	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	extracellu region
0	Q5VWG9	TAF3_HUMAN	Homo sapiens	Transcription initiation factor TFIID subunit 3	18.379595	NaN	S183;S199;S229;S243;S291;S297;S301;T501;S667;S755	30379171	MCESYSRSLLRVSVAQICQALGWDS VQLSACHLLTDVLQRYLQQLGRGCH RYSELYGRTDPILDDVGEAFQLMGV SLHELEDYIHNIEPVTFPHQIPSFPVS KNNVLQFPQPGSKDAEERKEYIPDY LPPIVSSQEEEEEEQVPTDGGTSAEA MQVPLEEDDELEEEEIINDENFLGK RPLDSPEAEELPAMKRPRLLSTKGD TLDVVLLEAREPLSSINTQKIPPMLS PVHVQDSTDLAPPSPEPPMLAPVAK SQMPTAKPLETKSFTPKTKTKTSSPG QKTKSPKTAQSPAMVGSPIRSPKTVS KEKKSPGRSKSPKSPKSPKVTTHIPQ TPVRPETPNRTPSATLSEKISKETIQV KQIQTPPDAGKLNSENQPKKAVVAD KTIEASIDAVIARACAEREPDPFEFSS GSESEGDIFTSPKRISGPECTTPKAST SANNFTKSGSTPLPLSGGTSSSDNS WTMDASIDEVVRKAKLGTPSNMPP NFPYISSPSVSPPTPEPLHKVYEEKT KLPSSVEVKKKLKKELKTKMKKKEK QRDREREKDKNKDKSKEKDKVKEK EKDKETGRETKYPWKEFLKEEEADP YKFKIKEFEDVDPKVKLKDGLVRKEK EKHKDKKKDREKGKKDKDKREKEK VKDKGREDKMKAPAPPLVLPPKELA LPLFSPATASRVPAMLPSLLPVLPEK LFEEKEKVKEKEKKKDKKEKKKKKE KEKEKKEKEREKEKREREKREKEKE KHKHEKIKVEPVALAPSPVIPRLTLR VGAGQDKIVISKVVPAPEAKPAPSQN RPKTPPPAPAPAPGPMLVSPAPVPLP LLAQAAAGPALLPSPGPAASGASAKA PVRSVVTETVSTYVIRDEWGNQIWIC PGCNKPDDGSPMIGCDDCDDWYH WPCVGIMTAPPEEMQWFCPKCANK KKDKKHKKRKHRAH	None	None	None	None	None	None	None	None	None