

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
070194	EIF3D_MOUSE	Mus musculus	Eukaryotic translation initiation factor 3 subunit D	28.986578	NaN	S161;S528;S529	29187734;37453647;34887587	MAKFMTPIQDNPSGWGPCAVPEQ FRDMPYQPFSGDRLGKVADWTGA TYQDKRYTNKYSSQFGGGSQYAYFH EEDETSFQLVDTARTQKTAYQRNRM RFAQRNLRRDKDRRNMVQFNLQTL PKSAKQKERERIRLQKKFKQFQFVGR QKWDQKSQKPRDSSVEVRSDEWEK EEMDFPQMKMRYLEVSEPPQDIEC CGALEYYDKAFDRITRSEKPLRSIK RIFHTVTTTTDDPVIRKLAKTQGNVFA TDAILATLMSCTRSVYSWDIVVQRVVG SKLFFDKRDNDFDILLTVSETANEP PQDEGNSFNSPRNLAMEATYINHN FSQQCLRMGRERYNFPNPNPFVED DMDKNEIASVAYRYRRWKLGDIDL IVRCEHDGVMGTGANGEVSFINIKTL NEWDSRHCNGVDWRQKLDSSQRGA VIATELKNNSYKLARWTCCALLAGS EYLKLGYSRYHVKDSRHHVILGTQ QFKPNEFASQINLSVENAWGILRCVI DICMKLEEGKYLILKDPNKQVIRVYS LPDGTFFSSEDEDEEEEEEEEE ET	False	False	3.16	2.276	1.428	1.46	0.761	1.048	1.136