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NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...	FVAAEEVQSN	SLKSIDFRNTNKDKDAVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1				0
NTDB id 180775	A4203	RS02085	WP 015702713.1				0
NTDB id 1347	HI 0435	AAC22094.1					0
NTDB id 1373	NTHI	RS02645	WP 011272046.1				0
consensus							

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	TLTKGKYLEVV	ISK	KADEKPKPKSVLEKEG	156
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	TLTKGKYLEVV	ISK	KADEKPKPKSVLEKEG	156
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLLNTSVKDDQLYLLDVKDFSTVSVESVEVFRDTSTTRLVAYIDEDYSHD	YRLTG	RYLEV	KGKIKANEKAPDKSILDKEG	157
NTDB id 1381	A4U84	RS02010	WP 021113424.1					17
NTDB id 180775	A4203	RS02085	WP 015702713.1					28
NTDB id 1347	HI 0435	AAC22094.1						29
NTDB id 1373	NTHI	RS02645	WP 011272046.1					29
consensus								
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logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGV	PWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	236
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGV	PWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	236
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGV	PWQQVLDIILQVKGLDKRVDGNVIL	IAPKDEL	DLRE	237
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAEIISYLAEEETGKNITISDEIKDTKNFRVEKSHFDE	ILNSLIKTHQLNLKKENG	IYYIHQAQEH		93
NTDB id 180775	A4203	RS02085	WP 015702713.1	EHFYLRLLKQAPLVEMLQYLALQHQD	LLIDDHLEGTLSLQMKKTTFEKCLQSIARMKQLELHQEGKSYLT	TSPSG		103
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRL	LQIIAKSKHLT	LNKDDG	IYYLNGSQSG....	105
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRL	LKIIAKSKHLT	LNKDDG	VYYLNGSQSG....	105
consensus								
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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	.KQHHTAQLVNALPKLITKTIKLHYSKASEVIESLT	.KGQGNLLSESGYLHFD	DRSNSIIVKDS	AASVKNFTQLIESLDK	171	
NTDB id 180775	A4203	RS02085	WP 015702713.1	...VAANDTHHTPSLMTSSIKLHF	AKAEVVKSLT	.SGQGSLLSVGGSLSF	DERTNLLLIQDEPQSIQRIKALVAEMDK	178	
NTDB id 1347	HI 0435	AAC22094.1		.KGQVAGNLT	TNEPHLVSH	TVKLHFAKASELMKSLT	.TSGSGLLS	PAGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK	183
NTDB id 1373	NTHI	RS02645	WP 011272046.1	.KQQVAGNLT	TNEPHLVSH	TVKLHFAKASELMKSLT	.TSGSGLLS	PAGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK	183
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