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NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQVEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQVEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...FVAAEEVQSNLSKSIDFRNTKDKDAVIVVELVSPAAIVDIKRVQVEGLSI	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1		0
NTDB id 367734	FEE42	RS07405	WP 059368781.1		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	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGGKYLEVVISKLKADEKPKPKSVLEKEG	156
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGGKYLEVVISKLKADEKPKPKSVLEKEG	156
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLLNTSVKDDQLYLLDVKDFSTVSVESVEVFRDSTTTRLVAYIDEDYSHDYRLTGRYLEVKGKLANEKAPDKSILDKEG	157
NTDB id 1381	A4U84	RS02010	WP 021113424.1	MRYLFLFFFA.....TFPVLAN	17
NTDB id 367734	FEE42	RS07405	WP 059368781.1	MKVWIKKCGLFFLCFCCGN.....EVAAE.FKEN	28
NTDB id 1347	HI 0435	AAC22094.1		MKKYFLKCGYFLVCFCLPL.....IVFANPKTDN	29
NTDB id 1373	NTHI	RS02645	WP 011272046.1	MKKYFLKCGYFLVCFCLPL.....IVFANPKTDN	29
consensus					

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILIAPKEELDLRE	236
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILIAPKEELDLRE	236
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NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAEIISYLAEEETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENGIIYYIHQAQE.....	92
NTDB id 367734	FEE42	RS07405	WP 059368781.1	QVFSIRLKQAPMVSAQLQELALVQGANLIIDDALDDSLSLQLDNVDFDRVLRSAKMKGLSLRLEDDIYYLGGKDDVVFSET	108
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRISQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRLQIIAKSKHLTLNKKDDGIYYLNGSQ.....	103
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRISQAPLAQTLEQLAFQQDVNLVIGERLEGNISLKLNNIDMPRLKIIAKSKHLTLNKKDDGVYYLNGSQ.....	103
consensus					

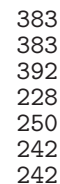
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NTDB id 367734	FEE42	RS07405	WP 059368781.1	SLQPTENEVLANIKQIEPSLVSKTIKLNFAKASEVMKSLT.TGSGSLLSASGSIAFDDRSNLLIIQDDAKSVKNLVKLI A	187
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consensus

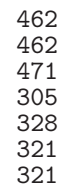
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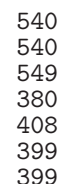
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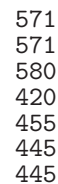
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consensus

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