

NTDB id 1186	GTF74 RS01320 WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFS	AESEATANQLENIDFRVNKEKA	AAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC RS12105 WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFS	AESEATANQLENIDFRVNKEKA	AAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67 RS13980 WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...	FVAAEEVQSNLSKSIDFRTNKDKDA	IVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 1381	A4U84 RS02010 WP 021113424.1				0
NTDB id 309780	CH639 RS02995 WP 011272046.1				0
NTDB id 1373	NTHI RS02645 WP 011272046.1				0
NTDB id 1347	HI 0435 AAC22094.1				0
consensus					

NTDB id	Accession	Protein	Sequence	Position
NTDB id 1186	GTF74	RS01320 WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGGEFQHDYTLKGYKYLEVVISK.LKADEKPKPKSVLEKE	155
NTDB id 1175	A1552VC	RS12105 WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGGEFQHDYTLKGYKYLEVVISK.LKADEKPKPKSVLEKE	155
NTDB id 1408	DSB67	RS13980 WP 010643500.1	DLLNTSVKDDQLYLLDVKDFSTVSVESVEVFRDSTTTRLVAYIDEDYSHDYRLTGRYLEVKGK.LKAN EKAPDKSILDKE	156
NTDB id 1381	A4U84	RS02010 WP 021113424.1	MRYLFLFFATF.....PVLA	16
NTDB id 309780	CH639	RS02995 WP 011272046.1	MKKYFLKCGYFLVCFCLPLIVFAN.....PKTD	28
NTDB id 1373	NTHI	RS02645 WP 011272046.1	MKKYFLKCGYFLVCFCLPLIVFAN.....PKTD	28
NTDB id 1347	HI 0435	AAC22094.1	MKKYFLKCGYFLVCFCLPLIVFAN.....PKTD	28
consensus			* * * * * * * *	

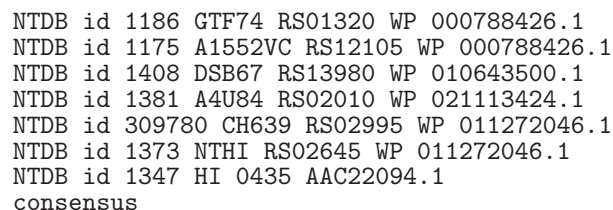
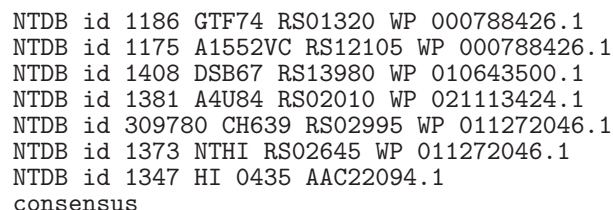
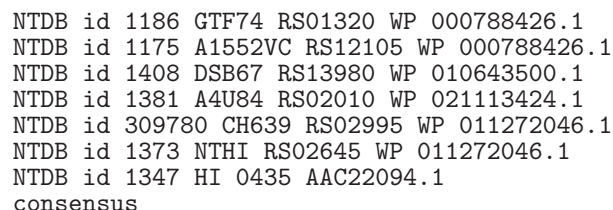
[illegible]

NTDB id 1186	GTF74	RS01320	WP 000788426.1	EKQALEKARLAEEIGDLKSEIKINFAKASDI	AAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREIIESLD	315
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	EKQALEKARLAEEIGDLKSEIKINFAKASDI	AAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREIIESLD	315
NTDB id 1408	DSB67	RS13980	WP 010643500.1	EKQKLEKQKLEKIGELSSSEIKVNF	AKASDI AEMINGEINISMLDRGSMITIDER	TNSLLIRELPDNIEVIREIIESLD
NTDB id 1381	A4U84	RS02010	WP 021113424.1	..KQHTTAQLVNALPKLITIKLHYSKASEVIESLT	.KGGNNLLSESGYLHFDDRSNIIIVKDS	SAASVKNFTQLIESLD
NTDB id 309780	CH639	RS02995	WP 011272046.1	..KQQVAGNLLTTNEPHLVSHTVKLHFAKASELMKSLT	.TGSGSLLSSAGSITFDDRSNLLVIQDEPRFVQNIKKLIAEMD	182
NTDB id 1373	NTHI	RS02645	WP 011272046.1	..KQQVAGNLLTTNEPHLVSHTVKLHFAKASELMKSLT	.TGSGSLLSSAGSITFDDRSNLLVIQDEPRFVQNIKKLIAEMD	182
NTDB id 1347	HI 0435	AAC22094.1		..KQQVAGNLLTTNEPHLVSHTVKLHFAKASELMKSLT	.TGSGSLLSPAGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMD	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 309780 CH639 RS02995 WP 011272046.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
NTDB id 1347 HI 0435 AAC22094.1
consensus

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