

NTDB id	1029	TT RS02230 WP 011228203.1	MPVYQYKARDPQGRLLVEATTEAEDLRATAARLRLDRGLFVAETKEPGKGLQAEVRIPALER	60
NTDB id	1169	A1552VC RS11075 WP 000648511.1	MKATQTLP LKNYRWKGINSNGKKVSGQMLAISEIEVRDKLKDQHITIKKLKKSVSLLAR....LTH	63
NTDB id	1402	DSB67 RS12670 WP 010643257.1	MKTIA PQLKNYRWKGINSNGKK.TSGNVLAMTEIEVRERLDAQHIKILKKLKGSISFITK....ISH	62
NTDB id	1112	NAGFG RS09215 WP 003689811.1	MAKNGGFSLF AKKEKR FIFEGRHSASD KLVNGEVS AFTEEEARKK LAKRGIRPLQITRVK.....T....SSKR	65
NTDB id	1113	AAA85695.1 219..1451()	MAKNGGFSLF AKKEKR FIFEGRHSASD KLVNGEVS AFTEEEARKK LAKRGIRPLQITRVK.....T....SSKR	65
NTDB id	1134928	ACLVWC RS05190 WP 039525808.1	MSVARNAVNKPQPSRN TSLQQPFVMEGT D KRGVK.MKGEQTARNAN MLRAELRRQG ITPTV VAKPK PLFGS....AGK	74
NTDB id	1252	GC085 RS07730 WP 011213805.1	MDKN SPPI LTFHYCGINKAGQK.MEGDIQARS LAIAKAD LRKQGI VTNKVVKRK PLFDR....KNK	62
NTDB id	1198	PSJM300 03950 AFN76868.1	MAQKA IKNSVF TWEGLD RQGAI.LKGELSGVSPAL VKAQ LRKQGIN PQKVRK KSISLFS.....AGK	61
NTDB id	1016	ACIAD RS01680 WP 004920476.1	MAAKKTQVM PVFAYEGVD RKGIK.LKGE LPARNMAL AKVT LRKQG ITIRT IREKR KNIFEG...LLKK	64
NTDB id	1059	ABD1 RS01610 WP 000279215.1	MAVKKA QMMPTFA YEGVD RKGVK.LKGE LPAKNMAL AKVT LRKQGV TRVNR IREKR KNILEG...LFKK	64
consensus			* * * * *	

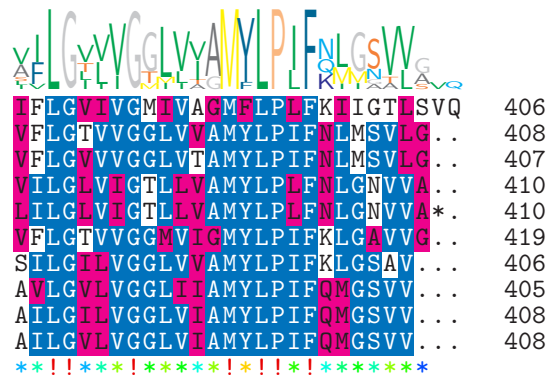
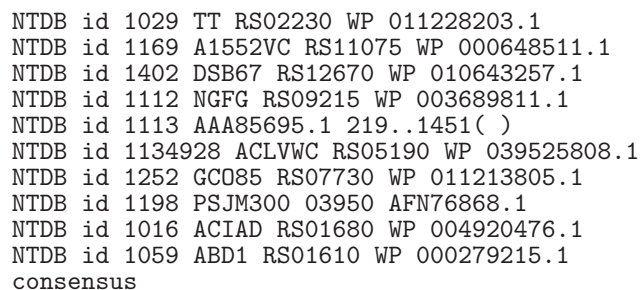
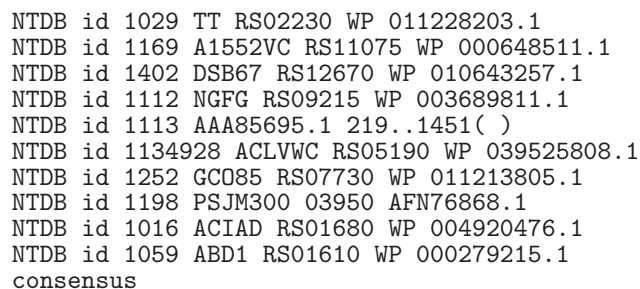
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NTDB id	1029	TT RS02230 WP 011228203.1	TSGGLDLILDRLASFLKELELRGKIRSAMTYPVIVFVFAVGVAFLLTGTVPQFAQLITDLCSELPLLTRFLIAVSDDL	219
NTDB id	1169	A1552VC RS11075 WP 000648511.1	MSCNLPEVFERLATYREKSEQLRAKVIKALIYPSMVVLVALGVSYLMLTMVTIPEFESMFKGFGAELPWFTQQVKLSHWV	223
NTDB id	1402	DSBG RS12670 WP 010643257.1	QSGNLSQVFERLATYREKSEQLRAKVIKALIYPMVVLVALGVSYLMLTRVTIPEFEKMFTGFGADLPFTAQTQVMVNLNSAT	222
NTDB id	1112	NCGF RS09215 WP 003689811.1	TGGVLESLLDKLAITYEKTQAIRKKVKTALTYPVSVIAVAIGLVFVMMIFVLPAFKEYVANMGAELPPLTQTVDMSDF	225
NTDB id	1113	AAA85695.1 219..1451()	TGGVLESLLDKLAITYEKTQAIRKKVKTALTYPVSVIAVAIGLVFVMMIFVLPAFKEYVANMGAELPALTQTVDMSDF	225
NTDB id	1134928	ACLVWC RS05190 WP 039525808.1	GAGVLETVLDTVATYKENIEALKGKIKKALFYPPAMVVAVAITVSAILLIFVVPQFEVFKSFGAELPAFTQIIIVNASRFM	234
NTDB id	1252	GC085 RS07730 WP 011213805.1	KSGSLDIMLDKVATYKEKTEIKKKIKKALTYPVIAVMVALLVTAGLLIYVVPQFESLFKGFGDLPAMTRAVITMSEFM	222
NTDB id	1198	PSJM300 03950 AFN76868.1	QSGSLETILDRVATYKEKTEALKAKIKKANYPVAVVLVAVIVTALLIKVVPQFDQVFANFGAELPAFTLMVIGLSEAL	221
NTDB id	1016	ACIAD RS01680 WP 004920476.1	QSGSLETIMLDRAVAYKEKSEVLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFKDLFTSFEGADLPFTAQTLVNMNSWM	224
NTDB id	1059	ABD1 RS01610 WP 000279215.1	QSGALETMLDRVATYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFKDLFTSFEGADLPFTAQTMVNMSKWM	224
consensus			* ! * ! * ! * ! * ! * * * * * * * * * * * * * * * * ! * * * * * ! * * * * * ! * ! * * *	

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NTDB id 1029 TT RS02230 WP 011228203.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
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NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus

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- X non conserved
- X similar
- X > 50% conserved