

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 222264	A6J57	RS02480	WP 005757388.1				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	DLLNTSVKDDQLYLLDVKDFSTVSVESVEVFRDSTTTRLVAYIDEDYSHD	YRLTG	RYLEV	KGKIKANEKAPDKSILDKEG	157
NTDB id 1381	A4U84	RS02010	WP 021113424.1					17
NTDB id 222264	A6J57	RS02480	WP 005757388.1					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	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	LRLDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	236
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	LRLDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	236
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	LRLDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	237
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAEIISYLAEEETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENG	IYYIHQAQEH			93
NTDB id 222264	A6J57	RS02480	WP 005757388.1	EHFYLRLLKQAPLVEMLQYLALQHQD	LLIDDHLEGTLSLQMKKTTFEKCLQSIARMKQLELHQEGKSYLTSPSG			103
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRL	LQIIAKSKHLT	LNKDDG	IYYLNGSQSG	105
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRL	LQIIAKSKHLT	LNKDDG	IYYLNGSQSG	105
consensus								
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINF	FAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREII	IESLDI	316
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINF	FAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREII	IESLDI	316
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKARLAEELGDLKSEI	IKINF	FAKASDIAAMIGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	AVIREII	IESLDI	317
NTDB id 1381	A4U84	RS02010	WP 021113424.1	.KQHTTAQLVNALPKLITKT	IKLHYSKASEVIESLT	.KQQGNLLSESGYLHFD	DRSNSITIVKDSAAS	VKNFTQL	IESLDK	171
NTDB id 222264	A6J57	RS02480	WP 005757388.1	...VAANDTHHTPSLMTSS	IKLHF	FAKAEVVKSLT	.SGQGSLLSVGGSLSF	DERTNLLVIQDEPQSI	QRIKALVAEMDK	178
NTDB id 1347	HI 0435	AAC22094.1		.KQQVAGNLT	TNEPHLVSH	TVKLHFAKASELMKSLT	.TSGSGLLS	PAGSITFDDRSNLLVIQDEPR	SVQNIKKLIAEMDK	183
NTDB id 1373	NTHI	RS02645	WP 011272046.1	.KQQVAGNLT	TNEPHLVSH	TVKLHFAKASELMKSLT	.TSGSGLLS	PAGSITFDDRSNLLVIQDEPR	SVQNIKKLIAEMDK	183
consensus										
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consensus



☒ non conserved
☒ similar
☒ ≥ 50% conserved