

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q80UV9	TAF1_MOUSE	Mus musculus	Transcription initiation factor TFIID subunit 1	9.739272	T392;T397	S328;S1690;S1693;S1799;S1802;S1820;S1847	34418053	MGPGWAGLLQDKGGGSPSVVMSDT DSDEESAGGGPFSLTGFLLFGNINGA GQLEGESVLDDECKKHLAAGLALGL GSLITELTANEELSGSDGALVNDG WIRSREDAVDYSDINEVAEDESRRY QQTMGSLQPLCHTDYDEDDYDADC EDIDCKLMPPPPPPGPKKEKDQD DITGVSEDEGEGILPSIIAPSSLASEKV DFSSSSDSESEMGPQDAAQSESKDG QLTLPLAGIMQHDATAKLLPSVTELEFP EFRPGKVLRLFLFGPGKNVPSVWR SARRKRKKKHRELIQEGQVQEEEC VELEVNQKSLWNYDYAPPPLPDQCL SDDEITMMAPVESKFSQSTGDTDKV MDTKPRVAEWRYGPARLWYDMLGV PEDGSGFDYGFKMKKTEHESTIKCN IMKCLRKLEENSGVDLLADENFLMV TQLHWEDDIIWDGEDVKHKGTKPQ RASLAGWLPSSMTRNAMAYNVQQG FTATLDDDKPWYSIFPIDNEDLVYGR WEDNIIWDAQNMPRILEPPVLTLDL NDENLILEIPDEKEEATSNSPSKENK KESLKKSRILLGKTGVIKEEPQQNM SQPEVKDPWNLSNDEYYYPKQQGL RGTFFGNIIQHSIPAVELRQPFPTH MGPIKLRQFHRPPLKKYSFGALSQP GPHSVQPLLKHIKKAKMREQERQA SGGEMFFMRTPQDLTGKDGDLILA EYSEENGPLMMQVGMATKIKNYK RKPGKDPGAPDCKYGETVYCHTSPF LGSHPGQLLQAFENNLFRAPIYLH KMPESDFLIIRTRQYFIRELVDIFVV GQCPLFEVPGPNSKRANTHIRDFL QVFIYRLFVWKSADRPRRIRMEDIKKA FPSHSESSIRKRLKLCADFKRTGMD SNWWVLKSDFLPTEEEIRAMVSPE QCCAYYSMIAAEQRLKDAGYGEKSF FAPEEENEEDFQMKIDDEVRTAPW NTTRAFIAAMKGGKCLEVTGVADPT GCGEGFSYVKIPNKPTQKDDKEPQ PVKKTVTGTDADLRRLSLKNAKQLL RKFGVPEEEIKLSRWEVIDVVRTM STEQARSSEGPMKSFARGSRFSVAE HQERYKEECQRIFDLQNKVLSSTEV LSTDTDSSSAEDSDFEEMGKNIENM LQNKKTSSQLSREREEQERKELQR MLLAAGSAAAGNNHRDDDTASVTS LNSSATGRCLKIYRTFRDEEGKEYVR CETVRKATVIDAYVRIRTTKDEEFIRK

							FALFDEQHREEMRKERRRIQEQLRR LKRNOEKEKLGKGPPEKKPKMKERP DLKLCGACGAIGHMRTNKFCLYY QTNAPPSNPVAMTEEQEEELEKTVI HNDNEELIKVEGTKIVLGKQLIESAD EVRRKSLVLKFPKQQLPPKKRRVG TTVHCDYLNRPKHSIHRRTDPMVT LSSILESIINDMRDLPNTYPFHTPVN AKVVKDYKIIITRPMDLQTLRENVK RLYPSREEFREHLELIVKNSATYNGP KHSLTQISQSMLDLCDEKLKEKEDK LARLEKAINPLDDDDQVAFSILDN IVTQKMMAVPDSWPFHHPVNKKFV PDYKVVIVSPMDLETIRKNISKHKYQ SRESFLDDVNLILANSVKYNGPESQ YTKTAQEIVNVCHQTLTEYDEHLTQ LEKDICTAKEAALEEALESIDPMT GPYTPQPPDLYDNNTSLSVSRDASV YQDESNSVLDIPSATSEKQLTQEGG DGDGDLADEEEGTQQQASVLYED LLMSEGEDDEEDAGSDEEGDNPF AIQLSESGSDSDVESGLRQPRVL QENTRMGMENEESMMSYEGDGGD ASRGLEDSNISYGSYEEDPKSNTQ DTSFSSIGGYEVSEEEDEEEQRSGP SVLSQVHLSEDEEDSEDFHSIAGDT DLDSDE
--	--	--	--	--	--	--	--