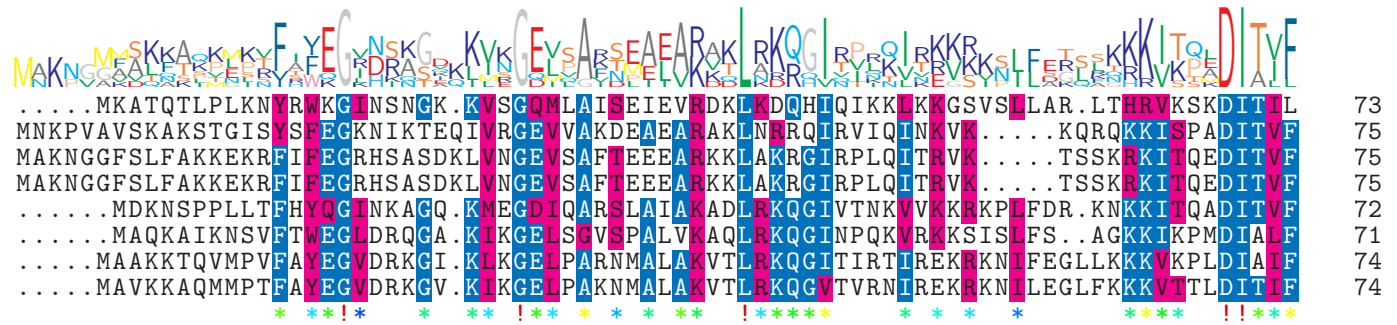


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NTDB id 1015659 ABV523 RS00650 WP 100156468.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus
```



logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1	VYRDTAAAGMPMYIAMRNTEAFPEMVLQMVMIGEESGQLDDMLNKVATIYEFYEDVNTVDNLGKILEPLIIVFLGTVVGGGLV	393
NTDB id 1015659 ABV523 RS00650 WP 100156468.1	IRAKVNQGISLTSAMQATELFPNMVMQMASIGEESGSLDDMLNKSAEFYEDVDTAVSQLSLMEPIIMVVLGLIVGVIL	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	IREEVATGQQMFIAMENTHLFPNMVIQMVAIGEESGALDKMLSKVADFYEEDVNAVDALSSLLEPTIMSILGILVGGGLV	392
NTDB id 1198 PSJM300 03950 AFN76868.1	IKSDVSSGMQLNFSMRTTGTFPSMAVQMTAIGEESGALDEMLSKVATFYEDVDNMVDGLTALMEPMIMAVLGVLVGGGLI	391
NTDB id 1016 ACIAD RS01680 WP 004920476.1	IREDEVSTGQQLNFMAMRVSNRFPTMAIQMVAIGEESGALDEMLDKVANYYEEDVHVDGLTSMMEPLIMAILGILVGGGLV	394
NTDB id 1059 ABD1 RS01610 WP 000279215.1	IREDEVATGQQQLQFAMRISNRFPSMAIQMVAIGEESGALDSMLDKVATYYENEVDNAVVDGLTSMMEPLIMAILGVLVGGGLV	394
consensus	*** ** !* * *!* * !! !*** !!!!! !*** !* !* !*!***!*** !* ***!*** *!***!***	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	VAMYLPIFNIMSVLG.	408
NTDB id 1015659 ABV523 RS00650 WP 100156468.1	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