

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGRL	VEATIEAEDLRTAARLLRDRGLFVAEIKPEPGKGLQAEVRIPALERGPGLK	65																
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKAT.....QTLPLKNYRWKGINSGKK	VSGQMLAISEIEVRDKLKDQHIQIKKLKKGVSLLAR.....LTHRVKSK	68																
NTDB id 1402	DSB67 RS12670 WP 010643257.1	.MKT.....IAPQLKNYRWKGINSSGKK	TSGNVLAMTEIEEVRERLDAQHIKIKKLKKGVSIFITK.....ISHRVKKG	67																
NTDB id 1112	NGFG RS09215 WP 003689811.1	.MAKNGGFSLF	AKKEKRFIFEGRHSASDKLVNCEVS	AFTEEEARKKLAKRGIRPLQITRV.....KTS	SSKRKITQE	70														
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NTDB id 1252	GC085 RS07730 WP 011213805.1	.MDKN.....SPPLLTFHYQGINKAGQK	MEGDIQARSLATAKADLRKQGI	VTNKVVKKRKL	LFDR.....KNKKITQA	67														
NTDB id 1198	PSJM300 03950 AFN76868.1	.MAQK.....AIKNSVFTWEGLD	RQGA	IKGELSGVSPALVKAQLRKQGIN	PQVRKKSISLFS.....AGKKIKPM	66														
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKK.....TQVMPVFAYEGV	DRKG	IKLKGELPARNMALAKVTLRKQGIT	IRTI	IREKRKN	IFEG....LLKKVK	KPL	69											
NTDB id 110209	ACAK30 RS01695 WP 000279216.1	MAVKK.....AQMMPTFAYEGV	DRKG	IKGELPAKNMALAKVTLRKQGV	TVRN	IREKRKN	ILEG....LFKKKV	TTL	69											
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKK.....AQMMPTFAYEGV	DRKG	IKGELPAKNMALAKVTLRKQGV	TVRN	IREKRKN	ILEG....LFKKKV	TTL	69											
consensus		***	*	*	*	*****	*	*	*	*	*	*****	!	*****	*	*	*****	*	*	*****

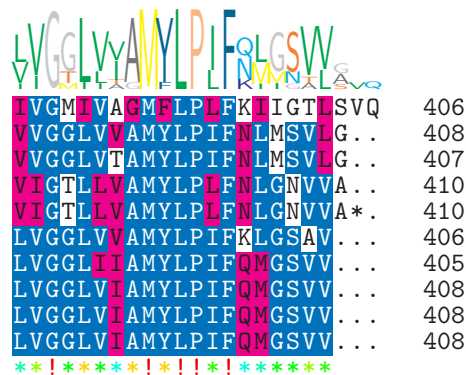
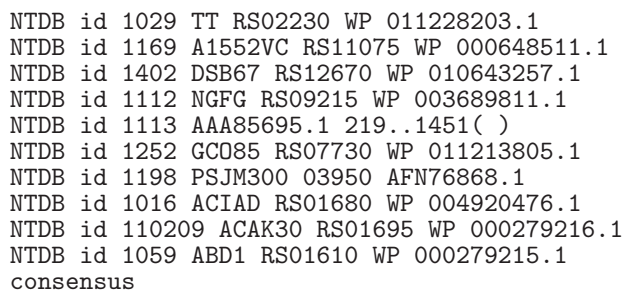
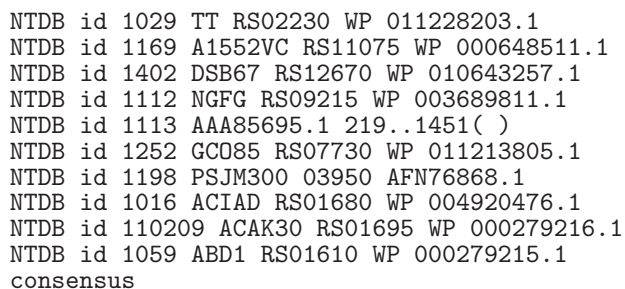
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NTDB id	1169 A1552VC RS11075 WP 000648511.1	DITILLTRQLATMTTTGVPIVQAALKLVGDNHRKAEMKSILAQITKSVEAGTPLSKAMRTASAHFDTLVVDL VETGEM SGNL	148
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NTDB id	1112 NGFG RS09215 WP 003689811.1	DITVFTRQLSTMIKAGLPLMQAFEIVARGHNPSMTEMLMEIRGQQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGETGGVL	150
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consensus		!*****!*!!**!*!!!!*!!***!!****!!*****!*****!*****!*****!*****!*****!	

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consensus			* * * * * ! * * ! ! * * * * ! * * * * ! * ! * ! * ! * * ! * * * * * * * * ! * ! * * * * * ! * * * * ! * * ! * * ! * * ! * * ! * * * * *	

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NTDB id 1029 TT RS02230 WP 011228203.1
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consensus

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- non conserved
- similar
- >50% conserved