIKDFSWSPGNNIAFWVPEDKDIPA RVTLMLQPTRQEIIRVNLFNVDCK LHWQKNGDYLCVKVDRTPKGTQGV VTNFEIFRMREKQVPVDDVEMKETII AFAWEPNGSKFAVLHGEAPRISVSF YHVKSNGKIELKMFDDKQANTIFW SPQGGFVVLAGLRSMNGALAFVDT DCTVMNIAEHYMASDVEWDPDTPG VVTSVSWWSHKVDNAYWLWTFQG RLQKNNKDRFCQLLWRPRPPTLLS ODQIKQIKKDLKYSKIFEQKDRLSQ SKASKELVERRRTMMEDFRQYRKM AQELYMKQKNERLELRGGVDTDEL DSNVDDWEEETIEFFVTEEVIPLGSQ E	False	False	3.388	3.765	1.613	1.61	0.792