

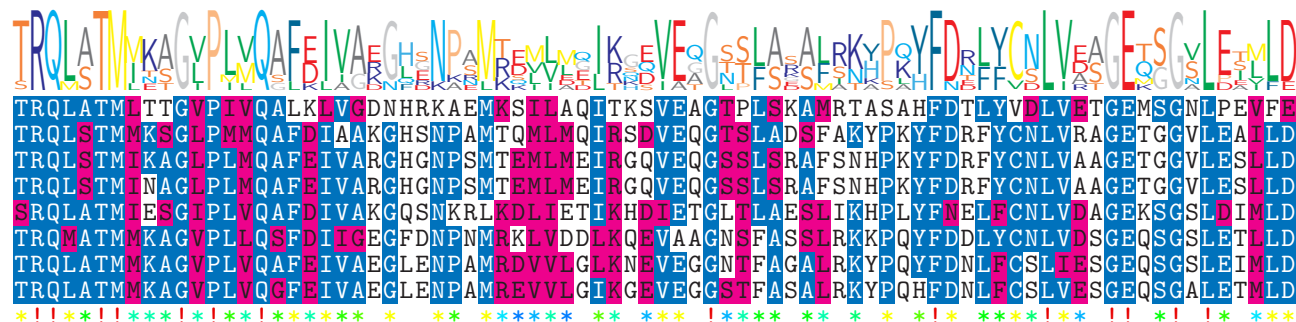
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NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1020495 ABU615 RS07820 WP 100156468.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451()
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus



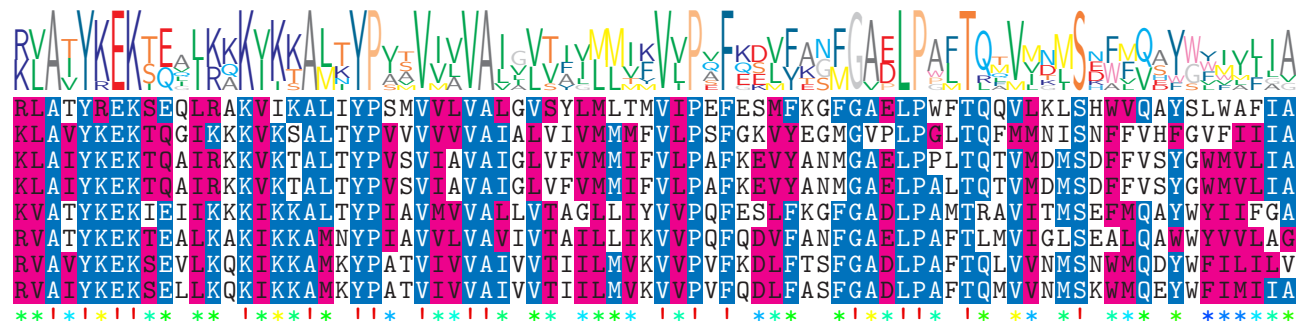
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NTDB id 1169 A1552VC RS11075 WP 000648511.1	VYRDTAAGMPMYIAMRNTEAFPEMVLQMVMIGEESGQLDDMLNKVATIYEFYEDVNTVDNLGKILEPLIIVFLGTVVGGLV	393
NTDB id 1020495 ABU615 RS07820 WP 100156468.1	IRAKVNQGISLTSAMQATELFPNMVMQMASIGEESGSLDDMLNKSAEFYEDVDTAVSQSLSLMEPIIMVVLGLIVGVIL	395
NTDB id 1112 NGFG RS09215 WP 003689811.1	IRTRVIQGLSMTSCMRATELFPNMMLQMSSIGEESGSLDDMLNKAAEFYEDVDNAVGRLSAMMEPIIIVILGLVIGTLL	395
NTDB id 1113 AAA85695.1 219..1451()	IRTRVIQGLSMTSCMRATELFPNMMLQMSSIGEESGSLDDMLNKAAEFYEDVDNAVGRLSAMMEPIIILILGLVIGTLL	395
NTDB id 1252 GC085 RS07730 WP 011213805.1	IREEVATGQQMFIAMENTHLFPNMVIQMVAIGEESGALDKMLSKVADFYEEDVNAVDALSSLEPIIMSILGILVGGLV	392
NTDB id 1198 PSJM300 03950 AFN76868.1	IKSDVSSGMQLNFSMRTTGTFPSMAVQMTAIGEESGALDEMLSKVATFYEDVDNMVDGLTALMEPMIMAVLGVLVGGLI	391
NTDB id 1016 ACIAD RS01680 WP 004920476.1	IREDEVSTGQQLNFMARVSNRFPTMAIQMVAIGEESGALDEMLDKVANYYESEVDHVDGLTSMMEPLIMAILGILVGGLV	394
NTDB id 1059 ABD1 RS01610 WP 000279215.1	IREDEVATGQQQLFAMRISNRFPSMAIQMVAIGEESGALDSMLDKVATYYENEVDNAVVDGLTSMMEPLIMAILGVLVGGLV	394
consensus	*** * !* * *!* * !! !*** !!!!! !*** !* *!*!***!* !* **!*!* *!*!***!***	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	VAMYLPIFNIMSVLG.	408
NTDB id 1020495 ABU615 RS07820 WP 100156468.1	VAMYLPLFNIGNAVG.	410
NTDB id 1112 NGFG RS09215 WP 003689811.1	VAMYLPLFNLGNVVA.	410
NTDB id 1113 AAA85695.1 219..1451()	VAMYLPLFNLGNVVA*	410
NTDB id 1252 GC085 RS07730 WP 011213805.1	VAMYLPIFKLGSAV..	406
NTDB id 1198 PSJM300 03950 AFN76868.1	IAMYLPIFQMGSVV..	405
NTDB id 1016 ACIAD RS01680 WP 004920476.1	IAMYLPIFQMGSVV..	408
NTDB id 1059 ABD1 RS01610 WP 000279215.1	IAMYLPIFQMGSVV..	408
consensus	*!!!!!!*!*****	

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