

logo

NTDB id 1029 TT RS02230 WP 011228203.1	MPVYQYKARDHQGRL..VEATTEAEDLRTAARLLRDRGTFVAETKEPFGKIQAEVRIPALER	60
NTDB id 1402 DSB67 RS12670 WP 010643257.1	MKTIA PQLKNYRWKGINSSGKK.TSGNVLAMTEIEVRERLDAQHIKIKKKLKKGSISFITK....ISH	62
NTDB id 1169 A1552VC RS11075 WP 000648511.1	MKAQTLP LKKNYRWKGINSGKK.VSGQMLATSEIEVRDKLQDQHIQIKKKLKKGSVSLAR....LTH	63
NTDB id 1112 NGFG RS09215 WP 003689811.1	MAKNGGFSLF AKKEKRFIFEGRHSASDKLVNGEVSATFEEDARKKLAKRCIRPLQITRVK.....T....SSKR	65
NTDB id 1113 AAA85695.1 219..1451()	MAKNGGFSLF AKKEKRFIFEGRHSASDKLVNGEVSATFEEDARKKLAKRCIRPLQITRVK.....T....SSKR	65
NTDB id 997940 ABFB11 RS17745 WP 049429912.1	MSVSRSAIKKEPVARATMELQPFVVEGTDKRGVK.MKGEQLAKNANILRABLRQGINPGQVKPKPKPLFGA....AGS	74
NTDB id 1252 GC085 RS07730 WP 011213805.1	MDKNSPPLLTFHYGGINAKGQK.MEGDIQARSLAIAKADLRKQGVITNKVVKRKPPLFDR....KNK	62
NTDB id 1198 PSJM300 03950 AFN76868.1	MAQKA IKNVSFTVEGLDRQGAK..IKGELSGVSPALVMAQLRKQGINPQKVRKKSISLFS.....AGK	61
NTDB id 1016 ACIAD RS01680 WP 004920476.1	MAAKKTQVMPVFAYEGVDRKGIK..IKGELPARNMALAKVTLRKQGITIRTIKPKKNIFEG....LLKK	64
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MAVKKAQMPPTFAYEGVDRKGIK..IKGELPAKNMALAKVTLRKQGVTVRNIREKKNILEG....LFKK	64
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NTDB id 1029 TT RS02230 WP 011228203.1	GPGLKDLAIFSRQLATMIGACITLLQALAILERQTENRKFRRLKQVHTDVEGCMAFSEALSCHK.IFSRLYVNLVRAGE	139
NTDB id 1402 DSB67 RS12670 WP 010643257.1	RVKGGK DITIFTRQIATMLMTGVPIVQALKLVSENHKKAEMKSLMSVTRAVEACTPMSKAMRTASTHFDPLYTDLIATGE	142
NTDB id 1169 A1552VC RS11075 WP 000648511.1	RVKSKDITILTRQLATMITTGVPIVQALKLVGDNRHKAEMKSLAQITKSVBACTPLSKAMRTASAHFDPLYVDLVEITGE	143
NTDB id 1112 NGFG RS09215 WP 003689811.1	KITQEDITV FTRQLSTMIAKAGIPLMQAFEIVARGHGNPSMTEMLMEIRGQVEQCSSLRAFSNHPKYFDRFYCNLVAAAGE	145
NTDB id 1113 AAA85695.1 219..1451()	KITQEDITV FTRQLSTMIAKAGIPLMQAFEIVARGHGNPSMTEMLMEIRGQVEQCSSLRAFSNHPKYFDRFYCNLVAAAGE	145
NTDB id 997940 ABFB11 RS17745 WP 049429912.1	PVKPKDIAFFSRQMATMMKSGVPIVSALDITASGHKNPRMKKLVDITRTDIEGCSMYYAVSKHPVQFDELYRNLVRAGE	154
NTDB id 1252 GC085 RS07730 WP 011213805.1	KITQADITVFSRQLATMIESGIPLVQAFDIVAKGQSNKRLKDLTETIKHDIETGLTLAESLIKHPLYFNEFLCNLVDAGE	142
NTDB id 1198 PSJM300 03950 AFN76868.1	KIKPMDIALFTRQMATMMKAGVPLIQSFDIIGEGFDNPNMRKLVDLQKQEVAAACNSFASSLRKPKQYFDDLYCNLVDSGE	141
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NTDB id 1059 ABD1 RS01610 WP 000279215.1	KVTTLDITIFTRQLATMMKAGVPLVQCFEIVABGLENPAMREVVLGKGEVEGCSTFASALRKYPQHFDNLFCSLVESGE	144
consensus	*****	

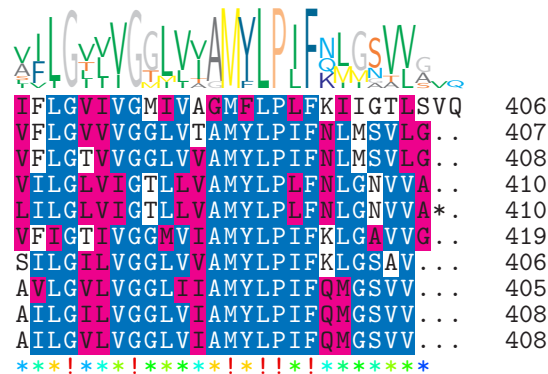
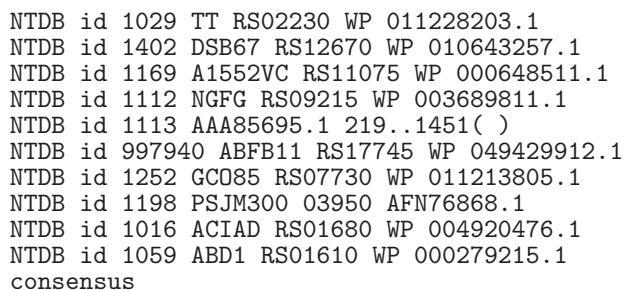
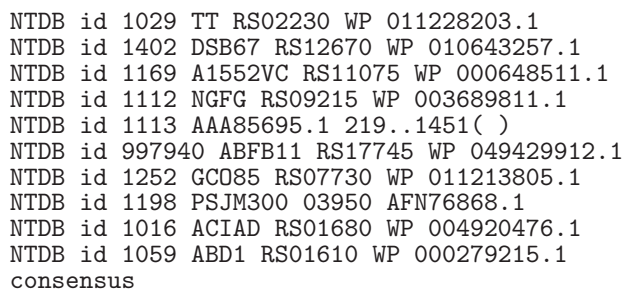
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NTDB id 1029 TT RS02230 WP 011228203.1	TSGGLDLILDRLASFLEKELELRGKIRSAMTYPVIVFVFVGVAFLLTGIVPQFAQLTLDLIGSELPLLTRFLIAVSDLL	219
NTDB id 1402 DSB67 RS12670 WP 010643257.1	QSGNLSQVFERLATYREKSEQLRKAVIKALIYPAMVVLVALCVSYLMLTRVIPLEFEKMTGFGADLPFAQTQVNLNSAWT	222
NTDB id 1169 A1552VC RS11075 WP 000648511.1	MSGNLPEVFERLATYREKSEQLRKAVIKALIYPAMVVLVALCVSYLMLTMVIPLEFESMFKGFGAELPWFTQQVLKLSHWV	223
NTDB id 1112 NGFG RS09215 WP 003689811.1	TGGVLESLLDKLATYKEKTQAIRKKVKLTALTYPVSVIATAICGVFVMMIFVLPFAFKEVYANMGAEPLPPTQTQVMDMSDF	225
NTDB id 1113 AAA85695.1 219..1451()	TGGVLESLLDKLATYKEKTQAIRKKVKLTALTYPVSVIATAICGVFVMMIFVLPFAFKEVYANMGAEPLPPTQTQVMDMSDF	225
NTDB id 997940 ABFB11 RS17745 WP 049429912.1	GAGVLETVLDITVATYKENIEALKGKIKKALFYPIPMVVVVANLVSGIMLVFVVPQFEDVFKSFGAELPAFTQLVVNLSRFM	234
NTDB id 1252 GC085 RS07730 WP 011213805.1	KSGSLDIMLDKVATYKEKIEIKKKIKKALTYPVAVMVVALLVITAGLLIYVVPQFESLFGKFGADLPAMTRAVITMSEFM	222
NTDB id 1198 PSJM300 03950 AFN76868.1	QSGSLETILDRVATYKEKTEALKAKIKKANYPVAVVIVAVIVTATILLIKVVPQFQDVFAANFGAELPAFTLVVIGLSEAL	221
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NTDB id 1059 ABD1 RS01610 WP 000279215.1	QSGALETMLDRVAVYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVQDLFASFGADLPFAFTQMVVNMSKWM	224
consensus	*****	

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NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus

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