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NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...FVAAEEVQSNLSKSIDFRNTKDKDAVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1		0
NTDB id 222244	A6J55	RS04430	WP 005754751.1		0
NTDB id 1347	HI 0435	AAC22094.1			0
NTDB id 1373	NTHI	RS02645	WP 011272046.1		0
consensus					

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLTKGKYLEVVISKIKADEKPKPKSVLEKEG	156
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLTKGKYLEVVISKIKADEKPKPKSVLEKEG	156
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLLNTSVKDDQLYLLDVKDFSTVSVESVEVFRDTSTTRLVAYIDEDYSHDYRLTGRYLEVKGKIKANEKAPDKSILDKEG	157
NTDB id 1381	A4U84	RS02010	WP 021113424.1	MRYLFL.....F.....FATFPVLAN	17
NTDB id 222244	A6J55	RS04430	WP 005754751.1	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					

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPPWQQVLDIILQVKGLDKRVDGNVILVAPKEELDRE	236
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPPWQQVLDIILQVKGLDKRVDGNVILVAPKEELDRE	236
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NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAEIISYLAEEETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENGITYYIHQAQEH....	93
NTDB id 222244	A6J55	RS04430	WP 005754751.1	EHFYLRLLKQAPLVEMLQYLALQHQDLLIDDHLEGTLSLQMKKTTFEKCLQSIARMKQLELHQEGKSYLTSPSG....	103
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NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQQDVNLVIGERLEGNISLKLNNIDMPRLKIIAKSKHLTLNKKDDGIVYYLNQSQSG....	105
consensus					

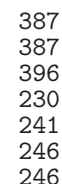
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NTDB id 1381	A4U84	RS02010	WP 021113424.1	.KQHHTTAQLVNALPKLITKTIKLHYSKASEVIESLT.KGQGNLLSESGYLHFDDRSNSIIVKDSAAASKNFTQLIESLDK	171
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consensus					

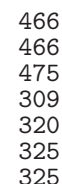
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consensus

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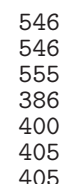
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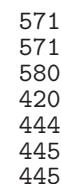
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consensus

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-  non conserved
-  similar
-  >50% conserved