

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPS	AVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPS	AVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...	FVAAEEVQSN	SLKSIDFR	TNKKDKDAVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1					0
NTDB id 1347	HI 0435	AAC22094.1						0
NTDB id 304500	DV384	RS06010	WP 114934598.1					0
NTDB id 1373	NTHI	RS02645	WP 011272046.1					0
consensus								0

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	TLKGKYLEVVIS	KLKADEKPKPKSVLEKEG	156	
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	TLKGKYLEVVIS	KLKADEKPKPKSVLEKEG	156	
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLNNTSVKDDQLYLLDVKDFSTVSVESVEVFRDTSTTRLVAYIDEDYSHD	YRLTGRLGYLEV	KVGKLANEKAPDKSILDKEG	157	
NTDB id 1381	A4U84	RS02010	WP 021113424.1					17
NTDB id 1347	HI 0435	AAC22094.1						29
NTDB id 304500	DV384	RS06010	WP 114934598.1					29
NTDB id 1373	NTHI	RS02645	WP 011272046.1					29
consensus								29

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSV	VEGNLT	RLNNVPWQRV	DKGGVYYLNPSQE	236
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSV	VEGNLT	RLNNVPWQRV	DKGGVYYLNPSQE	236
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSV	VEGNLT	RLNNVPWQRV	DKGGVYYLNPSQE	237
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAEIISYLAEEETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENGITYYIHQAQE				92
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRLQTI	AKSKH	LTNLKDDG	IYYLNGSQS	104
NTDB id 304500	DV384	RS06010	WP 114934598.1	ERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRLQTI	AKSKH	LTNLKDDG	IYYLNGSQS	104
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRLQTI	AKSKH	LTNLKDDG	IYYLNGSQS	104
consensus								104

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINFAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREIIESLDI	316
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINFAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREIIESLDI	316
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKARLAEELGDLKSEI	IKINFAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREIIESLDI	317
NTDB id 1381	A4U84	RS02010	WP 021113424.1	HKQHTTAQLVNALPKLITKTIKLHYSKASEVIESLT	.KGQGNLLSES	GYLHFD	DRSNSIIVKDSAASVKNFTQLIESLDK	171
NTDB id 1347	HI 0435	AAC22094.1		GKQGVAGNLTITNEPHLVSHTVKLHFAKASELMKSLT	.TGSGSLLSPAGS	ITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK		183
NTDB id 304500	DV384	RS06010	WP 114934598.1	GKQGVAGNLTITNEPHLVSHTVKLHFAKASELMKSLT	.TGSGSLLSPAGS	ITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK		183
NTDB id 1373	NTHI	RS02645	WP 011272046.1	GKQGVAGNLTITNEPHLVSHTVKLHFAKASELMKSLT	.TGSGSLLSPAGS	ITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK		183
consensus								183

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consensus

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