

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQ	TLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AAVLIVELASPSAVVDVQKVQ	EGLSI	76	
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQ	TLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AAVLIVELASPSAVVDVQKVQ	EGLSI	76	
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...	FVAAEEVQSN	SLKSIDFR	TNKKDKDAVIVVELVSPA	AI	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1					0	
NTDB id 1347	HI 0435	AAC22094.1						0	
NTDB id 219973	BV139	RS02995	WP 005686897.1					0	
NTDB id 1373	NTHI	RS02645	WP 011272046.1					0	
consensus									

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	YTLKGYLEVVIS	KLKADEKPKPKSVLEKE	EG	156
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	YTLKGYLEVVIS	KLKADEKPKPKSVLEKE	EG	156
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLNNTSVKDDQLYLLDVKDFSTVSVESVEVFRDTSTTRLVAYIDEDYSHD	YRLTGRYLEVVKVGK	KLKAN	EPDKS	157
NTDB id 1381	A4U84	RS02010	WP 021113424.1					17
NTDB id 1347	HI 0435	AAC22094.1						29
NTDB id 219973	BV139	RS02995	WP 005686897.1					29
NTDB id 1373	NTHI	RS02645	WP 011272046.1					29
consensus								

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	RLRDGVPWQQVLDIILQVKG	LDKRV	DGNVIL	IAPKE	EELDLRE	236
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	RLRDGVPWQQVLDIILQVKG	LDKRV	DGNVIL	IAPKE	EELDLRE	236
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	RLRDGVPWQQVLDIILQVKG	LDKRV	DGNVIL	IAPKE	EELDLRE	237
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAEIISYLAEEETGKNITISDEIK	DTKNFRVEKSHFDEILNS	LKTHQ	LN	LKKENG	TIYYIHQAQEH....	93
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILE	NKISLKLNNIDMPRL	LQIIAKSKHL	TLN	KDDG	IYYLNGSQSG....	105
NTDB id 219973	BV139	RS02995	WP 005686897.1	ERFFIRLSQAPLAQTLEQLAFQHDVNLVMGER	LEGNISLKLNNIDMPRL	LKIIAKSKHL	TLN	KDDG	VYYLNGSQSG....	105
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQQDVNLVMGER	LEGNISLKLNNIDMPRL	LKIIAKSKHL	TLN	KDDG	VYYLNGSQSG....	105
consensus				*****	*****	*****	*****	*****	*****	*****

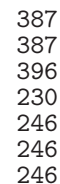
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NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINFAKAS	DIAAM	IGGEG	NVNML	SERGSIS	IDERTNS	LLIRELP	DNIAVIRE	IIIESLDI	316
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKARLAEELGDLKSEI	IKINFAKAS	DIAAM	IGGEG	NVNML	SERGSIS	IDERTNS	LLIRELP	DNIAVIRE	IIIESLDI	317
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NTDB id 219973	BV139	RS02995	WP 005686897.1	.KQQVAGNLT	TNEPHLV	SHTVKL	HFAKASELM	KSLT	.TGSG	LLSPAGS	ITFDDRS	NLLVIQ	DEPRSV	183
NTDB id 1373	NTHI	RS02645	WP 011272046.1	.KQQVAGNLT	TNEPHLV	SHTVKL	HFAKASELM	KSLT	.TGSG	LLSPAGS	ITFDDRS	NLLVIQ	DEPRSV	183
consensus										*****				

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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 1347 HI 0435 AAC22094.1
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NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

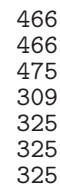
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consensus

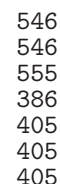
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NTDB id 1381 A4U84 RS02010 WP 021113424.1
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consensus

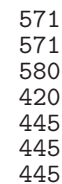
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NTDB id 1186 GTF74 RS01320 WP 000788426.1
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NTDB id 1408 DSB67 RS13980 WP 010643500.1
NTDB id 1381 A4U84 RS02010 WP 021113424.1
NTDB id 1347 HI 0435 AAC22094.1
NTDB id 219973 BV139 RS02995 WP 005686897.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

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- X non conserved
- X similar
- X >50% conserved