

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLVIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLVIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...FVAAEEVQSNLSKSIDFRNTKDKDAVIVVELVSPAIVDIKRVQEGLSI	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1		0
NTDB id 1347	HI 0435	AAC22094.1			0
NTDB id 71036	K6J68	RS02915	WP 005686897.1		0
NTDB id 1373	NTHI	RS02645	WP 011272046.1		0
consensus					

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGYLEVVISKLKADEKPKPKSVLEKEG	156
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGYLEVVISKLKADEKPKPKSVLEKEG	156
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLLNTSVKDDQLYLLDVKDFSTVSVESVEVFRDSTTTRLVAYIDEDYSHDYRLTGRYLEVKGKLANEKAPDKSILDKEG	157
NTDB id 1381	A4U84	RS02010	WP 021113424.1	MRYLFLFFAT.....FPVLAN	17
NTDB id 1347	HI 0435	AAC22094.1		MKKYFLKCGYFLVCFCLPLIVF.A....NPKTDN	29
NTDB id 71036	K6J68	RS02915	WP 005686897.1	MKKYFLKCGYFLVCFCLPLIVF.A....NPKTDN	29
NTDB id 1373	NTHI	RS02645	WP 011272046.1	MKKYFLKCGYFLVCFCLPLIVF.A....NPKTDN	29
consensus					

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILVAPKEELDRLRE	236
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILVAPKEELDRLRE	236
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILVAPKDELDLRE	237
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAEIISYLAEEETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENGITYYIHQAQEH....	93
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRLQTIAKSKHLTLNKKDDGVIYLLNGSQSG....	105
NTDB id 71036	K6J68	RS02915	WP 005686897.1	ERFFIRLSQAPLAQTLEQLAFQHDVNLVMGERLEGNISLKLNNIDMPRLKIIAKSKHLTLNKKDDGVIYLLNGSQSG....	105
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQQDVNLVMGERLEGNISLKLNNIDMPRLKIIAKSKHLTLNKKDDGVIYLLNGSQSG....	105
consensus					

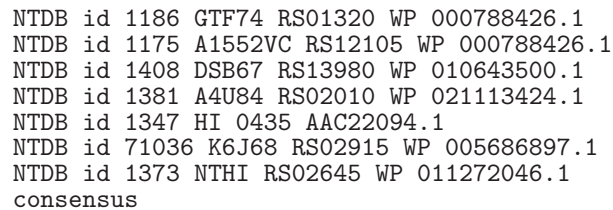
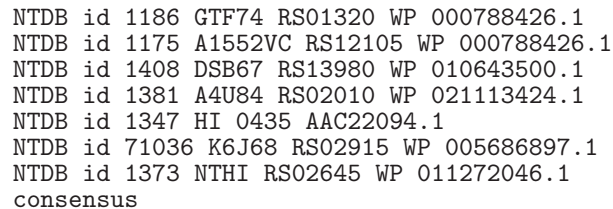
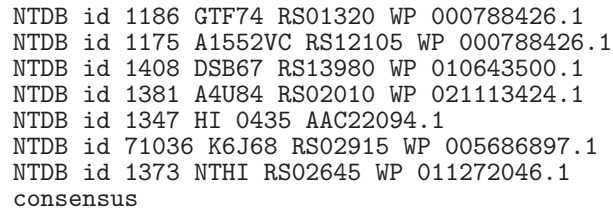
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KQALEKARLAEELGDLKSEI IKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNI AVIREIIESLDI	316
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEELGDLKSEI IKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNI AVIREIIESLDI	316
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKARLAEELGDLKSEI IKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNI AVIREIIESLDI	317
NTDB id 1381	A4U84	RS02010	WP 021113424.1	.KQHHTTAQLVNALPKLITKTIKLHYSKASEVIESLT.KGQGNLLSESGLYLFHDDRNSNIIIVKDSAASVKNFTQLIESLDK	171
NTDB id 1347	HI 0435	AAC22094.1		.KQQVAGNLTITNEPHLVSHTVKLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK	183
NTDB id 71036	K6J68	RS02915	WP 005686897.1	.KQQVAGNLTITNEPHLVSHTVKLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK	183
NTDB id 1373	NTHI	RS02645	WP 011272046.1	.KQQVAGNLTITNEPHLVSHTVKLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK	183
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consensus

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- ☐ non conserved
- ☒ similar
- ☒ $\geq 50\%$ conserved