

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFS	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFS	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLLAP...FVAAEEVQSNLSKSIDFR	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1		0
NTDB id 1071092	ACIRXL	RS02790	WP 410679840.1		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	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY..TLKGYLEVVIS.KLKADEKPKPKSVLE	153
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY..TLKGYLEVVIS.KLKADEKPKPKSVLE	153
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLNNTSVKDDQLYLLDVKDFSTVVESVEVFRDSTTRLVAYIDEDYSHDY..RLTGRYLEVKKV.G.KLKANEKAPDKSILD	154
NTDB id 1381	A4U84	RS02010	WP 021113424.1	MRYLFFLFFATF.....PV	14
NTDB id 1071092	ACIRXL	RS02790	WP 410679840.1	MKLLNIIKNILLFFLILSYPF.R.....LVNAS	27
NTDB id 1347	HI 0435	AAC22094.1		MKKY..FLKCGYFLVCFCLPLIV.....FANPK	26
NTDB id 1373	NTHI	RS02645	WP 011272046.1	MKKY..FLKCGYFLVCFCLPLIV.....FANPK	26
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KEGKLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILVAPKEELD	233
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KEGKLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILVAPKEELD	233
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KEGKLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILVAPKDELD	234
NTDB id 1381	A4U84	RS02010	WP 021113424.1	LANQQISLSIRNAPTAEIISYLAEEETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENGITYYIHQAQE.H	93
NTDB id 1071092	ACIRXL	RS02790	WP 410679840.1	DENNTFSVHLKQAPLVDTVQYLAGYQKINLVIDDELSCSISLQLQNTTLLQLLKSIAELKHLRFWENNGIYYLSQSQA.T	106
NTDB id 1347	HI 0435	AAC22094.1		TDNERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRLQITIAKSKHLTLNKDDGIYYLNGSQS.G	105
NTDB id 1373	NTHI	RS02645	WP 011272046.1	TDNCEFFIRLSQAPLAQTLEQLAFQQDVNLVIGERLEGNISLKLNNIDMPRLKIIAKSKHLTLNKDDGVYYLNGSQS.G	105
consensus					

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	LRKQALEKARLAEEELGDLKSEI IKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREITIES	313
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	LRKQALEKARLAEEELGDLKSEI IKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREITIES	313
NTDB id 1408	DSB67	RS13980	WP 010643500.1	LRKQALEKQKLEKEIGELSSSEI IKVNFAKASDIAEMINGEGNISMLSDRGSMTIDERTNSLLIRELPENIEVIREITIES	314
NTDB id 1381	A4U84	RS02010	WP 021113424.1	KQHTTA...QLVNALPKLITKTIKLHYSKASEVIESLT.KGQGNLLSESGYLHFDDRNSNIIVKDSAAASVKNFTQLIES	168
NTDB id 1071092	ACIRXL	RS02790	WP 410679840.1	NQENVVLNSSTDIATSSADMLITSIKLHYAKASEVMKALT.SGNSLLTVNGSISFDERNNLLIIQDEAKKIKIRKLIVAD	185
NTDB id 1347	HI 0435	AAC22094.1		KGQVAG...NLTITNEPHLVSHITVKLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRNSNLLVIQDEPRSVQNIKKLIAE	180
NTDB id 1373	NTHI	RS02645	WP 011272046.1	KGQVAG...NLTITNEPHLVSHITVKLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRNSNLLVIQDEPRFVQNIKKLIAE	180
consensus					

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	LDIPVKQVQIEARIVTVKEGNLEELGVRWGVMS	384
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	LDIPVKQVQIEARIVTVKEGNLEELGVRWGVMS	384
NTDB id 1408	DSB67	RS13980	WP 010643500.1	LDIPVKQVQIEARIVTINEGNMDELGVRWGFTS	393
NTDB id 1381	A4U84	RS02010	WP 021113424.1	LDKPTEQIAIEARIVTISSEHLQLQGVWGLFSP	228
NTDB id 1071092	ACIRXL	RS02790	WP 410679840.1	MDKPIQIIIEARIATISDES	248
NTDB id 1347	HI 0435	AAC22094.1		MDKPIEQIAIEARIVTITDES	243
NTDB id 1373	NTHI	RS02645	WP 011272046.1	MDKPIEQIAIEARIVTITDES	243
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	TSANASSIAFQVAKLGS	463
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	TSANASSIAFQVAKLGS	463
NTDB id 1408	DSB67	RS13980	WP 010643500.1	TSPNATSI	472
NTDB id 1381	A4U84	RS02010	WP 021113424.1	.VNPSAS	306
NTDB id 1071092	ACIRXL	RS02790	WP 410679840.1	SSNPAGS	327
NTDB id 1347	HI 0435	AAC22094.1		TITPAGS	322
NTDB id 1373	NTHI	RS02645	WP 011272046.1	TITPAGS	322
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	LKVTPQITPDNR	543
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	LKVTPQITPDNR	543
NTDB id 1408	DSB67	RS13980	WP 010643500.1	LKVTPQITPDNR	552
NTDB id 1381	A4U84	RS02010	WP 021113424.1	LEVTPHISKQ	383
NTDB id 1071092	ACIRXL	RS02790	WP 410679840.1	LEVTPHILKHNG	407
NTDB id 1347	HI 0435	AAC22094.1		LEVTPHISKDNN	402
NTDB id 1373	NTHI	RS02645	WP 011272046.1	LEVTPHISKDNN	402
consensus					

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	LGALFRRTIEQMGKSELLIFVTPKVV	571
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	LGALFRRTIEQMGKSELLIFVTPKVV	571
NTDB id 1408	DSB67	RS13980	WP 010643500.1	LGALFRRSYENVGKSELLIFVTPKVV	580
NTDB id 1381	A4U84	RS02010	WP 021113424.1	IKRLFSQSQDKISKREL	420
NTDB id 1071092	ACIRXL	RS02790	WP 410679840.1	IKRLFSKEVERHQKREL	440
NTDB id 1347	HI 0435	AAC22094.1		IKRLFSKESERHQKREL	445
NTDB id 1373	NTHI	RS02645	WP 011272046.1	IKRLFSKESERHQKREL	445
consensus					

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved