

NTDB id 1186	GTF74 RS01320 WP 000788426.1	..MKNGLKT..YVAQTLTLWVGLALCASSMVFSASATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC RS12105 WP 000788426.1	..MKNGLKT..YVAQTLTLWVGLALCASSMVFSASATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67 RS13980 WP 010643500.1	MKMRHGLTLTPRKLTLGLFSLWFSLLLAP...FVAAEEVQSNLSKSIDFRTNKDKDAVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 1381	A4U84 RS02010 WP 021113424.1		0
NTDB id 379932	FWK55 RS03480 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	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHD	YTLKGYLYEVVIS.KLKADKPKPKSVLEK	155
NTDB id 1175	A1552VC	RS12105	WP	000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHD	YTLKGYLYEVVIS.KLKADKPKPKSVLEK	155
NTDB id 1408	DSB67	RS13980	WP	010643500.1	DLLNTSVKDDQLYLLDVKDFSTVSVESVEVFRDTSSTRLVAYIDEDYSHD	YRLTGRYLEVVKVG.KLKANEKAPDKSILDK	156
NTDB id 1381	A4U84	RS02010	WP	021113424.1	MRYLFLFFATF.....P..VLA		16
NTDB id 379932	FWK55	RS03480	WP	012340255.1	MLIKNFLY..FFLFLFT.....		15
NTDB id 1347	HI 0435	AAC22094.1			MKKYFLKCGYFLVCFCLPLIVFAN....P..KTD		28
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consensus					*		

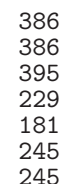
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NTDB id	1408	DSB67	RS13980	WP	010643500.1	GKLTISIN	FQDIPVRNVLQLIADYNGFNLVVSDSVEGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILIA	PKKDELD	LDR	236										
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consensus						***	* * *	*****	* * *	*	*****	* * *	*****	* * *	*	*****	* * *	*****	* * *	*

[illegible]

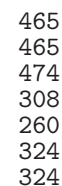
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NTDB id 1186 GTF74 RS01320 WP 000788426.1
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NTDB id 379932 FWK55 RS03480 WP 012340255.1
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consensus

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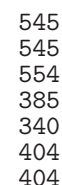
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NTDB id 379932 FWK55 RS03480 WP 012340255.1
NTDB id 1347 HI 0435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus



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NTDB id 1186 GTF74 RS01320 WP 000788426.1
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NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

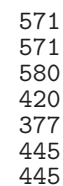
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NTDB id 1186 GTF74 RS01320 WP 000788426.1
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NTDB id 379932 FWK55 RS03480 WP 012340255.1
NTDB id 1347 HI 0435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

```



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