

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQRL	VEATTEAEDLRTAARLRDRGLFVAETKEPGKGLQAEVRIPALERGPGLKD	66
NTDB id 1402	DSB67 RS12670 WP 010643257.1	MKTIAPQIKNYRWKGINSSGKK	TSGNVLAMTEIEVRRIDAQHIKIKKLLKKSISFITKI	68
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKATQTLPLKNYRWKGINSGKK	VSGQMLAISEIEVRDKLKDQHIQIKLLKKSVSLLARL	69
NTDB id 1112	NGFG RS09215 WP 003689811.1	MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVS	AFTEEEARKKLAKRGIRPLQITRVKT	71
NTDB id 1113	AAA85695.1 219..1451()	MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVS	AFTEEEARKKLAKRGIRPLQITRVKT	71
NTDB id 1252	GC085 RS07730 WP 011213805.1	MDKNSPPLTFHYQGINKAGQK	MEGDIQARSLATAKADLRKQGVITNKVVKRKP	68
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKKTQVMPVFAYEGVDRKGIK	LKGELPARNMAKAVTLRKQGITTIRIREKRKNIFEG	70
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKKAQMMPTFAYEGVDRKGIVK	IKGELPAKNMAKAVTLRKQGVTVRNIREKRKNILEGL	70
NTDB id 998046	ABDX87 RS17565 WP 346829021.1	MATKAVKTPKVAKPVVYVMEGVDKGA	KVTGELTGHSPALIKAQLRKQGVNPLKVRKKSVSIFSA	73
NTDB id 1198	PSJM300 03950 AFN76868.1	MAQKAIKNSVFTWEGLDKQGA	IKGELSGVSPALVKAQLRKQGINPQKVRKKSISIFSA	67
consensus		* * * * *	* * * * *	

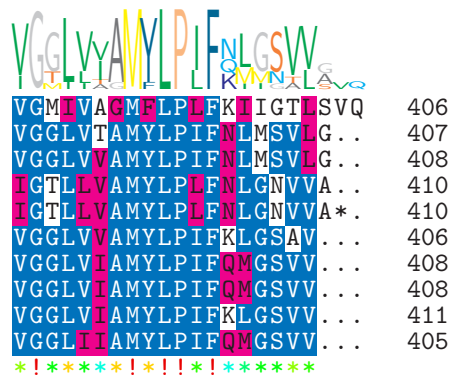
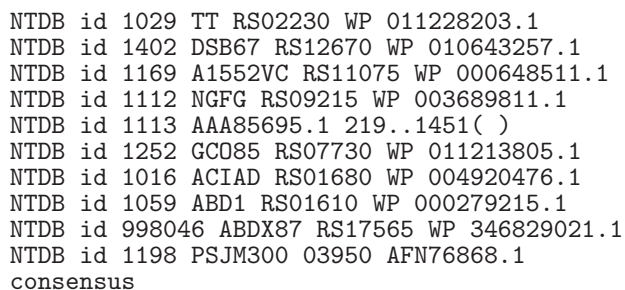
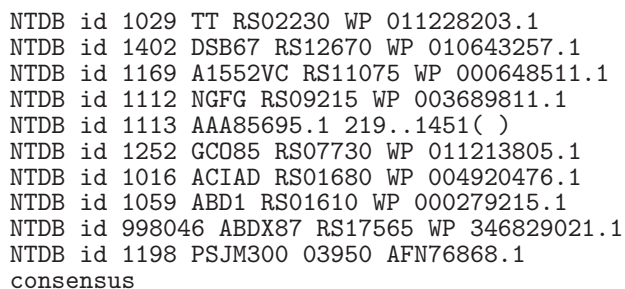
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NTDB id 1402	DSB67 RS12670 WP 010643257.1	QVFERLATYREKSEQLRAKVIKALIYPAMVVLVALGVSYLMLTRVIPEFEKMTFTGFGADLPFTQMVNLNSAWTQNWGP	228
NTDB id 1169	A1552VC RS11075 WP 000648511.1	EVFERLATYREKSEQLRAKVIKALIYPSMVVLVALGVSYLMITMVIPEFESMFKGFGAELPWFQTQQVILKLSHWVQAYSLW	229
NTDB id 1112	NGFG RS09215 WP 003689811.1	SLLDKLATYKEKTAIRKKVKKTALTYPVSVIAVAIGLVFVMMIFVLPAFKEVYANMGAELPPLTQTVMDSDFVSYGWM	231
NTDB id 1113	AAA85695.1 219..1451()	SLLDKLATYKEKTAIRKKVKKTALTYPVSVIAVAIGLVFVMMIFVLPAFKEVYANMGAELPPLTQTVMDSDFVSYGWM	231
NTDB id 1252	GC085 RS07730 WP 011213805.1	IMLDKVATYKEKTEIKKKIKKALTYPVIMVVALVLTAGLLIVVVPQFESLFGKFGADLPAMTRAVITMSEFMQAYWYI	228
NTDB id 1016	ACIAD RS01680 WP 004920476.1	IMLDRVATYKEKSEVLKQKIKKAMKYPATVIVVAIVVTIILMVKKVVPVFKDLFTSFGADLPFTQLVNVMSNMQDYWFI	230
NTDB id 1059	ABD1 RS01610 WP 000279215.1	IMLDRVATYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKKVVPVFKDLFTSFGADLPFTQMVVNMSKWMQDYWFI	230
NTDB id 998046	ABDX87 RS17565 WP 346829021.1	SLLDRVATYKEKTESLKAKIKKAMTYPVAVVIVAFIVTGILLIKVVPQFQSLFAGFGAELPAFTVMVINLSQYLQDYWFI	233
NTDB id 1198	PSJM300 03950 AFN76868.1	TLLDRVATYKEKTEALKAKIKKANYPVAVVLVAIVVTAILLIKVVPQFQDVFAFGAELPAFTLMVIGLSEALQAWYV	227
consensus		*****LILDRLASFLEKELELRGKIRSAMTYPVIVFVFAVGVAIFLLTGIVPQFAQILTDLGSSELPLLTRFLIAVSDLLRAATLP*****	

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NTDB id 1029 TT RS02230 WP 011228203.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1252 GCO85 RS07730 WP 011213805.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 998046 ABDX87 RS17565 WP 346829021.1
NTDB id 1198 PSJM300 03950 AFN76868.1
consensus

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- non conserved
- similar
- >50% conserved