

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
P55884	EIF3B_HUMAN	Homo sapiens	Eukaryotic translation initiation factor 3 subunit B	14.49512	S743	S78;S81;S83;S85;S119;S125;S152;S154;S164;S239	37217939;16408927;38665916;33214551;23301498;35138101;34019948;30397120;26853435;34725712;35008409;33465208;28510447	MQDAENVVPEAAEFERAEPGQQP AAEPPPAEGLLRPAGPGAPEAAGTEA SSEEVGIAEAGPESEVRTEPAAEAEA ASGPSESPSPAEEELPGSHAEPVP AQGEAPGEQARDERSDSRAQAVSED AGGNEGRAAAEAPRALENGDADEPS FSDPEDFVDDVSEELLGDLVKDRP QEADGIDSVIVVDNVPQVGPDRLEK LKNVIHKIFSKFGKITNDFYPEEDGK TKGYIFLEYASPAHAVDAVKNADGYK LDKQHTFRVNLFTDFDKYMTISDEW DIPEKQPFKDLGNLRYWLEEAECRD QYSVIFESGDRTSIFWNDVKDPVSIE ERARWTEYVVRWSPKGTYLATFHQR GIALWGGEKFKQIRFESHQVQLID FSPCERYLVTFSPMLDQDDPQAIH WDILTGHKKRGFHCESAHWPFK WHDGKFFARMTLDTLSIYETPSMG LLDKKSLKISGIKDFSWSPGGNIAF WVPEDKDIPARVTLMQLPTRQEI RNLFNVDCKLHWQKNGDYLCVKV DRTPKGTQGVVTNFEIFRMREKQVP VDVVMKETIAFAWEPNGSKFAVL HGEAPRISVSFYHVKNNGKIELIKMF DKQANTIFWSPQGQFVVLGLRS MNGALAFVDTSDCTVMNIAEHYMA SDVEWDPTGRYVTVSWSWSHKVD NAYWLWTFQGRLLQKNNKDRFCQL LWRPRPPTLLSQEQIKKIDLKKYS KIFEQKDRLSQSKASKELVERRRTM MEDFRKYRKMAQELYMEQKNERLE LRGGVDTDELDSNVDDWEEETIEFF VTEIIPLGNQE	True	False	4.784	4.236	1.699	1.641	0.868