

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane
P82094	TMF1_HUMAN	Homo sapiens	TATA element modulatory factor	24.602422	S156	S72;S77;S112;S136;S199;S217;S328;S330;S333;S338;S344;S413;S542;S925;S928;T929;S933	23301498;40307207;30379171	MSWFNASQLSSFAKQALSQAQKSID RVLDIQEEEEPSIWAETIPYGEPISSP VSGGWDSTWGLKSNTEPQSPPIAS PKAITKPVRRRTVDESENFSAFLSP TDVQTIQKSPVVSPPAKSQRPEEEV KSSLHESLHIGQSRTPETTESQVKDS SLCVSGETLAAGTSSPKTEGKHEETV NKESDMKVPTVSLKVSSEVIDVKIT MESISNTSTQSLTAETKDIALEPKEQ KHEDRQSNTPSPPVSTFSSGTSTTS DIEVLDHESVSESSASSRQETTDISK SSLHLMQTSFQLLSASACPEYNRLD DFQKLTESCCSSDAFERIDSFVQSL DSRSVSEINSDDDELSGKGYALVPIIV NSSTPKSKTVESAEGKSEEVNETLVI PTEEAEMEESGRSATPVNCEQPDIL VSSTPINEGQTVLDKVAEQCEPAES QPEALSEKEDEVCKTVEFLNEKLEKR EAQLLSLSKEKALLEEAFDNLKDEM FRVKESSSISLKDFTQRIAEAEK KVQLACKERDAAKKEIKNIKEELATR LNSSETADLLKEKDEQIRGLMEEGE KLSKQQLHNSNIIKLRADKKNEN MVAKLNKKVKELEELQHLKQVLD GKEEVEKQHRENIKLNISMVERQE KDLGRLQVDMDELEEKNRSIQAALD SAYKELTDLHKANAADSEAQEAAL SREMKAKEELSAALEKAEARQQQ ETLAIQVGDRLRLALORTEQAAARKE DYL RHEIGELQQRLEAENRNQELS QSVSSTTRPLLRIENLQATLGSQTS SWEKLEKNLSDRLGESQTLAAAVE RERAATEELLANKIQMSSMESQNSL LRQENSRFQAQLESEKNRLCKLEDE NNRYQVELENLKDEYVRTLEETRKE KTLNLSQLEMERMKVEQERKKAIFT QETIKEKERKPFVSVSSTPTMSRSSI SGVDMAGLQTSFSLQDESHDHSFG PMPISANGSNLYDAVRMGAGSSIE NLQSQLKLRGEITHLQLEIGNLEKT RSIMAEELVKLTNQNDLEEKVKEIP KLRTQLRDLQRYNTILQMYGEKAE EAELRLDLEDVKNMYKTQIDELLR QSL	True	True	4.112	5.0	False	3.0	5.0	False