

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPSAVVDVQKVQEGLSI	76		
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPSAVVDVQKVQEGLSI	76		
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...	FVAAEEVQSN	SLKSIDFR	TNKKDKDAVIVVELVSPA	IVDIKRVQEGLSI	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1						0
NTDB id 310090	CH602	RS03015	WP 011272046.1						0
NTDB id 1373	NTHI	RS02645	WP 011272046.1						0
NTDB id 1347	HI 0435	AAC22094.1							0
consensus									0

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	TLKGYLEVVISK	..LKADEKPKPKSVLEKE	155	
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	TLKGYLEVVISK	..LKADEKPKPKSVLEKE	155	
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLLNTSVKDDQLYLLDVKDFSTVSVESVEVFRD	TSTTRLVAYIDEDYSHDY	RYLTGRYLEVKGK	..LKANEKAPDKSILDK	156
NTDB id 1381	A4U84	RS02010	WP 021113424.1					16
NTDB id 310090	CH602	RS03015	WP 011272046.1					28
NTDB id 1373	NTHI	RS02645	WP 011272046.1					28
NTDB id 1347	HI 0435	AAC22094.1						28
consensus								28

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	GKLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	LRLDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLR	235
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	GKLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	LRLDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLR	235
NTDB id 1408	DSB67	RS13980	WP 010643500.1	GKLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	LRLDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLR	236
NTDB id 1381	A4U84	RS02010	WP 021113424.1	NQQISLSIRNAPTAEIISYLAETGKNITISDEIKD	TKNFRVEKSHFDEILNSLIKTHQLNLKKENG	IYYIHQAQEH...		93
NTDB id 310090	CH602	RS03015	WP 011272046.1	NECFFIRLSQAPLAQTLEQLAFQQDVNLVMGERLEGNISLKLNNIDMPRL	LKIIAKSKH	LTlnKDDGVYYLNGSQSG...		105
NTDB id 1373	NTHI	RS02645	WP 011272046.1	NECFFIRLSQAPLAQTLEQLAFQQDVNLVMGERLEGNISLKLNNIDMPRL	LKIIAKSKH	LTlnKDDGVYYLNGSQSG...		105
NTDB id 1347	HI 0435	AAC22094.1		NERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRL	LQIIAKSKH	LTlnKDDGVYYLNGSQSG...		105
consensus								

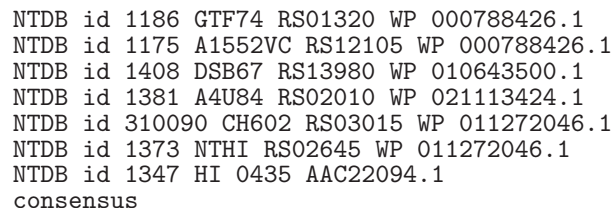
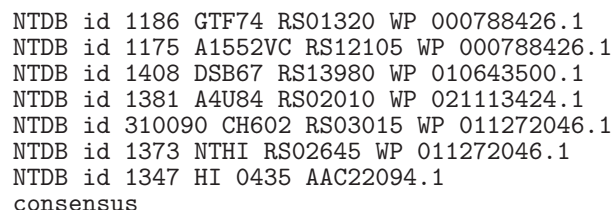
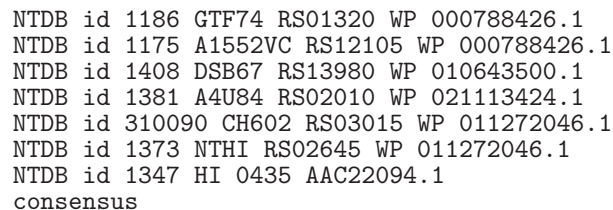
logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	EKQALEKARLAEELGDLKSEI	IKINF	AKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	IAVIREIIIESLD	315
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	EKQALEKARLAEELGDLKSEI	IKINF	AKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	IAVIREIIIESLD	315
NTDB id 1408	DSB67	RS13980	WP 010643500.1	EKQQLEKQKLEKEIGELSSEI	IKVNF	AKASDIAEMINGEGNISMLSDRGSMT	IDERTNSLLIRELPENIEVIREIIIESLD		316
NTDB id 1381	A4U84	RS02010	WP 021113424.1	..KQHTTAQLVNALPKLITKT	IKLHYSKASEVIESLT	..KGQGNLLSES	GYLHFDDRSNSITIVKDS	AASVKNFTQLIESLD	170
NTDB id 310090	CH602	RS03015	WP 011272046.1	..KQQVAGNLT	TNNEPHLVSHTVKLHFAKASELMKSLT	..TGSGSLLSS	SAGSITFDDRSNLLVIQDEPRFVQNIKKLIAEMD		182
NTDB id 1373	NTHI	RS02645	WP 011272046.1	..KQQVAGNLT	TNNEPHLVSHTVKLHFAKASELMKSLT	..TGSGSLLSS	SAGSITFDDRSNLLVIQDEPRFVQNIKKLIAEMD		182
NTDB id 1347	HI 0435	AAC22094.1		..KQQVAGNLT	TNNEPHLVSHTVKLHFAKASELMKSLT	..TGSGSLLSP	AGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMD		182
consensus				***	***	***	*****!	*****!	

```

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 310090 CH602 RS03015 WP 011272046.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
NTDB id 1347 HI 0435 AAC22094.1
consensus

```



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