

logo

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 92488	MFE95	RS12770	WP 015702713.1				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	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	DLNNTSVKDDQLYLLDVKDFSTVSVESVEVFRDSTTTRLVAYIDEDYSHD	YRLTG	RYLEV	KGKIKANEKAPDKSILDKEG	157
NTDB id 1381	A4U84	RS02010	WP 021113424.1					17
NTDB id 92488	MFE95	RS12770	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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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 92488	MFE95	RS12770	WP 015702713.1	EHFYLRLLKQAPLVEMLQYLALQHQHQLLIDDHLEGTLSLQMKKTTFEKCL	QSIARMKQLELHQEGKSYLT	TSPSG....		103
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRL	LQIIAKSKHLTLNKKDDG	IYYLNGSQSG....		105
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRL	LQIIAKSKHLTLNKKDDG	IYYLNGSQSG....		105
consensus								
				*** * **!* ***** *! * ***** ***** * * *! ***** !*** ** ***** *				

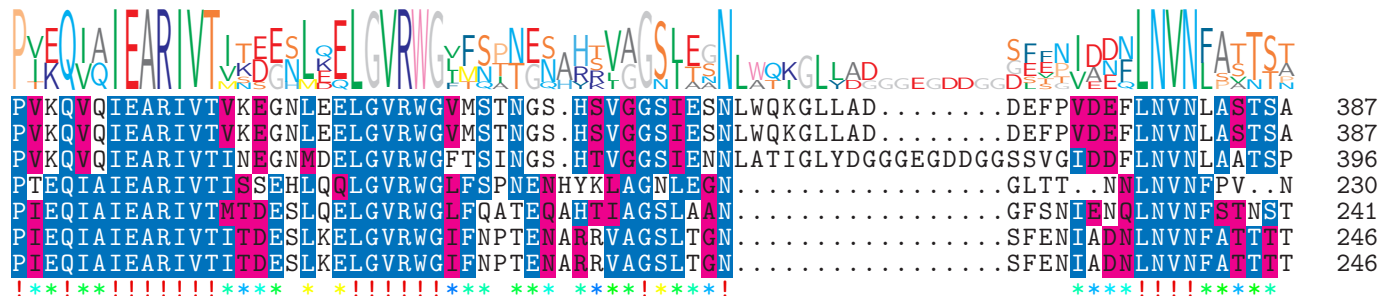
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINFAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREIIESLDI	316
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINFAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREIIESLDI	316
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKARLAEELGDLKSEI	IKINFAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREIIESLDI	317
NTDB id 1381	A4U84	RS02010	WP 021113424.1	.KQHHTTAQLVNALPKLITKT	IKLHYSKASEVIESLT.KGQGNL	LSSEGYLHFD	DRSNSIIVKDSAASVKNFTQLIESLDK	171
NTDB id 92488	MFE95	RS12770	WP 015702713.1	...VAANDTHHTPSLMTSS	IKLHFAKAAEVVKS	LT.SGQGSLLSVGGS	LSFDERTNLLVIQDEPQSIQRIKALVAEMDK	178
NTDB id 1347	HI 0435	AAC22094.1		.KGQVAGNLT	TNEPHLVSH	TVKLHFAKASELMKSLT.TGSGSLLSPAGS	ITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK	183
NTDB id 1373	NTHI	RS02645	WP 011272046.1	.KGQVAGNLT	TNEPHLVSH	TVKLHFAKASELMKSLT.TGSGSLLSPAGS	ITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK	183
consensus								
				* * * * * !** **!* ***** !*****! * !*****!***** * * ** *****!*				

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consensus
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-  non conserved
-  similar
-  >50% conserved