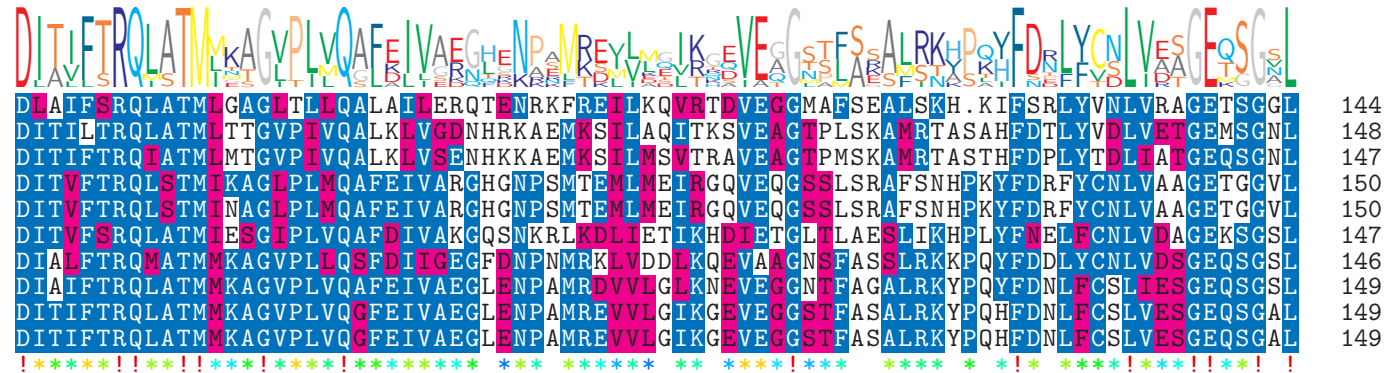


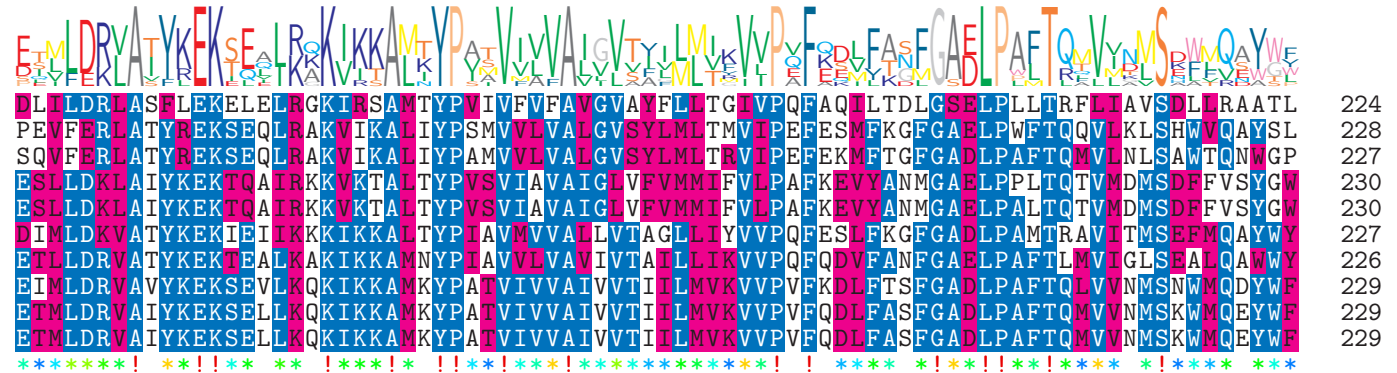
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NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1014153 ABWX03 RS17220 WP 000279216.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus

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NTDB id 1029 TT RS02230 WP 011228203.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
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NTDB id 1112 NGFG RS09215 WP 003689811.1
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NTDB id 1014153 ABWX03 RS17220 WP 000279216.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus
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NTDB id 1029	TT RS02230 WP 011228203.1	PLLLLAVALFFAYRW.YYGTIPQGRRV	IDRLKLRPLVFGNINRKTAVARFSRTLALLSSSGVNIVEALDITKGTAGNSVVE	303
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NTDB id 1252	GC085 RS07730 WP 011213805.1	IIIFGALGGVVVSFFYAKN	HSLEFAQTTIDRVMLKLPVIGPILEKAAIARFARTLSITFAAGLPLVEALKSVAGATGNNIYA	307
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NTDB id 1016	ACIAD RS01680 WP 004920476.1	ILILVIGAATAGFLEARKRSKR	FRDQLDRLVLRPLVIGDLVYKAI	309
NTDB id 1014153	ABWX03 RS17220 WP 000279216.1	IMIATIGAVIAAFLEAKKRSK	KFRDGLDKLALKLPIFGDLVYKAI	309
NTDB id 1059	ABD1 RS01610 WP 000279215.1	IMIATIGAVIAAFLEAKKRSK	KFRDGLDKLALKLPIFGDLVYKAI	309
consensus		*****	*****!*****!*	309

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NTDB id 1029	TT RS02230 WP 011228203.1	EIVEAAKLKIQQGDPLNLT	LTAQHPFVFPPMVSSMVAIGEETGALD	383
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NTDB id 1402	DSB67 RS12670 WP 010643257.1	LAIEEVYRD	TAAGMPMYIAMR.HCHVFPELVLMQVM	386
NTDB id 1112	NGFG RS09215 WP 003689811.1	EATREIRTRV	IQGLSMTSGMR.ATELFPNMMLQMSS	389
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NTDB id 1014153	ABWX03 RS17220 WP 000279216.1	KAVMKIRE	DVATGQQLQFAMR.VSNRFPSMAIQ	388
NTDB id 1059	ABD1 RS01610 WP 000279215.1	KAVMKIRE	DVATGQQLQFAMR.ISNRFPSMAIQ	388
consensus		***	*****!*****	388

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NTDB id 1059	ABD1 RS01610 WP 000279215.1	LVGGLV	IAMYLPIFQMGSV..	408
consensus		***!*****!*	*****	

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved