

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...	FVAAEEVQSNLSKSIDFRTNKKDKDAVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 1381	A4U84	RS02010	WP	021113424.1			0
NTDB id 324223	PmCQ2	RS10890	WP	015702713.1			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	YTLKGYKYLEVVISK	LKADEKPKPK	SVLEK	EG	156																			
NTDB id 1175	A1552VC	RS12105	WP	000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGGEFQHD	YTLKGYKYLEVVISK	LKADEKPKPK	SVLEK	EG	156																			
NTDB id 1408	DSB67	RS13980	WP	010643500.1	DLLNTSVKDDQLYLLDVKDFSTVVESVEVFRDTSSTRLLVAYIDEDYSHD	YRLTGRYLEVVKVGK	LKANEKAPDK	SILDK	EG	157																			
NTDB id 1381	A4U84	RS02010	WP	021113424.1	MRYFL.....F.....FATFPVLN					17																			
NTDB id 324223	PmCQ2	RS10890	WP	015702713.1	MWRAFRKIS.FVYFLCGVAY....VGSSQAQDA					28																			
NTDB id 1347	HI	0435	AAC22094.1	MKKYFLKCGYFLVCFCLPLI....VFANPKTDN						29																			
NTDB id 1373	NTHI	RS02645	WP	011272046.1	MKKYFLKCGYFLVCFCLPLI....VFANPKTDN					29																			
consensus						Y	K	L	V	I	S	K	L	K	A	D	E	K	P	K	P	K	S	V	L	E	K	E	G

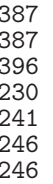
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consensus		*** * * * ! * * * * *	* ! * * *	*****	***** * * * ! * * * * *	*****

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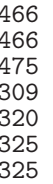
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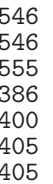
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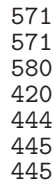
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consensus



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consensus



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NTDB id 324223 PmCQ2 RS10890 WP 015702713.1
NTDB id 1347 HI 0435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus



- X non conserved
- X similar
- X $\geq 50\%$ conserved