

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLLAP...FVAAEEVQSNLSKSIDFRTNKDKDAVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1		0
NTDB id 379833	FWK47	RS03475	WP 012340255.1		0
NTDB id 1347	HI 0435	AAC22094.1			0
NTDB id 1373	NTHI	RS02645	WP 011272046.1		0
consensus					

NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVSVEVFRKEPSTQLVVTVDGEFQHD	YTLKGYLEVVVIS.KLKADKPKPKSVLEK	155
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVSVEVFRKEPSTQLVVTVDGEFQHD	YTLKGYLEVVVIS.KLKADKPKPKSVLEK	155
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLNNTSVKDDQLYLLDVKDFSTVVSVEVFRDTSSTRLVAYIDEDYSHD	YRLTGRYLEVKG.KLKANEKAPDKSILDK	156
NTDB id 1381	A4U84	RS02010	WP 021113424.1	MRYLFLFFATF.....P..VLA		16
NTDB id 379833	FWK47	RS03475	WP 012340255.1	MLIKNFLY..FFLFLFT.....		15
NTDB id 1347	HI 0435	AAC22094.1		MKKYFLKCGYFLVCFCLPLIVFAN....P..KT		28
NTDB id 1373	NTHI	RS02645	WP 011272046.1	MKKYFLKCGYFLVCFCLPLIVFAN....P..KT		28
consensus					*****	

NTDB id 1186 GTF74 RS01320 WP 000788426.1 GKLISINFDQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPPWQQVLDITILQVKGLDKRRVDGNVILIAPEKELDRLR 235
NTDB id 1175 A1552VC RS12105 WP 000788426.1 GKLISINFDQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPPWQQVLDITILQVKGLDKRRVDGNVILIAPEKELDRLR 235
NTDB id 1408 DSB67 RS13980 WP 010643500.1 GKLISINFDQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPPWQQVLDITILQVKGLDKRRVDGNVILVAPKDELDRLR 236
NTDB id 1381 A4U84 RS02010 WP 021113424.1 NQQISLSIRNAPTAEIISYLAETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTQHNLNKKENGIIYIHHQAQEHK.. 94
NTDB id 379833 FWK47 RS03475 WP 012340255.1VSVELSARDPFDK.....RQRSIKKETNIQL.....D.. 42
NTDB id 1347 HI 0435 AAC22094.1 NERFFTIRLSQAPLAQTLEQLAFQGDVNLVIGDILENKISLKLNNIDMPRLLIQITIAKSKHLLTNKKDDGIYYLNGSQSGK.. 106
NTDB id 1373 NTHI RS02645 WP 011272046.1 NECFFTIRLSQAPLAQTLEQLAFQGDVNLVIGERLEGNISLKLNNIDMPRLLIQITIAKSKHLLTNKKDDGIYYLNGSQSGK.. 106
consensus *** *

NTDB id	1186	GTF74	RS01320	WP	000788426.1	EKQALEKARLAEEIGDLKSEITKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREIIIESLD	315
NTDB id	1175	A1552VC	RS12105	WP	000788426.1	EKQALEKARLAEEIGDLKSEIKNFPAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREIIIESLD	315
NTDB id	1408	DSB67	RS13980	WP	010643500.1	EKQQLEKKQKLEKEIGELSSSEIKVNFAKASDIAEMINGEGNISLSDRGSMTIDERTNSLLIRELPENIEVIREIIIESLD	316
NTDB id	1381	A4U84	RS02010	WP	021113424.1	...QHHTAQLVNALPKLIKTIKLHYSKASEVIESLT.KGGQNLLSESGLYHFDDRSNSIIVKDSAASKVNFTQLIESLD	170
NTDB id	379833	FWK47	RS03475	WP	012340255.1	...TKENLSKSTNEPNLTTTSIKLHYAKAADVMKSLT.TGQGSWLSNGIFSFDERSNLLIQDNQETLNALKKLKLIKEMD	118
NTDB id	1347	HI_0435	AAC22094.1			...CQVAGNLTTNEPHLVSHTVKLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRNSLLVIQDEPRSVQNIKKLIAEMD	182
NTDB id	1373	NTHI	RS02645	WP	011272046.1	...CQVAGNLTTNEPHLVSHTVKLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRNSLLVIQDEPRFVQNIKKLIAEMD	182
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 379833 FWK47 RS03475 WP 012340255.1
NTDB id 1347 HI 0435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

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