**Text S2.** **Multiple sequence alignment of MTA and MTB predicted proteins**.

**A. MTA protein**

MTA3 -----------------------------------MMILIILICSL-FGALRSAKSLDQG

MT6 -------------------------------MLFLLVILNISIQ--QTLCDWRFVQILDG

MTA7 MKFYLLTFLVKCFIAFQIQFTIQAKNNLCYHLIFYQSLKQILVSRAFSYFDFNYFNIDRG

MTA2 ----------------------------------------MIVT---IVICSDIHFIRNG

MTA5 MKVLLLVA-----------------------------LLNIFIGLVK--PITRFYYTDLG

MTA4 ------------------------------------------------------------

MTA3 IIIEVKH-GDYKFIKIVTNDESSIKNFNGKLDQLNSKIFTINQIVFQQPKE-DYCFLPSP

MT6 LILQIKNDLSLAHVQIVTNDESAITNSTVNQDQF-QNVQIVNQLWFWAKNQ-QNYLLPFS

MTA7 IIFQFYT-LDLIYIQITTNDFASLNNSTL-NNYFSSKMMILSQPSFSIKQN-QTCFLPLA

MTA2 IMFQYTN-QNAKSFEIITNDMSPLSSATTQPDQYKQNVYVVNSLKFPQQQNNKYFFLPFP

MTA5 IVVQVYN-DTLTNVQIITNDEESILNNTIKSDQLNRNIQQINQLWYQQKDS-SFYLLPFS

MTA4 -MIQAFN-QGLTYCEVITNDEASILNQTKSFQNFNDKVYKVNQLAFWSKNL-NNTFFPFA

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MTA3 QNTFGRIISQEFYDENMNLINSAISNIHGQVLIKNTTQKLIDAGNGQKWLSVILNAY-QQ

MT6 SNTFKRRMTYNFSKDQLMGASEASS--------SQLLQTLINN----LKLSMT------Q

MTA7 QNTYNRVVGLDIYDRNNNLLYNLQVKNKVLIKLT--TQKLVYNQDNSMFLSLTFNVF-EQ

MTA2 PNTFNRLLSVKCLDQNENIIDQYFLQNIV--TISSQIQQLINSEDNNKYLQLTLSSSNNA

MTA5 LNTFKRTIIFNYYDSNMVLKGTGGISNYIKAKLNQILIGEVQS--NQAQLLMQISTI-KQ

MTA4 QNSFERNISLNFYDASMNFLGTSYIYSK--VMIKKVLQNYIQDANNNKMLQISLQTY-GQ

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MTA3 FQIISNQINSIFQIYTTIPNLTTNQSAQSIFSLLIDGKKMNCFFNQTNTGIKNGTLFTQN

MT6 MKITIS----------------KSKYK----------QQINSFQSCQS---NMLFSICQC

MTA7 FRIIDPNLQEY-FQIQISPQNKSQTYPQYQFRLLADNQEISSYQTVQGY-----SYFAKQ

MTA2 FRAIQSSTSGN-FKVQFTVENTLKSYPQNFINLIIDGQNITCIQSSYQITFRNQVQYTQY

MTA5 FKIINNNQQDDFQVSFKLSNTVQFPYPDKIFDFMIDGQKMAYSQCFKQVLYNNLVVFNQI

MTA4 FQIIDKTNFYDFFQIKFYTNSALLSSPQDLFSLQADGQTINSYSSYTEL-FITQVIYVKN

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MTA3 FYLSDFPTQFQDLEIQFFNPFFPTFSLSISS-----GADFLQFQISNQLYSSYLLYNSNP

MT6 FKQTDFPQQFKNLEFVFQQYLISSIQIKMQLYNQVSSFESIDISTFNQFYSVQYLLSNDM

MTA7 YNQSSFPTQFTNLEIIFSDYSFQTVYLQLAL-----GAAQTSMTIYNQYYSVQQLQIANF

MTA2 FNYSEFPSQFQNFGFQFYGYFFLGFNLQFSV-----GYDFNVIQGWNQFYSIYVIQNQTN

MTA5 FQQSDIPISFQKMQLVFNQYQFQQFTIKIIF-----GSDMIQFDNFNQHFSTQFLMSISL

MTA4 FNQSDFPDKFQNLQLIFSEYRFQNINLQILF-----GSTNIKVTDYNQYYSALQLQNNNP

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MTA3 SAMIIPQS-DNLIANQLVFILQSDIFSQCYVQQLDLPSNMNTQFSFSQPVLNVQNPQNRS

MT6 QALVRPSSLGKLLAYELAFNLNFTIFVNCDLYQIILPPSISQQFTKAAPTLMAQFQTNSN

MTA7 NALIKPAPQTNFLGNEIDVQLQFSIYQQCSLSQIIFPPEISAQFSQTQPVLQLQNVINNQ

MTA2 LAYVRPVQYGDFIANELAFNIQQSAFNFCNLQQIVLPQSISDQFQAIDPILLLQNTVNQN

MTA5 NATVNPSSQFNQIYNQIYFNIDQTVFFQCSIKQIILPPLLQSQYTNQLPSLIITNSQNSN

MTA4 QAIIRPSSFLSQIKNELAFNITDVVFLQCLLSSILLPSEFQNEFITAEPSLFVFYPTNKT

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MTA3 QSSIIDSNFY-NGTISFVRNQNLRVLPSQVSYSDTLVAIS--SFLDVIGFDNNLTMTFVC

MT6 -FTEQVTDYYNNGTLILQKSLSFKSSNTLN---NIQILIQQFEFQDQIDDNSSSSIVFNC

MTA7 -IQYLETTFD-SRTLNLKQKQNTLQQQD-N---SNVLVIQ--SFNDVVGINKNLSFVFNC

MTA2 NFTSQEIIYVNNGTIMF-KNQNLQIISQLN---PIPFILK--QFVDVVGFNQNLSLTFSC

MTA5 -TTLMSTDYN-NGIISLQQTQVFQQFKYQA---YYSIIIT--PFNDVVGQNMNLTATFNC

MTA4 -FQNQETYYM-QGNISLKVSQNFFPNRANN---QIQILIS--PFSDIVGQNKQLNVAFNC

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MTA3 SQNLIQKKSIRIIKNEYLNANQTQVSIQVNNLVGTKNISIQISIPSVTIQNTSLIMQLPQ

MT6 SQNNLFVKNFTISKIQLPNANLTQTQIVLNNTD-IPTLNIIIQIPPINILSSVLTLELPN

MTA7 FQNSQFIKNVTLSATTYSNSNQTSALLQFYNYD-QKNLTINISIMKIYIQNTSLFLRLPQ

MTA2 NQNQLITKNYSIIQNEFLNANLTQALVQLNNNGNTPNLTIQIQIPVVQVQNSSFYLKLPL

MTA5 PQDVQVSKNITIYQSEFPNANQTITQIQFNNSG-IKNLTISIQIPRVNVLNTSLYLQLPQ

MTA4 QNNIQTIKNISISSNELLNANQTQTQIKFDNNQ-VKMLSINIQIPQVQVSNTSLFIQLPY

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MTA3 SILFQDNTQQSLDVQGFVYSNFVWQQNNLAFFGANFSSTILRLSFTNIDVKNQTSNVDIL

MT6 GVQCLNQSQTNILTNGFVYSQYQSIDNSISFLSTNFTSASISITFTKVNKID-----GNQ

MTA7 GISYLDQTQNSILVNGFVFSQFQWQQGIISFSQANFSSTSIQISIKNVQLNVQSN-G-GY

MTA2 GINFLAMDQSQVSVQGFSFSQLNWSQNIISFLQSNFTSTSILISVSKVELQKQSN-G-KQ

MTA5 EVDYQQQTQQSIIVQGISFSQFNWNEQIIYFGQTNFSSNVISFQILNVQLKNSSN-QTSF

MTA4 GIDYLQMNQSSLSVQGFAYSQFNWDQNLISFYSTNFTSSSILISIKQVQLSSSSN-N-IQ

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MTA3 NSTFIIAKCIFNQTFVFYVDSNTNQTVKIIQPT-PLPRSQITLSSFNQTKVNSLDVPESR

MT6 NLTKINVKCIFNSTVLFYVDANTQQTITIIQPPSPPPQTQIQINSFNQTSFSSLSIYENR

MTA7 NISTILVKCIQNSSFVFQVDSNSKQSYQIFELP-SPPTTLIKINSFNQTNANSLNILENR

MTA2 QITKIIAKCVFNSTFVFYVDENTNQTLQIIQPS-PTPITQIQINSFKQTSDVSLDIPEDK

MTA5 KINKIMAKCIFNQTFVFYVDSNTNQTTQIIQPT-KVPITLIQINNFNQTTVNSLSIPENR

MTA4 NVTKIIAKCILNQAFVFYVDSNSNQTIQILQPQ-PLPTTSIQINTFNQTQSTSLSSPQNR

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MTA3 TSVQSQIAFSFSVLNFLDTCSWLIVQIPFEFTIGFIQMSQFTLQDCFGNIYSYLQGSSIL

MT6 TSVLSQLAFSFSILKFLETCSWLIVSLPLEFTTSFQQNSSFNLFGCHGNNYQYTQVNPLS

MTA7 TSVQSQLIFSFSILQFLDTCSWLIVTIPNRFTIGFLKQSQFNLQDCVGNSYSYLQGNPIS

MTA2 TSMMSQLVISFSILKFQSTCSWIIIDLPQDFRIGFIQQSQFSLQDCFGNSYTYLQGSSIN

MTA5 QSVQSQLAFSFSILQFQDSCSWLIVSLPLEFTLNFMQQSVFNFQDCYGNTYSYQNGSPLS

MTA4 TSLQSQLAFSFSILNFQDTCSWLIVSLPYSFTIGFLDKSQFTLFDCFGNTYSYTQGNPIS

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MTA3 PQSMISYTDNNNCMYISCSSLRLNSQM---NSSSQCMNNTVIIQSIKSPDQAVQTSSLNI

MT6 AQSVIGYTQNNNCIYISCLALQQFSQLNNNNNNNYCLNNTVTIQNVKSYDLATQTQGLKI

MTA7 YQSVISFTDNNNCIYISCSALKLVSQS---NNNNQCQNNIITIDNVKSPDLPTQTESLNL

MTA2 SQQTLSYTDNNSCIYISCQTLRQIAQN---HNTDQCLNNALTIKNVKSPDYATQTDNLNL

MTA5 NQSPIGYTDSNQCIYISCLNLRILSQA---NQSNQCLNNTVIIQQVRSPDLPIKTWSLDF

MTA4 SQSTIGYTDNNNCIYISCKALRSVSSS---HAANNCLNNTVTIRNVRSPDFPLQTAGIQF

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MTA3 YIANQNSSAPSNIGPPTFFTQNLNLKNNSLPYFFIPTEVYSHVGINITEADFSDLNFNIS

MT6 FVANRNSSNKQDLNPPTFFSENLDLNNNSLPYYFTNSEVSTYQGINLTEVDLSYLAFNTI

MTA7 IIANSKTPAQANNFPPTFYTSSLNLSDITLPYLFFSSEQQRYQGISLQQVDLSSLSFYAS

MTA2 YIANSYSSTPQNNYQPTFFNSSLNFTDSSLPYYFNPEEVFSYQGINVKPFNLSALSFQIS

MTA5 FVANSNSSMQSNNNPPIQFNRSLDLQNNSLPYFFIPTEISTYQGINITNFDLSQFSSQIS

MTA4 LIANSNTSSQPNSDPPTFFNQSTDLTNSSLPYFFIGSEVFTSKGINISQVDLSTLSTQIS

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MTA3 SNYYGDVFNLTTTISYPFYLYQQHQISIQIPVRIL-ANQSVVCYPESFLSCSIKIPINLN

MT6 STYLRDIFNFTMTFSYPIYLWEKHHLNIQLPYQFL-GSNNITCQPASFVQCSLQMGTQQN

MTA7 SNYLGDVFNFSMTFSFPIYFSQQHKINVDLPVKLL-ANQTVQCFPQSKIQCSLQIGDQQI

MTA2 SNYYRDIFNFSMIFSYPIYLFPQHKVIVLLPFKVL-INQNLTCFPA-NIQCSLSIISQDI

MTA5 SNYQGDVFDFQMILAFPIYFWDQHLISVQVPFKIL-GNNSLSCQPSTNVICSILNIKGLD

MTA4 SNYFGDVFNFSMILAYPIYFWEQHQIDVQLPFRLFAGKSDIECQPISTVVCSLQASRDPQ

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MTA3 F-TTILIQILQQVSPNTKIAFQLNQVVASQSLLQGKQAFALIQVSSTSKIINTKNKTIDV

MT6 Y-STLNLLILQQVLPDTKITIEVYQVVVVAENTI-QQKQAVIKLFLENRLINTKSKIIDV

MTA7 Q-TTIQIIFVDQVIPDIKITLQLFCAALQQSNQQNQQFFSLIRIFQQNQIINTQNKTIDV

MTA2 YPTSVLIQFTNQIL----------------------------------PSINSQNKTLDV

MTA5 F-TNVQIQIINQMKPNTQIILNLQEVAVSQ---LNQTQNMTISVLLSNRIINTLNKTIDA

MTA4 S-TIFRIQVLNLTAPNTQIGLQVSYVVLSQNFQLDQISFLIISVSLSNRIINYQNKTIDA

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MTA3 LNNLKLQSWINIDAIQPLQISNTLLGQEGVNYTFAILSLQIPSNQLDNYYLSLKIDQSLS

MT6 LSDLMPQNFIVFSTSSPLQLSNSLLGLSDVNYTFFIKQMNLPPKINDNYYLSLTLDQSFL

MTA7 FSNFTTTQWITFST-TPLQISNSLLGIQGNNYTFLISKLFTPPNLTENYFLGLELDSSLQ

MTA2 LTNSTQQNWINFENNLPLQLSNSLLGISGVNYTFFIQSLYLPQNLNSNYYVKINLDQNIQ

MTA5 LKSLTFQQWITFQKDKQLEVSNTLLGFQRVNYTFAIQSLQLPKNLNSNYYLSLTLDPNIQ

MTA4 FKNLAIQNLITLQNQQPLQISNPLLGITGVNYTFNIQQLQIPQNINNYYYLSLSLDPSFQ

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MTA3 YNSSNSNCYSLTQVT---GDNSQREEKVNCINFSSNTFLIPISKN-QISQPFQLRITGLR

MT6 LVSNNYNCSILTQGE----NQIYLEQQLKCLFYFPNNFIFSISSFAIMSQPFQLRISGLR

MTA7 FNQTSLNCYLLNITLDTS---LINENKIDCFFSSMNMLILPLSKI-LTSQSFQLRISGLR

MTA2 FNSSEISFFQIIQTQDG--KMTLSENQITNFYYASQAFFIPLINL-NLSKAFQLRISGLR

MTA5 FNTTEVYCSQLILTTDSSGVNQVKEISLNCIYNFFNQFLIAIQSI-SVSQPFQLRISGLR

MTA4 YNSSQIAFYQLIQSQ---NLNTFIENRIDFISNSSNLFLIPITSL-VPSYPFQLRISGLR

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MTA3 NPSGIVEQ---NVAQTYNFQLIWSTQQGNSFKNVWVIQSVSLPITSKYTCSPNCQACASN

MT6 NPSGIVEQNEQNVAQTYNFQLIWSTQQGNSFKNVWLIQSVSLPITSKYTCSPNCQACASN

MTA7 NPSGIVEQNEQNVAQTYNFQLIWSTQQGNSFKNVWLIQSVSLPITSKYTCSPNCQACASN

MTA2 NPSG-------NVAQTYNFQLIWSTQQGNSFKNVWVIQSVSLPITSKYTCSPNCQACASN

MTA5 NPSGIVEQNEQNVAQTYNFQLIWSTQQGNSFKNVWVIQSVSLPITSKYTCSPNCQACASN

MTA4 NPSGIVEQNEQNVAQTYNFQLIWSTQQGNSFKNVWVIQSVSLPITSKYTCSPNCQACASN

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MTA3 YAACTACAPGYLKSQYNHHAVLACLPTCSPQYVAYNGTCLACQLKDPQCLSCSPSNLTQC

MT6 YAACTACAPGYLKSQYNHHAVLACLPTCSPQYVAYNGTCLACQLKDPQCLSCSPSNLTQC

MTA7 YAACTACAPGYLKSQYNHHAVLACLPTCSPQYVAYNGTCLACQLKDPQCLSCSPSNLTQC

MTA2 YAACTACAPGYLKSQYNHHAVLACLPTCSPQYVAYNGTCLACQLKDPQCLSCSPSNLTQC

MTA5 YAACTACAPGYLKSQYNHHAVLACLPTCSPQYVAYNGTCLACQLKDPQCLSCSPSNLTQC

MTA4 YAACTACAPGYLKSQYNHHAVLACLPTCSPQYVAYNGTCLACQLKDPQCLSCSPSNLTQC

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MTA3 SSCNQGYTLVPEFNGCVDSHLLQTARSRLLDYSLSTNLADNIDTTHPDDAQKSDKAQSLT

MT6 SSCNQGYTLVPEFNGCVDSHLLQTARSRLLDYSLSTNLADNIDTTHPDDAQKSDKAQSLT

MTA7 SSCNQGYTLVPEFNGCVDSHLLQTARSRLLDYSLSTNLADNIDTTHPDDAQKSDKAQSLT

MTA2 SSCNQGYTLVPEFNGCVDSHLLQTARSRLLDYSLSTNLADNIDTTHPDDAQKSDKAQSLT

MTA5 SSCNQGYTLVPEFNGCVDSHLLQTARSRLLDYSLSTNLADNIDTTHPDDAQKSDKAQSLT

MTA4 SSCNQGYTLVPEFNGCVDSHLLQTARSRLLDYSLSTNLADNIDTTHPDDAQKSDKAQSLT

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MTA3 RMTEESSSAKADQREGGESSSNTGSKVMGQLQDTVGALKGGGAIFIWIVLAALAVSVSQS

MT6 RMTEESSSAKADQREGGESSSNTGSKVMGQLQDTVGALKGGGAIFIWIVLAALAVSVSQS

MTA7 RMTEESSSAKADQREGGESSSNTGSKVMGQLQDTVGALKGGGAIFIWIVLAALAVSVSQS

MTA2 RMTEESSSAKADQREGGESSSNTGSKVMGQLQDTVGALKGGGAIFIWIVLAALAVSVSQS

MTA5 RMTEESSSAKADQREGGESSSNTGSKVMGQLQDTVGALKGGGAIFIWIVLAALAVSVSQS

MTA4 RMTEESSSAKADQREGGESSSNTGSKVMGQLQDTVGALKGGGAIFIWIVLAALAVSVSQS

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MTA3 VLRYTYNKLKGKSHSNGSSSGMRRSSSGSRSSSSGSSSAGRNASGKEHKGVREREKELRG

MT6 VLRYTYNKLKGKSHSNGSSSGMRRSSSGSRSSSSGSSSAGRNASGKEHKGVREREKELRG

MTA7 VLRYTYNKLKGKSHSNGSSSGMRRSSSGSRSSSSGSSSAGRNASGKEHKGVREREKELRG

MTA2 VLRYTYNKLKGKSHSNGSSSGMRRSSSGSRSSSSGSSSAGRNASGKEHKGVREREKELRG

MTA5 VLRYTYNKLKGKSHSNGSSSGMRRSSSGSRSSSSGSSSAGRNASGKEHKGVREREKELRG

MTA4 VLRYTYNKLKGKSHSNGSSSGMRRSSSGSRSSSSGSSSAGRNASGKEHKGVREREKELRG

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MTA3 QRIDCLMLFLLSVAEAIQMPYTLLWGFSVSGGDFESPVLQTLLGACALSTLLWIFDAYLL

MT6 QRIDCLMLFLLSVAEAIQMPYTLLWGFSVSGGDFESPVLQTLLGACALSTLLWIFDAYLL

MTA7 QRIDCLMLFLLSVAEAIQMPYTLLWGFSVSGGDFESPVLQTLLGACALSTLLWIFDAYLL

MTA2 QRIDCLMLFLLSVAEAIQMPYTLLWGFSVSGGDFESPVLQTLLGACALSTLLWIFDAYLL

MTA5 QRIDCLMLFLLSVAEAIQMPYTLLWGFSVSGGDFESPVLQTLLGACALSTLLWIFDAYLL

MTA4 QRIDCLMLFLLSVAEAIQMPYTLLWGFSVSGGDFESPVLQTLLGACALSTLLWIFDAYLL

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MTA3 GSILANSENTPKTSCLLNPLPSSKREGCLSFLIVPARLAFCLIPKSVSLTLTNIFSVEGW

MT6 GSILANSENTPKTSCLLNPLPSSKREGCLSFLIVPARLAFCLIPKSVSLTLTNIFSVEGW

MTA7 GSILANSENTPKTSCLLNPLPSSKREGCLSFLIVPARLAFCLIPKSVSLTLTNIFSVEGW

MTA2 GSILANSENTPKTSCLLNPLPSSKREGCLSFLIVPARLAFCLIPKSVSLTLTNIFSVEGW

MTA5 GSILANSENTPKTSCLLNPLPSSKREGCLSFLIVPARLAFCLIPKSVSLTLTNIFSVEGW

MTA4 GSILANSENTPKTSCLLNPLPSSKREGCLSFLIVPARLAFCLIPKSVSLTLTNIFSVEGW

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MTA3 FRYPYGEDVERATIVLTNFRKVLSSQIKSNVSGLCSLVSFLILQYPEVLSSYVQIYDLII

MT6 FRYPYGEDVERATIVLTNFRKVLSSQIKSNVSGLCSLVSFLILQYPEVLSSYVQIYDLII

MTA7 FRYPYGEDVERATIVLTNFRKVLSSQIKSNVSGLCSLVSFLILQYPEVLSSYVQIYDLII

MTA2 FRYPYGEDVERATIVLTNFRKVLSSQIKSNVSGLCSLVSFLILQYPEVLSSYVQIYDLII

MTA5 FRYPYGEDVERATIVLTNFRKVLSSQIKSNVSGLCSLVSFLILQYPEVLSSYVQIYDLII

MTA4 FRYPYGEDVERATIVLTNFRKVLSSQIKSNVSGLCSLVSFLILQYPEVLSSYVQIYDLII

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MTA3 FDIVMTILCLTNIRNVDKLISTLNQIQESEGEL

MT6 FDIVMTILCLTNIRNVDKLISTLNQIQESEGEL

MTA7 FDIVMTILCLTNIRNVDKLISTLNQIQESEGEL

MTA2 FDIVMTILCLTNIRNVDKLISTLNQIQESEGEL

MTA5 FDIVMTILCLTNIRNVDKLISTLNQIQESEGEL

MTA4 FDIVMTILCLTNIRNVDKLISTLNQIQESEGEL

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**B. MTB protein**

MTB3 ------MTDCKIFIWFYILYLISKFCKCQVDCYLASSTQIGQQIQPIAYSQRQGQT---S

MTB6 ---------MQVYTLVFVLILLIRKNSAQTICYQISADKNLKQIQQLPLNQRLEFY---N

MTB7 -----MSGLRK-VSLLMAIYQILCLSYGQLSYYLVSLNQNTKQLNPLPLNQRQISS---N

MTB2 -----MNRA---IEFLFLVILIIPIIKGQLQCYQVNLNQSQQQIKQLPLNQRDGNP---V

MTB5 MTSFFIKRKIQIVYCLFYIQCIIPLTKQQINCYKVLVSQSQTQLTPLPLNQRVEYTQVDY

MTB4 -----MNASQIIYLLIFVAALLIERSQATLSCYLINPTQSLKQIYQLPFNYRQGSD---A

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MTB3 FGAQYFFTIQLASNQTDIL-QGTSSVQFT-QYFLTSNKFVLSSMISKNM-GLLPTPQQYV

MTB6 KIPNYNFALDFSINNSDFLANYPPSLFFTYLFITTSNKQVLDKFTTFPS--IQQIQQNYG

MTB7 NLQQYYFVVEGN-ITQSND-LMGLGLDYTYRLLTTSNRVILYYFSNSD--QLSSIPTQFQ

MTB2 IVQTNYFGFEML-LGQDSD-LNQLTFTFTE-VFLTSNRNIISNQTIFYGNIVDPPYSIIN

MTB5 QSQLYYFALEYSQLFL-NN-LPSNDIQYTYQYLTTSNRQVIKIYDNTSQ--FFGVPDQFK

MTB4 APAQYNFALQYDQLSPDVQ-YTLQNIAYLKRYLITSNNQVISIIPNTIGDSLN--YIPPE

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MTB3 NSTYYVAAAFTYQ---QFISGALDFSSICNLDQYPLLTLKEFVQVNTQTQQMFVSVPLTS

MTB6 NQAFYITLVFDSSSFQQ-AIDSQALQAYSSITFFPMLTLPIFVQINSQSQQLMISSTLIS

MTB7 QNGFYINLLVSLQFSQGFNQVMQFINSNDALVQYDQLTLKKFVQLNTQSKQFLISTVLNS

MTB2 QVKYILILVNYQQFMQNF---QQSYSTLCGISQYSQLVQKEFVQVNSQNKQFIVSTVLSS

MTB5 QATFYVTQMINLK--NLNSNQLQNFFGICALNYFAQLHIKEFVLLSQQNQQFAISFDLIS

MTB4 LIQYTFNIVTIYQSIANFSDQPDFLTSYCQITQFPQMDLTTFRQVNKQNQQFLISAILLS

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MTB3 QINDVYYLTLFYPTSLASQPQNNNNLGCQLYTDYIVINQCSLVINS--TLTTVSFNLSSI

MTB6 QINDVSTLVLLYPTSFASQPQNSNNLGCQLYTDFVSINQCSFATNSSSGLSSISFNLSGQ

MTB7 QTDDISNLILQYPSSLATIAQNSQNIGCQLYTDYVVLNQCSLIANPSTGLTSISFNLSSI

MTB2 QISDQATLTIQYPTSFASQSQTTSNLGCQLYTDYGNLNSCSFSTNSSSGLSIVSFDLTNL

MTB5 QTDDLSYVVLQFPTSFAQKPQTNSNIGCQLYTDFITIDKCSLTTNATSGLTSVSFNLTNL

MTB4 QIEDLTSLVLSYPTSFALKSQNAANLGCQLYTDLGIINSCSFSLNAQQGLSLVTFNLTNF

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MTB3 NI---RNISATLVLSTATFNQNFIKQNQPFQIQLLNQLQNKIAQSPAFGAKDYRQCQLT-

MTB6 NV-SLQNISATIVLNSTTFIQNKLS-TVKFQMQLLNYYQSQIAQSPLFAIQDQRKCQQTT

MTB7 PI-PLTNISATVAINYSSFNQNFIQ-AQPFSLQLFNIYNIKISQSSNFGGVDQRVCQQS-

MTB2 NF-YQQNISASVILNYSTFNQQMLL-KPQFKLTLLNQYSYAIGGTPTFSILDQRICQQT-

MTB5 TQNNLQNISAALVLNYSTFNQNKFQ-NQQFQVQLLNYYSYLIANTTQFAIVDYRVCQQA-

MTB4 NI-SLQNISATIVINYTTFNQNLLS-NKQFSLQLLNQYQSQISQSSQFQIVDQSICQQT-

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MTB3 NFQAQLIFNNSYSISEQRLQVQFATASPAISRILIEFSSQILPKFNLTNVLVMVSDLSKL

MTB6 SFSANLIDNTMQFLQMKRLQVSFTPATQKITRLMIDFLQQVNVKTNSSNIVLILSDTSQA

MTB7 SFSAVLVNDTQAYNQRPIWQVRFTPT-TNITRILIEFSQPIIAKIKGNTIPFTLSDLSGN

MTB2 GFSSILSPSILKLNQQDKLQINFTPVSNNITRLIFEFNQQIIPKFNLTKIFITLADSSLT

MTB5 SFSAEIINNVQDFSQNERLKIKFTPASSNITRILIDLNNQAIIKINLTKIVIELSDLNQS

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MTB3 NPYAVFCQKAYQSIILCELFSPASPFDMSKGVILTLPYFQPMQAVPQNSYSFTVKYFTDS

MTB6 NSFVIFCNKVYQSLVLCDLTN-YQLFDFTKGILITLPYFQTVQQVQQPTFSFQIIYYGDL

MTB7 NQFLLSCNKIYKSIILCDLYG-YIPFDNSQPIILNFPQFDIISSTLQPQIQFSFKYFTDF

MTB2 NSVITQCYKAYLTMIVCDLEF-QLTFDITKSIYVILPYLQIPSLVTLNALTLSFKYFTDS

MTB5 NSLKILCSKVYQSLILCDLYN-AFDFDITQGILMTLPFFQLTQPLQLMNITISIKYYTDY

MTB4 TKFKVRCLKAYQFLLVCDLIG-GLPFDVSQGVYAELPYIQVKVPSYPQSLTFQIKYFTDL

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MTB3 TYQTCYNTNITIPFIPQD-IESPVAFLFNNDSLLLQFSNVINLETSTTINIQLPSQLSFA

MTB6 SYSNCYVTTLTVQYLPIPSLFSPIAYLYSNNTLQMSLFDKLSLDSSMFIQIQLPNLLSFS

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MTB5 SYKNCYFSNTTLQFQSET-SSSANAYLYSNNSFQISISNLIKLENQMTINIVLPSQLSFS

MTB4 SYQNCFVSNLTQKFQAAI-PTSPQAYLYNNSSLQIKLFDLIMLDNSTVINLSLPSQFSLS

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MTB3 PTSQLAQITGISSLSKILISSSQNSKISQVTQSLQQLTTNQGIQFQLLNVTRSSIFCSLT

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MTB2 VSSQLIQVSGISNLSQMTYKSQQSLQISQIAQQKSQQFA-YGISFQLTQLSQSSIFCNFS

MTB5 QNVQLNNAFGFNSTSRVQVVSQQQVKISQLVQSYQFIPTYSGISFQLINVTSNTVYCNLS

MTB4 SSTQINSIVGISNQSQITLSSQQNLQISQISLSLAQLMTYQGISFQLTKIIQNTVNCNLS

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MTB3 NIASLQIQIVNSFGVVIMSGTVPIQIQSYSIIVTSINSNQILNLQ-------Q-QDNDLT

MTB6 STNLVRIQITNTQGLLIYNITAALQIQPKQINIVSINTIYLTPDSQSQS--QQLNQYFNV

MTB7 STYQAQIQIINSLGTIILSGNISIQIQPYPIQIVSTNTIQVPQSP---------------

MTB2 NPYSVQIQIINSFGNLILTGSIPIQIQPYPITILSLNPIYMPNSDSQQM-----QNSSKI

MTB5 TQLSIQFQVINSFGLIIISGSIPVQIQPYPIQIQSTNQIYLTNPQQQ------QQNNNKN

MTB4 SLSQIQIEIVNSFGVIVLSGSIPLQIQPYPIYIQQMNAVFTPLSQQQLLQYQQQQNSYTV

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MTB3 IPVSLEVNFQLKARYFPQSSAFVIQLPLQLTRDNRFSQVTVEIVNEFNFFCKSQTNSTKF

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MTB7 --TILQIDFQLQINYLPYSSAFVISLPPQVIRDSSISNVSVDISNQFNFYCKSQTNQTQF

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MTB4 PFVTLEVNFQIQINYFPQSSAFVIYLPPQIIRDNRVNNISIEIANQFNFYCSSQTNSTQF

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MTB3 STSKVTYQNTVQQRDTLVIQCQFSGNQTINISSDVFTAKIQGYQLPMQVQKPTDRVIINL

MTB6 SNQTVNYQNKIQQREVISIICQQSGNTN---TQSVFTVTIQGYQLPRQLQKPTDRLIIDL

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MTB2 FSSSIQYDQNLQQRDTISVVCQFYGNTTFKDPQSIFTAKIQGYLLPKQVLKPTDRIIIDL

MTB5 SSQSIQFENQQQQRDSVSIICQFNGNDTLNSQQNLFTVRIQGYQLPEQLYKSTDRVIINL

MTB4 FSSNITFQNKIQQRDSISILCSFSGNITTNNPQYLFTAKIQGYLLPKEFQKPTDRIIITL

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MTB3 VDFSQNSFNEQYYCQSSENQNIPDSLKNIAVFVKTQSTNFIVSQYSYQIT-SYINQNKVS

MTB6 IDFSQKQNQPLPLCLTTENPDISNQVKSSWVFTNTKQTSFFVSQYDFKFNGS---SDQNT

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MTB2 IDFQQNSYQRTFQCQTSESLNNKPQLKNQWVFTNTQSSNFVQNLYFYQINTNSSNQSQAN

MTB5 FDYSQKQFNQQSLCQSSENPNIQLQIKDKWVFTNTQVTSFQFSQYSYEGN-----DSQNL

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MTB2 TSVISIINITTSNQIQDGDDIVVQFPKQQFLKYESNQLAS--SQYFSELKNLTCIAIFPQ

MTB5 SSLKSQINITTSQNIQDGDQIVIQFDKQNFLKLENISLTQNN---IIQNESLNCSSIMPL

MTB4 NVPYSLINLTTRYPIQDGDYIVVSFSKLQFLKNEVINGIQ--ALAYPELKNLNCSFLQPQ

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MTB3 PAQPNSVSFIPTCVLVEKASSFSIQFQLYPSQLGQVLNWNNKDVVLNISGLAFQENLQQN

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MTB7 SNGIASSSLLFSCTLSEGPSYYSINLGLRPSTAGVNSSWNNKDLVVNIIGIGFYPNFL-N

MTB2 ATL---QTNFISCILSEVSSYFSIQLTIQPNQL-SQIDWNKQDLVIQFTGVAFQPNLQ-N

MTB5 SDPFPSSNFISSCVLLGGPLSYSIQIVLYPSKLGQISSWNYRDLIISFTGFVFQPSTQ-C

MTB4 SSF---LNLFSTCNLSEGSSQYSINLQLYPTQLDSNLSWNYKDLALSVAGLQFFPNQQ-Y

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MTB3 YQSEVLFSHSSSDLYLLSQSNQTFNNQLSVTANSVYQVSQISTLSS--LAQDYLNYKFDF

MTB6 NNTKIQFTHFSSDQYLLSQSNYLLNS-QQTTSNVFYSINQISSVTQQTITSFVQTIKFDF

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MTB5 FNCIIYFQHYQNSNQLVSQSNFTYFTNQQLLTNLSYSINKIDYLSQ--GINSASSLQFDF

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MTB5 IQIQYLTYLSLPSSNSSITLFLDFSQQLSFGPQSFCSIDSTCSQKTFITQCDLVSNGLTI

MTB4 IQLQFQTTLNLSSSSQNLSLTLQFSQPIYLSNLSFCSIDVGCSQSTLISTCSLSADGLLV

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MTB3 TLDNIGSWITCS-SVLSSFNITIHNPEINNQNSTGGAAQSIAINWNLTSTTSSQTLLSGH

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MTB5 QINGIDKYVTCSTPNLTSFNITIHNPEINNQNSTGGAAQSIAISWNLTSTTSSQTLLSGH

MTB4 QINKLDQFSTCASSVLNNFNITIHNPEINNQNSTGGAAQSIAISWNLTSTTSSQTLLSGH

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MTB3 TSLTTNTSQCPQPHCATCTSPPSICIHCTQGYYLLPDQNSCVQTCPPPTVAHQQTATCQP

MTB6 TSLTTNTSQCPQPHCATCTSPPSICIHCTQGYYLLPDQNSCVQTCPPPTVAHQQTATCQP

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MTB2 TSLTTNTSQCPQPHCATCTSPPSICIHCTQGYYLLPDQNSCVQTCPPPTVAHQQTATCQP

MTB5 TSLTTNTSQCPQPHCATCTSPPSICIHCTQGYYLLPDQNSCVQTCPPPTVAHQQTATCQP

MTB4 TSLTTNTSQCPQPHCATCTSPPSICIHCTQGYYLLPDQNSCVQTCPPPTVAHQQTATCQP

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MTB3 CFQHQECLQCQSQNPAACTSCSPTYSLNSTLLPYCYVPLPPSSSASSSVTKDVVNRTPSN

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MTB4 CFQHQECLQCQSQNPAACTSCSPTYSLNSTLLPYCYVPLPPSSSASSSVTKDVVNRTPSN

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MTB6 STFSGALNRPEPGQPSQKQQQQQQQQQQQEQQQQQEQQQQEAQASDQRGFAHFLAQTKSY

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MTB4 STFSGALNRPEPGQPSQKQQQQQQQQQQQEQQQQQEQQQQEAQASDQRGFAHFLAQTKSY

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MTB3 TKGFILTLLIPLSILGACLTRLVTFCLKKREKKVHPPLPSESRSAQIAQNLDERQETQKD

MTB6 TKGFILTLLIPLSILGACLTRLVTFCLKKREKKVHPPLPSESRSAQIAQNLDERQETQKD

MTB7 TKGFILTLLIPLSILGACLTRLVTFCLKKREKKVHPPLPSESRSAQIAQNLDERQETQKD

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MTB4 TKGFILTLLIPLSILGACLTRLVTFCLKKREKKVHPPLPSESRSAQIAQNLDERQETQKD

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MTB3 GNGGDEEQMRASSRVDTANMCPLNSRRGEQLQLEGVQQQSDGGVGGGESEGNGYLAFCWI

MTB6 GNGGDEEQMRASSRVDTANMCPLNSRRGEQLQLEGVQQQSDGGVGGGESEGNGYLAFCWI

MTB7 GNGGDEEQMRASSRVDTANMCPLNSRRGEQLQLEGVQQQSDGGVGGGESEGNGYLAFCWI

MTB2 GNGGDEEQMRASSRVDTANMCPLNSRRGEQLQLEGVQQQSDGGVGGGESEGNGYLAFCWI

MTB5 GNGGDEEQMRASSRVDTANMCPLNSRRGEQLQLEGVQQQSDGGVGGGESEGNWYLAFCWI

MTB4 GNGGDEEQMRASSRVDTANMCPLNSRRGEQLQLEGVQQQSDGGVGGGESEGNGYLAFCWI

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MTB3 AVILLLGNVGDLVEVPYIIFSQQNSFSNKSTTNVFDLQFSEADMGQICCLSYIALNAVCY

MTB6 AVILLLGNVGDLVEVPYIIFSQQNSFSNKSTTNVFDLQFSEADMGQICCLSYIALNAVCY

MTB7 AVILLLGNVGDLVEVPYIIFSQQNSFSNKSTTNVFDLQFSEADMGQICCLSYIALNAVCY

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MTB5 AVILLLGNVGDLVEVPYIIFSQQNSFSNKSTTNVFDLQFSEADMGQICCLSYIALNAVCY

MTB4 AVILLLGNVGDLVEVPYIIFSQQNSFSNKSTTNVFDLQFSEADMGQICCLSYIALNAVCY

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MTB3 LICIVMMVKAIIFETGSGEPLFCIYEVKLSSSSSFCGEEMKKDKSEAAACKGDSNAQNEN

MTB6 LICVVMMVKAIIFETGSGEPLFCIYEVKLSSSSSFCGEEMKKDKSEAAACKGDSNAQNEN

MTB7 LICIVMMVKAIIFETGSGEPLFCIYEVKLSSSSSFCGEEMKKDKSEAAACKGDSNAQNEN

MTB2 LICIVMMVKAIIFETGSGEPLFCIYEVKLSSSSSFCGEEMKKDKSEAAACKGDSNAQNEN

MTB5 LICVVMMVKAIIFETGSGEPLFCIYEVKLSSSSSFCGEEMKKDKSEAAACKGDSNAQNEN

MTB4 LICVVMMVKAIIFETGSGEPLFCIYEVKLSSSSSFCGEEMKKDKSEAAACKGDSNAQNEN

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MTB3 GEKHKKVLSGRKLWKLIIDVFLRCLVVVGGKAFCMVYSNVANVKGWLTCQVDKNLRAFRL

MTB6 GEKHKKVLSGRKLWKLIIDVFLRCLVVVGGKAFCMVYSNVANVKGWLTCQVDKNLRAFRL

MTB7 GEKHKKVLSGRKLWKLIIDVFLRCLVVVGGKAFCMVYSNVANVKGWLTCQVDKNLRAFRL

MTB2 GEKHKKVLSGRKLWKLIIDVFLRCLVVVGGKAFCMVYSNVANVKGWLTCQVDKNLRAFRL

MTB5 GEKHKKVLSGRKLWKLIIDVFLRCLVVVGGKAFCMVYSNVANVKGWLTCQVDKNLRAFRL

MTB4 GEKHKKVLSGRKLWKLIIDVFLRCLVVVGGKAFCMVYSNVANVKGWLTCQVDKNLRAFRL

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MTB6 FYMTLCIHAIFNMISAVFFTLMLTHFSFASWTAVAQDSLTSQGSDGGVEFSFFVDILAFK

MTB7 FYMTLCIHAIFNMISAVFFTLMLTHFSFASWTAVAQDSLTSQGSDGGVEFSFFVDILAFK

MTB2 FYMTLCIHAIFNMISAVFFTLMLTHFSFASWTAVAQDSLTSQGSDGGVEFSFFVDILAFK

MTB5 FYMTLCIHAIFNMISAVFFTLMLTHFSFASWTAVAQDSLTSQGSDGGVEFSFFVDILAFK

MTB4 FYMTLCIHAIFNMISAVFFTLMLTHFSFASWTAVAQDSLTSQGSDGGVEFSFFVDILAFK

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MTB6 FLMSLICFLNCLHIQQLISACKSPNLPGPQNPVSHRLQSPSTPSSSASPADAVERDACKV

MTB7 FLMSLICFLNCLHIQQLISACKSPNLPGPQNPVSHRLQSPSTPSSSASPADAVERDACKV

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MTB4 FLMSLICFLNCLHIQQLISACKSPNLPGPQNPVSHRLQSPSTPSSSASPADAVERDACKV

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MTB3 SYFENTPNAAEKTTPTAVTLASYRQQETSGTGSQNLIEGRPRRKKPSKLSLLLKRDSGQK

MTB6 SYFENTPNAAEKTTPTAVTLASYRQQETSGTGSQNLIEGRPRRKKPSKLSLLLKRDSGQK

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MTB2 SYFENTPNAAEKTTPTAVTLASYRQQETSGTGSQNLIEGRPRRKKPSKLSLLLKRDSGQK

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MTB4 SYFENTPNAAEKTTPTAVTLASYRQQETSGTGSQNLIEGRPRRKKPSKLSLLLKRDSGQK

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MTB3 SGSSLSSRQQETPSPNLPSYSPNLYPSQAYI

MTB6 SGSSLSSRQQETPSPNLPSYSPNLYPSQAYI

MTB7 SGSSLSSRQQETPSPNLPSYSPNLYPSQAYI

MTB2 SGSSLSSRQQETPSPNLPSYSPNLYPSQAYI

MTB5 SGSSLSSRQQETPSPNLPSYSPNLYPSQACI

MTB4 SGSSLSSRQQETPSPNLPSYSPNLYPSQAYI

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