	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q9LD55	EIF3A_ARATH	Arabidopsis thaliana	Eukaryotic translation initiation factor 3 subunit A	27.144759	NaN	NaN	28154133	MANFAKPENALKRADELINVGQKQ DALQALHDLITSKRYRAWQKPLEKI MFKYLDLCVDLKRGRFAKDGLIQYRI VCQQVNVSSLEEVIKHFLHLATDKA EQARSQADALEEALDVDDLEADRKP EDLQLSIVSGEKGKDRSDRELVTPW FKFLWETYRTVLEILRNNSKLEALYA MTAHKAFQFCKQYKRTTEFRRLCEII RNHLANLNKYRDQRDRPDLSAPESL QLYLDTRFDQLKVATELGLWQEAFR SVEDIYGLMCMVKKTPKSSLLMVYY SKLTEIFWISSSHLYHAYAWFKLFSL QKNFNKNLSQKDLQLIASSVVLAAL SIPPFDRAQSASHMELENEKERNLR MANLIGFNLEPKFEGKDMLSRSALL SELVSKGVLSCASQEVKDLFHVLEH EFHPLDLGSKIQPLLEKISKSGGKLS SAPSLPEVQLSQYVPSLEKLATLRLL QQVSKIYQTIRIESLSQLVPFFQFSEV EKISVDAVKNNFVAMKVDHMKGVVI FGNLGIESDGLRDHLAVFAESLSKV RAMLYPVPSKASKLAGVIPNLADTVE KEHKRLLARKSIIEKRKEDQERQQLE MEREEEQKRLKLQKLTEEAEQKRLA AELAERRKQRILREIEEKELEEAQAL LEETEKRMKKGKKKPLLDGEKVTKQ SVKERALTEQLKERQEMEKKLQKLA KTMDYLERAKREEAAPLIEAAYQRRL VEEREFYEREQQREVELSKERHESD LKEKNRLSRMLGNKEIFQAQVISRR QAEFDRIRTEREERISKIIREKKQERD IKRKQIYYLKIEEERIRKLQEEEEARK QEEAERLKKVEAERKANLDKAFEKQ RQREIELEEKSRREREELLRGTNAPP ARLAEPTVTPVGTTAPAAAAAAAGAP AAPYVPKWKRQTTEVSGPSAPTSSET DRRSNRGPPPGDDHWGSNRGAAQ NTDRWTSNRERSGPPAEGGDRWGS GPRGSDDRRSTFGSSRPRPTQR	None	None	None	None	None	None	None	None	None