

NTDB id 1186	GTF74	RS01320	WP	000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP	000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP	010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...	FVAAEEVQSNLSKSIDFRTN	77
NTDB id 1381	A4U84	RS02010	WP	021113424.1			0
NTDB id 379907	FWK50	RS03475	WP	012340255.1			0
NTDB id 1347	HI	0435	AAC22094.1				0
NTDB id 1373	NTHI	RS02645	WP	011272046.1			0
consensus							0

NTDB id 1186	GTF74	RS01320	WP	000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGGEFQHD	YTLKGYKYLEVVVIS.KLKADEKPKPKSVLEKE	155
NTDB id 1175	A1552VC	RS12105	WP	000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGGEFQHD	YTLKGYKYLEVVVIS.KLKADEKPKPKSVLEKE	155
NTDB id 1408	DSB67	RS13980	WP	010643500.1	DLLNTSVKDDQLYLLDVKDFSTVVESVEVFRDSTTTRLVAYIDEDYSHD	YRLTGRYLEVKVG.KLKANEKAPDKSILDKE	156
NTDB id 1381	A4U84	RS02010	WP	021113424.1	MRYLFLLFFATF.....P..VLA		16
NTDB id 379907	FWK50	RS03475	WP	012340255.1	MLIKNFLLY..FFFLFT.....		15
NTDB id 1347	HI	0435	AAC22094.1		MKKYFLKCGYFLVCFCLPLIVFAN....P..KTD		28
NTDB id 1373	NTHI	RS02645	WP	011272046.1	MKKYFLKCGYFLVCFCLPLIVFAN....P..KTD		28
consensus						* * * * *	*

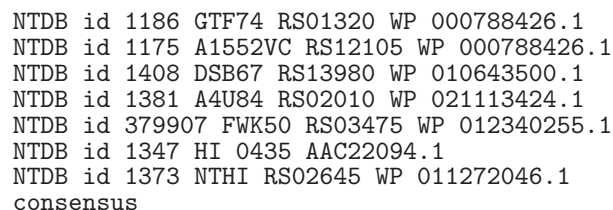
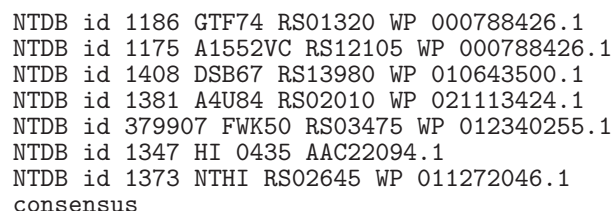
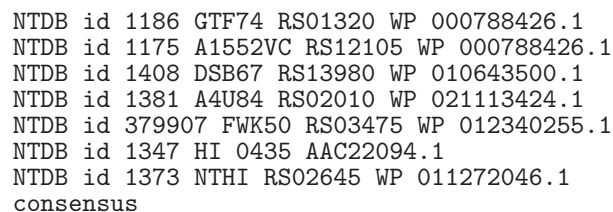
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NTDB id	1175	A1552VC	RS12105	WP	000788426.1	GKL	ISI	NFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILIAPKEELDRLR	235										
NTDB id	1408	DSB67	RS13980	WP	010643500.1	GKL	ISI	NFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILVAPKDELDRLR	236										
NTDB id	1381	A4U84	RS02010	WP	021113424.1	NQQ	ISL	SIRNAPTAEILSYLAETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENGIIYYIHQAQEHEK..	94										
NTDB id	379907	FWK50	RS03475	WP	012340255.1		VSVELSARDPFDK.....	RQRSIKKETNIQL.....D..	42										
NTDB id	1347	HI	0435	AAC22094.1	NERFF	I	RLSQAP	LIAQTLEQLAFQGDVNLVIGDILENKISLKLNNIDMPRLLLQIIAKSKHLLTNKDDGIYYLNGSQSGK..	106										
NTDB id	1373	NTHI	RS02645	WP	011272046.1	NECFF	I	RLSQAP	LIAQTLEQLAFQGDVNLVMGERLEGNISLKLNNIDMPRLLLQIIAKSKHLLTNKDDGIYYLNGSQSGK..	106									
consensus						***	*	* * *	* * *	* * *	* * *	*	* * *	* * *	* * *	* * *	* * *	* * *	*

[illegible]

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NTDB id 1186 GTF74 RS01320 WP 000788426.1
NTDB id 1175 A1552VC RS12105 WP 000788426.1
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NTDB id 1381 A4U84 RS02010 WP 021113424.1
NTDB id 379907 FWK50 RS03475 WP 012340255.1
NTDB id 1347 HI 0435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

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