

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGRLVEATTEAEDLRTAARLLRDGRGFLVAETKEPGKGQLAEVRIPALERGPGLK	65
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKAT.....QTLPKNRYRWKGINSNGKKVSGQMLAISEIEVRDCLKDQHIIQIKKLKKGVSLLARL.....THRVKSK	68
NTDB id 1402	DSB67 RS12670 WP 010643257.1	.MKT.....IAPQLKNRYRWKGINSNGKK.TSGNVLAMTEIEVRERLDAHQHKIKKLKKGVSISFITKI.....SHRVKGGK	67
NTDB id 1112	NFGF RS09215 WP 003689811.1	.MAKNGGFSLFAKKEKRIFIEGRHSASDKLVNGEVSATFEETEARKKLAKRGIRPLQITRVK.....TS....SKRKITQE	70
NTDB id 1113	AAA85695.1 219..1451()	.MAKNGGFSLFAKKEKRIFIEGRHSASDKLVNGEVSATFEETEARKKLAKRGIRPLQITRVK.....TS....SKRKITQE	70
NTDB id 1252	GCO85 RS07730 WP 011213805.1	.MDK.....NSPPLTLTFHYQGINKAGQK.MEGDIQARSLATAKADLRKQGIIVTNKVVKRRKP.LFDRK....NKKITQA	67
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAK.....KTQVMVPVFAYEGVDKGGIK.LKGELPARNMALAKVTLRKQGITTIRTIREKRKNIFEGL....LKKKVKPL	69
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVK.....KAQMMPPTFAYEGVDKGGVK.IKGELPAKNMALAKVTLRKQGVTVRNIREKRKNILEGL....FKKKVTTL	69
NTDB id 973784	AABM54 RS22070 WP 347902073.1	.MAV.....KAVKMSVYAMEGIDKKGKT.MTGELSCHNPALIKAKLRKQGINPKGVRKKKSASLLRA.....GQRKAH	66
NTDB id 1198	PSJM300 03950 AFN76868.1	.MAQ.....KAIKNSVFTFWGLDRQQAK.IKGELSGVSPALVKAQLRKQGINPKVRKKKSISLFSA.....GKKIKPM	66
consensus		***** * * **** * * * * *	

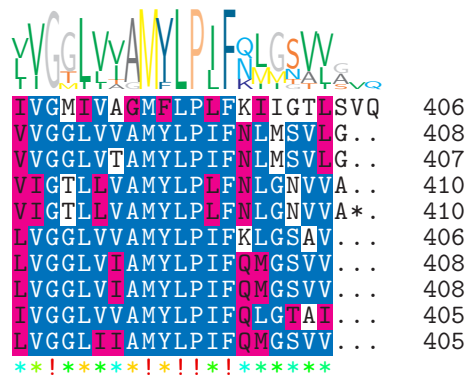
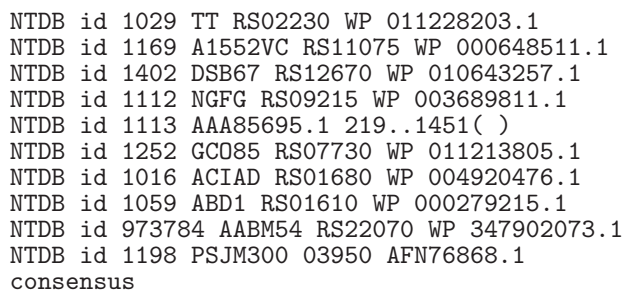
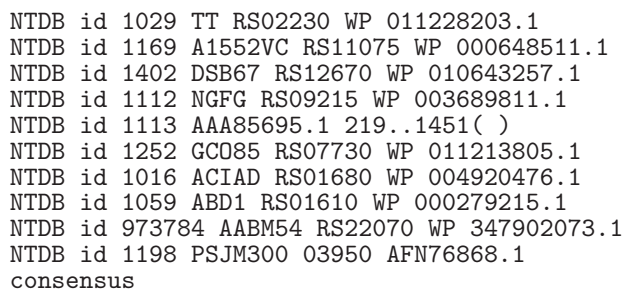
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NTDB id	1169	A1552VC RS11075 WP 000648511.1	PEVFERLATYREKSEQLRAKVIKALIYPSMVVLVALGVSYLMITMVIPEFESMFKGFGAELPWFTQQVLKLSHWVQAYSL	228
NTDB id	1402	DSB67 RS12670 WP 010643257.1	SQVFERLATYREKSEQLRAKVIKALIYPAMVVLVALGVSYLMLTRVIPEFEKMFTGFAGDLPFTAQMVLINLSANTQNNGP	227
NTDB id	1112	NGFG RS09215 WP 003689811.1	ESLCLKLATIYEKETQAIRKKVKKTALTYPVSVIAVAIGLVFVMIMIFVLPAFKEYVANMGALPPLTQTVMDMSDFFVSYGW	230
NTDB id	1113	AAA85695.1 219..1451()	ESLCLKLATIYEKETQAIRKKVKKTALTYPVSVIAVAIGLVFVMIMIFVLPAFKEYVANMGALPPLTQTVMDMSDFFVSYGW	230
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NTDB id	1016	ACIAD RS01680 WP 004920476.1	EIMLDRVAVIYEKSEVLKQIKKKAMKYPATVIVVAIVVTIIILMVKVVPVFQDLFTSFGADLPFTAQLVNMSNWMQDYWF	229
NTDB id	1059	ABD1 RS01610 WP 000279215.1	ETIMLDRVAVIYEKSELLKQIKKKAMKYPATVIVVAIVVTIIILMVKVVPVFQDLFASFGADLPFTAQMVVNMSKWMQEYWF	229
NTDB id	973784	AABM54 RS22070 WP 347902073.1	DTILERVATYKEKSESLKAKIKKKAMTYPLAVFVAVVVTGILLIKVVPQFQSVFAGFGAQLPVFTLMVIALSEFLTQTMWW	226
NTDB id	1198	PSJM300 03950 AFN76868.1	ETILDRVATYKEKETALAKIKKKAMNYPIAVVLVAIVITAILLIKVVPQFDVFANFGAELPAFTLMVIGLSEALQAWY	226
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
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 973784 AABM54 RS22070 WP 347902073.1
NTDB id 1198 PSJM300 03950 AFN76868.1
consensus

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