

logo

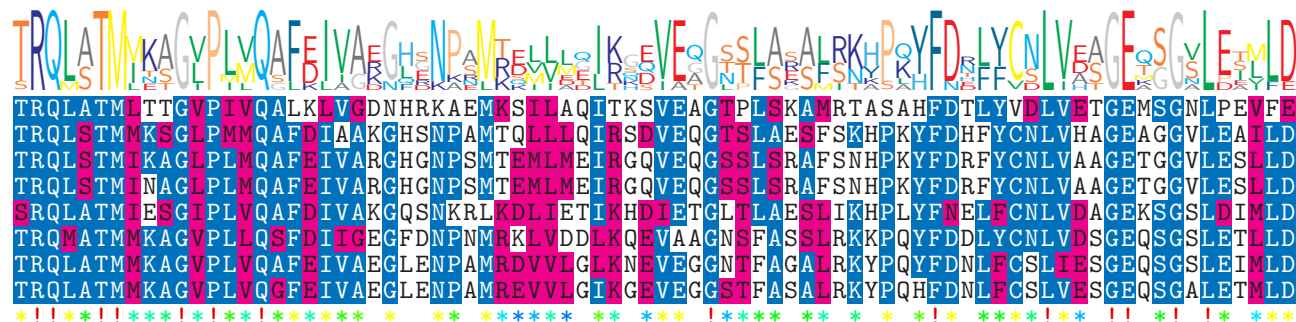
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NTDB id 1016697 ABW387 RS08970 WP 025329863.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451()
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus



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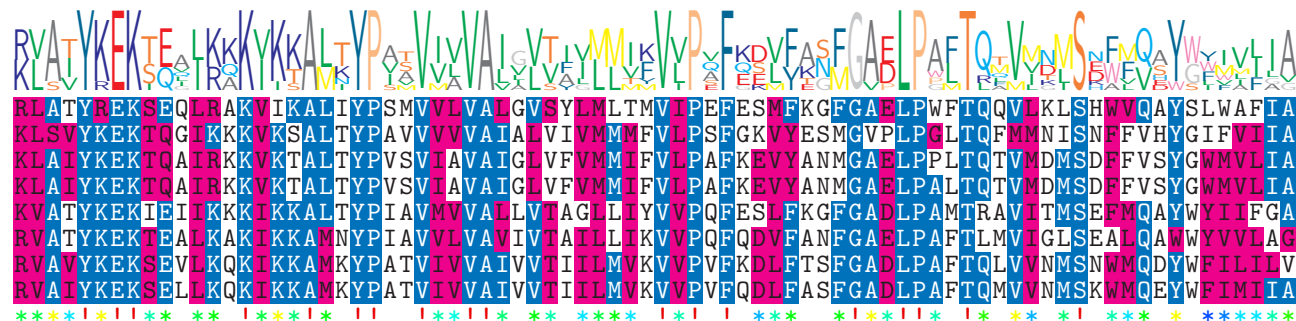
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NTDB id 1169 A1552VC RS11075 WP 000648511.1
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consensus



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NTDB id 1169 A1552VC RS11075 WP 000648511.1	VYRDTAAAGMPMYIAMRNTEAFPEMVLQMVMIGEESGQLDDMLNKVATIYEFYEVNDNTVDNLGKILEPLIIIVFLGTIVGGGLV	393
NTDB id 1016697 ABW387 RS08970 WP 025329863.1	IRAKVNQGISLTSAMQATELFPNMVMQMASIGEESGSLDDMLNKAAEFYEDVDTAVALSSLMETIIMVVLGGIVGTIL	395
NTDB id 1112 NGFG RS09215 WP 003689811.1	IRTRVIQGLSMTSCMRATELFPNMMLQMSSIGEESGSLDDMLNKAAEFYEDVDNAVGRLSAMMEPIIIVILGLVIGTLL	395
NTDB id 1113 AAA85695.1 219..1451()	IRTRVIQGLSMTSCMRATELFPNMMLQMSSIGEESGSLDDMLNKAAEFYEDVDNAVGWLSAMMEPIIILILGLVIGTLL	395
NTDB id 1252 GC085 RS07730 WP 011213805.1	IREEVATGQQMFIAMENTHLFPNMVIQMVAIGEESGALDKMLSKVADFYEVEVNAVDALSSLLEPTIMSILGILVGGGLV	392
NTDB id 1198 PSJM300 03950 AFN76868.1	IKSDVSSGMQLNFSMRTTGTFPSMAVQMTAIGEESGALDEMLSKVATFYEDVDNMVDGLTALMEPMIMAVLGVVLVGGGLI	391
NTDB id 1016 ACIAD RS01680 WP 004920476.1	IREDEVSTGQQLNFMARVSNRFPTMAIQMVAIGEESGALDEMLDKVANYYESEVDHVDGLTSMMEPLIMAILGILVGGGLV	394
NTDB id 1059 ABD1 RS01610 WP 000279215.1	IREDEVATGQQQLQFAMRISNRFPSMAIQMVAIGEESGALDSMLDKVATYYENEVDNAVDGLTSMMEPLIMAILGVVLVGGGLV	394
consensus	*** ** !* * *!* * !! !***! !!!!! !***! !* !*!***!***! !* ***!*** *!*!***!***	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	VAMYLPIFNIMSVLG.	408
NTDB id 1016697 ABW387 RS08970 WP 025329863.1	VAMYLPLFNIGNAVG.	410
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NTDB id 1113 AAA85695.1 219..1451()	VAMYLPLFNLGNVVA*	410
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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