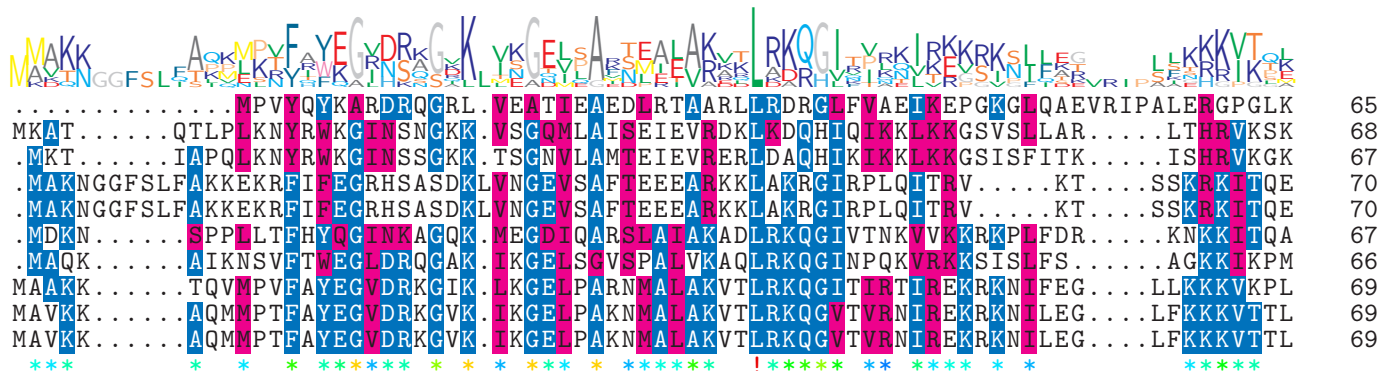


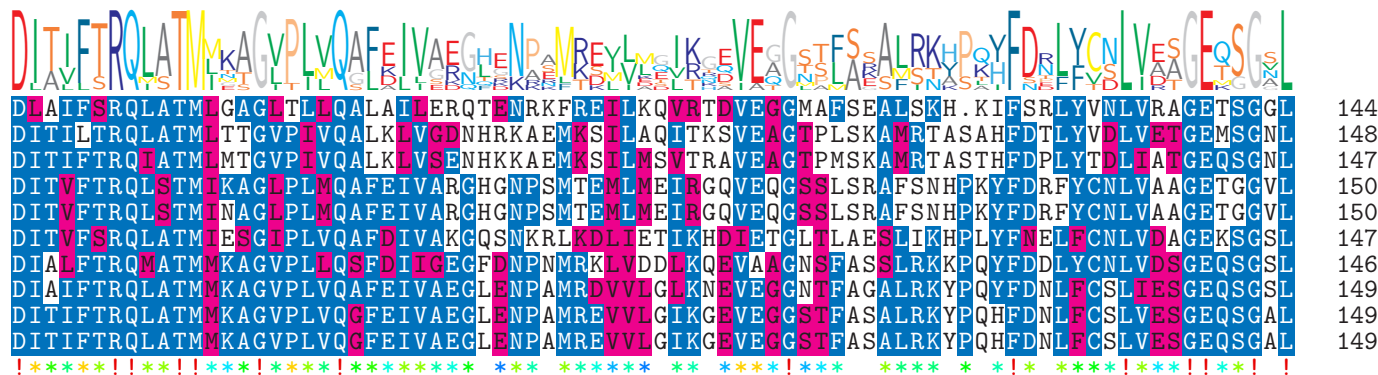
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NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
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NTDB id 1113 AAA85695.1 219..1451()
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
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NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus



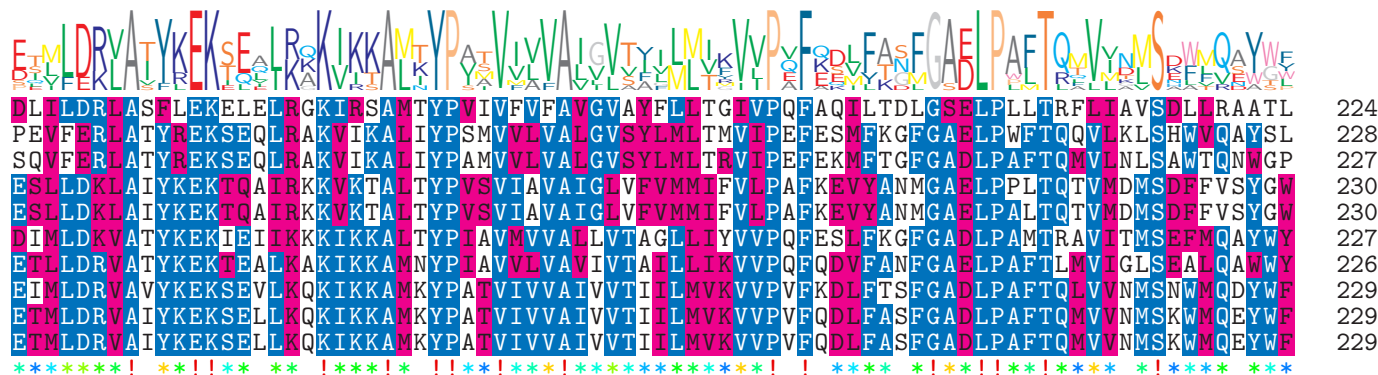
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NTDB id 1064226 ACG72M RS16805 WP 000279216.1
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consensus



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NTDB id 1064226 ACG72M RS16805 WP 000279216.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus



logo

NTDB id 1029	TT RS02230 WP 011228203.1	PLLLLAVALFFAYRW.YYGTIPQGRRV	IDRLKLRPLVFGNINRKTAVARFSRTLALLSSSGVNIVEALDITKGTAGNSVVE	303
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NTDB id 1402	DSB67 RS12670 WP 010643257.1	FIGLSFVSLISGKIMSQRSDS	FRLLAARSSLRVPILGPVLSKAAIAKFSRTLATSFSAGIPILTALKTT	307
NTDB id 1112	NGFG RS09215 WP 003689811.1	MVLIALGFATYGF	FLKLKARSIKIQRRMDAILLRMPIFGDIVRKGTIARWGRTTATLFAAGVPLVDVLD	310
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NTDB id 1059	ABD1 RS01610 WP 000279215.1	IMIATIGAVIAAFLEAKKRSKKFRDGLDKLAL	KLPIFGDLVYKAI	309
consensus		*****	*****!*****!*	309

logo

NTDB id 1029	TT RS02230 WP 011228203.1	EIVEAAKLKIQGGDP	NLTLAQHPFVFPMPVSSMVAIGEETGALD	383
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NTDB id 1402	DSB67 RS12670 WP 010643257.1	LAIEEVYRDTAAGMPMYIAMR	.HCHVFPELVLMQVMIGEESGRLDDMLNKIAAIYEF	386
NTDB id 1112	NGFG RS09215 WP 003689811.1	EATREIRTRVIQGLSMTSGMR	.ATELFPNMMLQMSSIGEESGSLDDMLNKAAEFY	389
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NTDB id 1252	GC085 RS07730 WP 011213805.1	KATDKIREEVATGQQMFIA	ME.NTHLFPNMVIQMV	386
NTDB id 1198	PSJM300 03950 AFN76868.1	SATNKKISDVSSGMQLNFSMR	.TTGTFPSPMAVQMTAIGEESGALDEMLS	385
NTDB id 1016	ACIAD RS01680 WP 004920476.1	QAVLKIREDEVSTGQQLNFAMR	.VSNRFPTMAIQMVAIGEESGALDEMLDKVAN	388
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consensus		***	*****!*****	388

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logo

NTDB id 1029	TT RS02230 WP 011228203.1	IVGMIVAGMFLPLFKIIGTISVQ	406
NTDB id 1169	A1552VC RS11075 WP 000648511.1	VVGGLVAMYLPIFNLMSVLG..	408
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NTDB id 1064226	ACG72M RS16805 WP 000279216.1	LVGGLVIAMYLPIFQMGSV..	408
NTDB id 1059	ABD1 RS01610 WP 000279215.1	LVGGLVIAMYLPIFQMGSV..	408
consensus		***!*****!***!*****	

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