

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPS	AVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPS	AVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...	FVAAEEVQSN	SLKSIDFR	TNKKDKDAVIVVELVSPA	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1					0
NTDB id 1373	NTHI	RS02645	WP 011272046.1					0
NTDB id 79562	HIY	RS03430	WP 005659796.1					0
NTDB id 1347	HI	0435	AAC22094.1					0
consensus								0

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	TLKGKYLEVVISK	..LKADEKPKPKSVLEKE	155
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	TLKGKYLEVVISK	..LKADEKPKPKSVLEKE	155
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLNNTSVKDDQLYLLDVKDFSTVSVESVEVFRDSTTTRLVAYIDEDYSHD	YRLTGRLGYLEVKGK	..LKANEKAPDKSILDKE	156
NTDB id 1381	A4U84	RS02010	WP 021113424.1		MRYLFLFFATF.....	P.....VLA	16
NTDB id 1373	NTHI	RS02645	WP 011272046.1		MKKYFLKCGYFLVCFCLP	..LIVFANP.....KTD	28
NTDB id 79562	HIY	RS03430	WP 005659796.1		MKKYFLKCGCFLVFFCLP	..LIVFANP.....KTD	28
NTDB id 1347	HI	0435	AAC22094.1		MKKYFLKCGYFLVCFCLP	..LIVFANP.....KTD	28
consensus					*	*	28

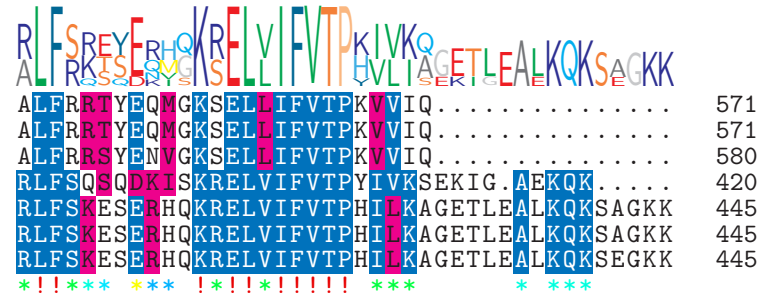
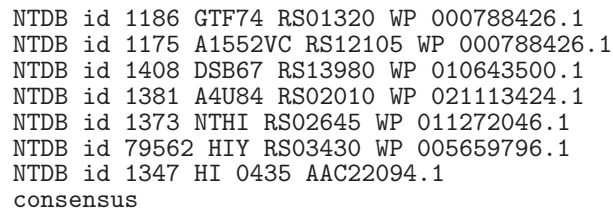
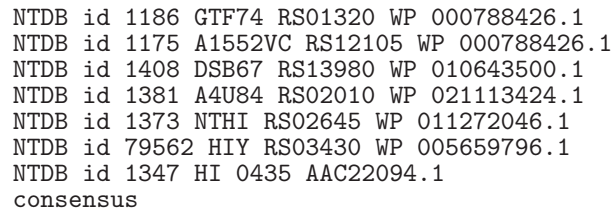
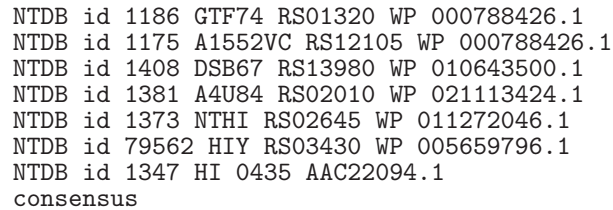
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	GKLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	LRLDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	235
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	GKLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	LRLDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	235
NTDB id 1408	DSB67	RS13980	WP 010643500.1	GKLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	LRLDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKDELD	236
NTDB id 1381	A4U84	RS02010	WP 021113424.1	NQKISLSIRNAPTAEIISYLAETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENG	IYYIHQAQEH...		93
NTDB id 1373	NTHI	RS02645	WP 011272046.1	NECFFIRLSQAPLAQTLEQLAFQQDVLNLMGERLEGNISLKLNNIDMPRL	LKTI	IAKSKH	105
NTDB id 79562	HIY	RS03430	WP 005659796.1	NERFFIRLSQAPLAQTVEQLAFQQDVLNLMGERLEGNISLKLNNIDMPRL	LKTI	IAKSKH	105
NTDB id 1347	HI	0435	AAC22094.1	NERFFIRLSQAPLAQTLEQLAFQQDVLNLMGERLEGNISLKLNNIDMPRL	LKTI	IAKSKH	105
consensus				*****	*****	*****	*****

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	EKQALEKARLAEELGDLKSEI	IKINF	AKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	315
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	EKQALEKARLAEELGDLKSEI	IKINF	AKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	315
NTDB id 1408	DSB67	RS13980	WP 010643500.1	EKQALEKARLAEELGDLKSEI	IKINF	AKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	316
NTDB id 1381	A4U84	RS02010	WP 021113424.1	..KQHTTAQLVNALPKLIT	KTIKLHYSKASEVIESLT	..KGQGNLLSES	GYLHFD	170
NTDB id 1373	NTHI	RS02645	WP 011272046.1	..KQQVAGNLT	TNNEPHLVSH	TVKLHF	AKASELMKSLT	182
NTDB id 79562	HIY	RS03430	WP 005659796.1	..KQVAGNLT	TNNEPHLVSH	TVKLHF	AKASELMKSLT	182
NTDB id 1347	HI	0435	AAC22094.1	..KQVAGNLT	TNNEPHLVSH	TVKLHF	AKASELMKSLT	182
consensus								182

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NTDB id 1186 GTF74 RS01320 WP 000788426.1
NTDB id 1175 A1552VC RS12105 WP 000788426.1
NTDB id 1408 DSB67 RS13980 WP 010643500.1
NTDB id 1381 A4U84 RS02010 WP 021113424.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
NTDB id 79562 HIY RS03430 WP 005659796.1
NTDB id 1347 HI 0435 AAC22094.1
consensus
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- ☐ non conserved
- ☒ similar
- ☒ $\geq 50\%$ conserved