

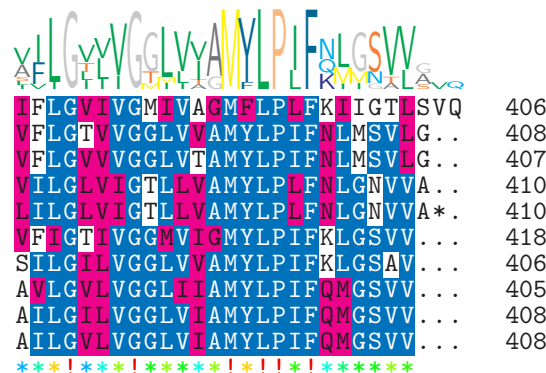
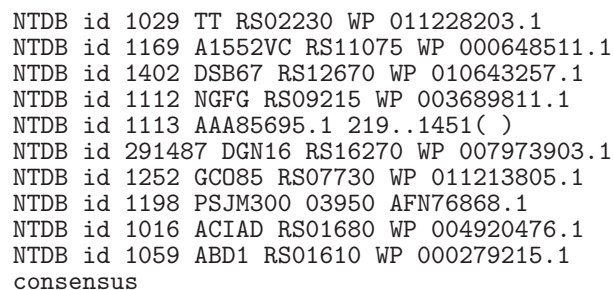
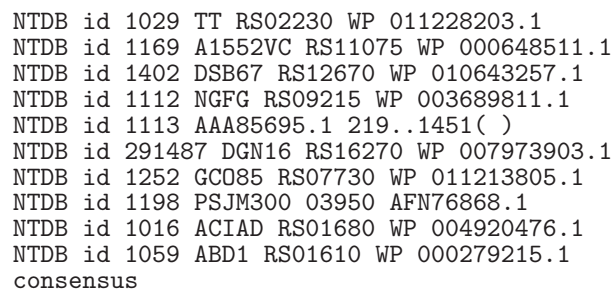
NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGR.L	VEATIEAEDLRTAARL	LRDRGLFVAET	IKPEPGKGLQAEVRIPALER	60
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKATQTLPLKKNYRWKGINSGKK	VSGQMLAISEIEVRDK	LKDQHIQIKK	LKKGVSLSLLAR....LTH	63
NTDB id 1402	DSB67 RS12670 WP 010643257.1	MKTIA	PQLKKNYRWKGINSSGKK	TSGNVLAMTEIEVRRLDAQH	IKKIKKKGKSISFITK....ISH	62
NTDB id 1112	NGFG RS09215 WP 00368911.1	MAKNGGFSLFAKKEKRF	FIEGGRHSASDKL	VNGEVS	AFTEEEARKKLAKRGIRPLQITRVK....T....SSKR	65
NTDB id 1113	AAA85695.1 219..1451()	MAKNGGFSLFAKKEKRF	FIEGGRHSASDKL	VNGEVS	AFTEEEARKKLAKRGIRPLQITRVK....T....SSKR	65
NTDB id 291487	DGN16 RS16270 WP 007973903.1	MSVARSSIIKKQPVDRNVSE	IPFVWEGTDKRGVK	MKGEQQARNANMLRAELRRQG	ITPTVVKTKPKPLFGA....AGK	74
NTDB id 1252	GC085 RS07730 WP 011213805.1	MDKN	SPPLTFHYQGINKAGQK	MEGDIQARSLAI	AKADLRKQGIVTNKKVKKRKPLFDR....KNK	62
NTDB id 1198	PSJM300 03950 AFN76868.1	MAQK	AIKNSVFTWEGLD	RQGAKIKGELSGVSPALVKAQ	LRKQGINPQKVRKKSISLFS....AGK	61
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKKTQV	MPVFAYEGVDRKG	IKLKGELPARNMALAKVT	LRKQGITTRTIREKRKNIFEG...LLKK	64
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKK	AQMPTFAYEGVDRKG	VVKLKGELPAKNMALAKVT	LRKQGVTVRNIIREKRKNILEG...LFKK	64
consensus			* * * * *	* * * * *	* * * * *	

[illegible][illegible]

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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 291487 DGN16 RS16270 WP 007973903.1
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