

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 180764	A4216	RS08875	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	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	YTLKG	KYLEVV	ISK	LKADEKPKPK	SVLEK	EG	156
NTDB id 1175	A1552VC	RS12105	WP	000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	YTLKG	KYLEVV	ISK	LKADEKPKPK	SVLEK	EG	156
NTDB id 1408	DSB67	RS13980	WP	010643500.1	DLLNTSVKDDQLYLLDVKDFSTPVESVEVFRDSTTRLVAYIDEDYSHD	YRLTG	RYLVEVK	VGK	LKANEKAPDK	SILDK	EG	157
NTDB id 1381	A4U84	RS02010	WP	021113424.1	MRYLFL.....F.....FATFPVLN							17
NTDB id 180764	A4216	RS08875	WP	015702713.1	MWRAFRKIS.FVYFLCGVAY.....VGSSQAQDA							28
NTDB id 1347	HI	0435	AAC22094.1	MKKYFLKCGYFLVCFCLPLI.....VFANPKTDN								29
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consensus												

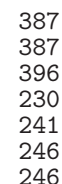
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NTDB id 1408	DSB67	RS13980	WP	010643500.1	KLISINFDQDIPVRNVLQLIADYNGFNLVVS	SVVGNLTLRLDGV	PWQQVLDIILQVKGLDKRV	DGNVIL	VAPKD	ELDLRE	237											
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NTDB id 1347	HI	0435	AA	C22094.1	ERFFIRLSGAPLAQTLEQLAFQGDVNLVI	GDILENKISLKLNNIDMPRL	LKIIAKSKHLLTNKDDG	IYYLNGSQSG	...		105											
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consensus					***	*	*	*	!	*	*****	*	*****	*	*	!	*	*****	*	*	*****	*

[illegible]

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NTDB id 1186 GTF74 RS01320 WP 000788426.1
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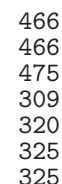
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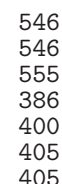
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consensus

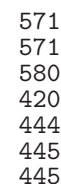
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NTDB id 180764 A4216 RS08875 WP 015702713.1
NTDB id 1347 HI 0435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

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-  non conserved
-  similar
-  >50% conserved