

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q14152	EIF3A_HUMAN	Homo sapiens	Eukaryotic translation initiation factor 3 subunit A	16.869789	S37;S1058	S492;S584;S881;S882;S895;S949;S1028;S1188;S1198;S1262;S1336;S1364	33214551;40596516;26853435;38665916;34725712;32119511;40914422;34019948;35138101;33465208	MPAYFORPENALKRANEFLEVGKK QPALDVLVDVMSKSKKHRWQKIHE PIMLKYLELCLVDRKSHLAKEGLYQY KNICQQVNIKSLEDVVRAYLKMMAEE KTEAAKEESQOMVLDIEDLDNIQTP ESVLLSAVSGEDTQDRTRDLLLTPW VKFLWESYRQCLDLLRNNSRVERLY HDIAQQAFKFCLOYTRKAEFRKLCD NLRMHLSQIQRRHINQSTAINLNNP ESQSMHLETRLVQLDSAISMELWQ EAFKAVEDIHGLFSLSKKPPKPLM ANYYNKVVTVFWKSGNALFHASTLH RLYHLSREMRKNLTQDEMORMSTR VLLATLSIPITPERTDIARLLDMDGIIV EKQRRLATLLGLQAPPTRIGLINDMV RFNVLQYVVPEVKDLYNWLEVEFNP LKLCEVTKVNLNVVREQPEKEPELQ QYVPQLQNNLTLRLLQVVSQIYQSIE FSRLTSLVPPVDAFQLERAIVDAARH CDLQVRIDHTSRTLSTFGSDLNYATR EDAPIGPHLQSMPEQIRNQLTAMS SVLAKALEVIKPAHILQEKEEQHOLA VTAYLKNSRKEHQRIARRQTIEERK ERLESNLIQREKEELEQREAELOKV RKAEEERLRQEAKEREKERILQEHE QIKKKTVRERLEQIKKTELGAFAKFD IDIEDLEELDPFIMAKQVEQLEKEK KELQERLKNQEKKIDYFERAKRLEEI PLIKSAYEEQRIKMDLWEQEEERI TTMQLEREKALEHKNRMSRMLEDR DLFVMRLKAARQSVYEEKLQKQFEER LAEERHNRLEERKRQRKEERRITYY REKEEEEQRRAEEOMLKEERERER AERAKREEELREYOERVKLEEVER KKRQRELEIEERERRREERRLGDS SLSRKDSRWGDRDSEGTWRKGPPEA DSEWRRGPPKEWRRGEGRDEDRS HRRDEERPRRLGDDEPREPLRPDD DRVPRRGMDDDRGPDRGPEEDRFS RRGADDDRPSWRNTDDDRPPRIA DEDRGNWRHADDPRPRRGLDEDER GSWRTADEDRGPRRGMDDDRCPRR GGADDERSSWRNADDDRGPRRGLD DDRGPRRGMDDDRGPDRGMDDDR GPRRGMDDDRGPDRGLDDDRGPW RNADDDRIPRRGAEDDRGPWRNMD DDRLSRADDDRFPRRGDDSRPGP WRPLVKPGWREKEKAREESWGPP RESRPSEEREWDREKERDRDNQDR EENDKPERERDRERDVEDRDRFR RPRDEGGWRRGPAEESSSWRDSSR RDDDRDRRRERDRDRDLRERRD LRDDDRRRGPPLRSEREVSSWRRRA DDRKDDRVEERDPPRRVPPPALS RERDRDREREKEKASWRAEKDR ESLRRTKNETDEDGWTTVRR	True	False	4.611	4.799	1.222	1.069	0.74