

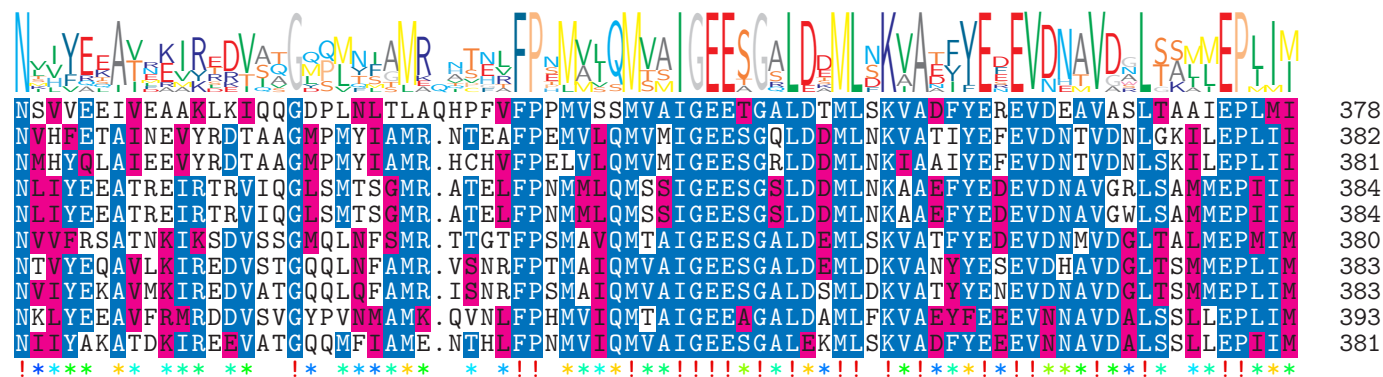
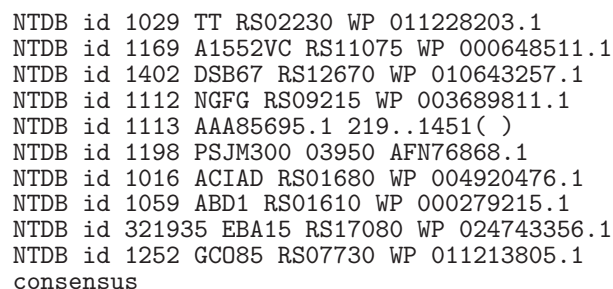
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NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKATQTLPLKNYRWKGINSGKK	VSGQMLAISEIEVRDKLKDQHIQIKKLKKGVSLSLLA....RLTH	63
NTDB id 1402	DSB67 RS12670 WP 010643257.1	MKTIAAPQLKNYRWKGINSSGKK	TSGNVLAMTEIEVRRLDAQHIKIKKLKKGKSISFIT....KISH	62
NTDB id 1112	NGAF RS09215 WP 003689811.1	MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVS	AFTEEEAARKKLAKRGIRPLQITRVK....T....SSKR	65
NTDB id 1113	AA85695.1 219..1451()	MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVS	AFTEEEAARKKLAKRGIRPLQITRVK....T....SSKR	65
NTDB id 1198	PSJM300 03950 AFN76868.1	MAQKAIAKNSVFTWEGLDLRQGAK	IKGELSGVSPALVKAQLRKQGINPQKVRKKSISLFS....AGK	61
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKKTQVMPVFAYEGVDRKGIK	LKGELPARNMALAKVTLRKQGITIRTIREKRKNIFEG...LLKK	64
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKKAQMMPTFAYEGVDRKGIVK	IKGELPAKNMALAKVTLRKQGVTVRNIREKRKNILEG...LFKK	64
NTDB id 321935	EBA15 RS17080 WP 024743356.1	MSVARSAIKKQPVDRNTMSLQTFVWEGADKRGVVK	MKGEGQARNANMLRAELRRQGIVPSMVKKQKPKPLLG....AAGK	74
NTDB id 1252	GC085 RS07730 WP 011213805.1	MDKNSPPLLTTHYQGINAKGQK	MEGDIQARSLAIKADLRKQGIVTNKVVKKRKPLFD....RKNK	62
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 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
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consensus

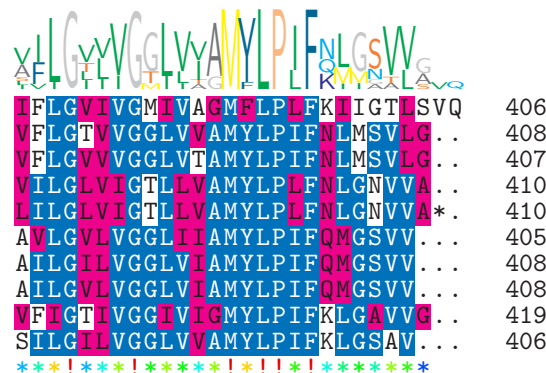
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NTDB id 1112 NGFG RS09215 WP 003689811.1
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NTDB id 1252 GC085 RS07730 WP 011213805.1
consensus

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