

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGR.L	VEATIEAEDLR	TAAARL	LRDRGLFVAET	IKPEPGKGL	QAEVRIPALER	60
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKATQTLPL	LKNYRWKGIN	SNGKK.VSGQMLA	ISEIEVRDKL	KDQHIQIKK	LKKGVSLLAR.....LTH	63
NTDB id 1402	DSB67 RS12670 WP 010643257.1	MKTII	APQLKNYRWK	GINSSGKK.TSGNV	LAMTEIEVRER	LDAQHIKIKK	LKKGKSISFITK.....ISH	62
NTDB id 1112	NGFG RS09215 WP 003689811.1	MAKNGGFSLF	AKKEKRFIF	EGRHSASDKL	VNGEVS	AFTEEEARKKL	LAKRGIRPLQITRVK.....TS....SKR	65
NTDB id 1113	AAA85695.1 219..1451()	MAKNGGFSLF	AKKEKRFIF	EGRHSASDKL	VNGEVS	AFTEEEARKKL	LAKRGIRPLQITRVK.....TS....SKR	65
NTDB id 1198	PSJM300 03950 AFN76868.1	MAQKA	IKNSVFTW	EGLDRQGAK.IKGEL	SGVSPALVKAQ	LRKQGINPQK	VRKKSISLFS.....AGK	61
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKKTQV	MPVFAYEGV	DRKGIK.LKGEL	PARNMALAKVT	LRKQGITIRT	IREKRKNIFEGL....LKK	64
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKKA	QMMPTFAYEGV	DRKGVK.IKGEL	PAKNMALAKVT	LRKQGVTVRN	IREKRKNILEGL....FKK	64
NTDB id 1000862	ABFU15 RS05320 WP 012437601.1	MSVARSAIKKQPVDRNTSMLQT	FIWEGADKR	GVK.MKGEQT	ARNANMLRAEL	LRKQGI	VPSVVKQKPKPLFGA....AGK	74
NTDB id 1252	GCO85 RS07730 WP 011213805.1	MDKN	SPPLTFHYQG	INKAKQK.MEGDI	QARSLAI	IAKADLRKQGI	VTVNKKRKPLFDR.....KNK	62
consensus			* * * * *	* * * * *	* * * * *	* * * * *	* * * * *	

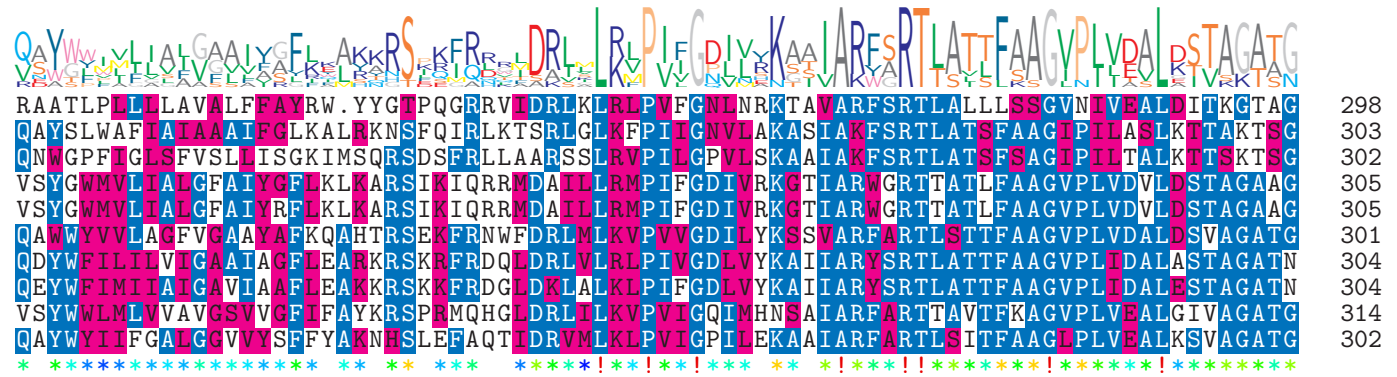
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NTDB id	1169	A1552VC RS11075 WP 000648511.1	RVKSKSDITILTTRQLATMLTGTGVPIVQAALKLVGDNRHKAEMKSILAQITKSVEAGTPLSKAMRTASAHFDTLVVDLETGE	143
NTDB id	1402	DSB67 RS12670 WP 010643257.1	RVKGKDITIFTTRQIATMIMTGVPINVQAALKLVSENHKKAEAKMSILMSVTRAWEAGTPMSKAMRTASTHFDPDYTDLIATGE	142
NTDB id	1112	NGFG RS09215 WP 003689811.1	KITQEIDITVFTRQLSTMIAKAGPLMQAFEIVARGHGPNPSMTLEMLEIRGQVEEQSSLSRAFSNHPKYFYDRFCYNLVAAGE	145
NTDB id	1113	AAA85695.1 219..1451()	KITQMEDIITVFTRQLSTMINAGLPLMQAFEIVARHGHNPSMTLEMLEIRGQVEEQSSLSRAFSNHPKYFYDRFCYNLVAAGE	145
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NTDB id	1016	ACIAD RS01680 WP 004920476.1	KVKPLDIAFITRQLATMMKAGVPLVQAFEIVAEGLENPAMRDVVVLGLKNVEVEGGNTFAAGLRKYPQFYDNLFCSLIESGE	144
NTDB id	1059	ABD1 RS01610 WP 000279215.1	KVTTLIDITIFTTRQLATMMKAGVPLVQGFEIVAEGLENPAMREVVLGIKEVEGGSTFASALRKYPQHFDNFLCSSLVESGE	144
NTDB id	1000862	ABFU15 RS05320 WP 012437601.1	KITPKDIATFFSRQMATMMKSGVPIVSSLEITIIEGHKNPRMKKMVGIRTDTIEGSSSLEYESTISAPVQFDELYRNLVAGE	154
NTDB id	1252	GCO85 RS07730 WP 011213805.1	KITQADIITVFSRQLATLMIESGIPLVQAQDIVAKGQSNKRLLKDIIETIKHDIIETGLTLAESLIIKHPLYNFELFCNLVDAGE	142
consensus			* * * * *	

[illegible]

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NTDB id 1029 TT RS02230 WP 011228203.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
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 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
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NTDB id 1252 GC085 RS07730 WP 011213805.1
consensus

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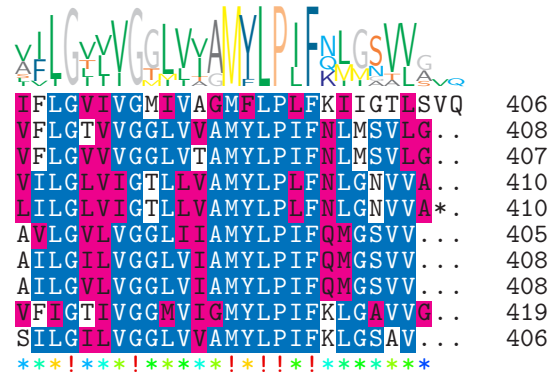
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NTDB id 1029 TT RS02230 WP 011228203.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NCFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 1000862 ABFU15 RS05320 WP 012437601.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
consensus

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 ≤ 0

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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 1000862 ABFU15 RS05320 WP 012437601.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
consensus
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- X non conserved
- X similar
- X $\geq 50\%$ conserved