

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	SLKSIDFRNTNKDKDAVIVVELVSPA	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1				0
NTDB id 421056	GV127	RS10260	WP 061406404.1				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	DLLNTSVKDDQLYLLDVKDFSTVSVESVEVFRDTSTTRLVAYIDEDYSHD	YRLTG	RYLEV	KGKIKANEKAPDKSILDKEG	157
NTDB id 1381	A4U84	RS02010	WP 021113424.1					17
NTDB id 421056	GV127	RS10260	WP 061406404.1					28
NTDB id 1347	HI 0435	AAC22094.1						29
NTDB id 1373	NTHI	RS02645	WP 011272046.1					29
consensus								
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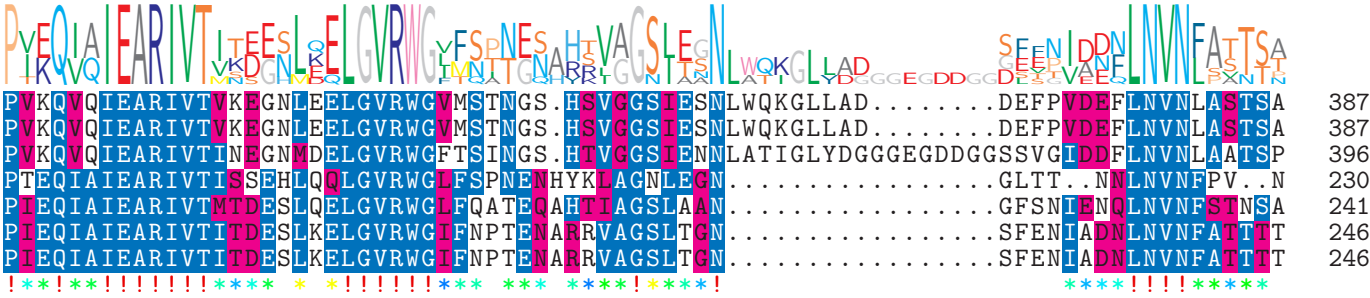
NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGV	PWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	236
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGV	PWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	236
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGV	PWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	237
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAEIISYLAEEETGKNITISDEIKDTKNFRVEKSHFDE	ILNSLIKTHQLNLKKENG	IYYIHQAQEH		93
NTDB id 421056	GV127	RS10260	WP 061406404.1	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	KQALEKARLAEEELGDLKSEI	IKINFAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREIIIESLDI	316
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEEELGDLKSEI	IKINFAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREIIIESLDI	316
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKARLAEEELGDLKSEI	IKINFAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREIIIESLDI	317
NTDB id 1381	A4U84	RS02010	WP 021113424.1	.KQHHTTAQLVNALPKLITKT	TIKLHYSKASEVIESLT.KGQGNLLSESGYLHFD	DRSNSIIVKDS	AASVKNFTQLIESLDK	171
NTDB id 421056	GV127	RS10260	WP 061406404.1	...VAANDTHHTPSLMTSS	IKLHFAKAAEVVKS	LT.SGQGSLLSVGGSLSF	DERTNLLVIQDEPQSI	178
NTDB id 1347	HI 0435	AAC22094.1		.KQQVAGNLT	TNEPHLVSH	TVKLHFAKASELMKSLT.TGSGSLLSPAGSIT	FDDRSNLLVIQDEPR	183
NTDB id 1373	NTHI	RS02645	WP 011272046.1	.KQQVAGNLT	TNEPHLVSH	TVKLHFAKASELMKSLT.TGSGSLLSPAGSIT	FDDRSNLLVIQDEPR	183
consensus								
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consensus



☒ non conserved
☒ similar
☒ ≥ 50% conserved