

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...FVAAEEVQSNLSKSIDFRTNKDKDAVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1		0
NTDB id 993527	AAIB12	RS11240	WP 010907082.1		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	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGKYLEVVISKKADEKPKPKSVLEKEG	156
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGKYLEVVISKKADEKPKPKSVLEKEG	156
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLNNTSVKDDQLYLLDVKDFSTVVESVEVFRDSTTRLVAYIDEDYSHDYRLTGKYLEVVKVGLKANEKAPDKSILDKEG	157
NTDB id 1381	A4U84	RS02010	WP 021113424.1	MRYLFLLF.....FAIFPVLAN	17
NTDB id 993527	AAIB12	RS11240	WP 010907082.1	MWRAFRLISLVYFICGWAY....VGSQAQDA	28
NTDB id 1347	HI 0435	AAC22094.1		MKKYFLKCGYFLVCFCLPLI....VFANPKTDN	29
NTDB id 1373	NTHI	RS02645	WP 011272046.1	MKKYFLKCGYFLVCFCLPLI....VFANPKTDN	29
consensus					

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILAPKEELDRE	236
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILAPKEELDRE	236
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILAPKDELDRE	237
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAEIISYLAETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKENGNYIYHQAQEH....	93
NTDB id 993527	AAIB12	RS11240	WP 010907082.1	EHFYLRLLKQAPLVEMLQYLALQHQDILLDDHLEGTLSLQMKKTTEKCLQSIARMKQLELHQECKSYVLTSPSG....	103
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLACTLEQLAFQCDVNLVIGDILENKISLKLNNIDMPRLQITIAKSKHLTLNKDDGIYYNGSQSG....	105
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLACTLEQLAFQCDVNLVIGDILENKISLKLNNIDMPRLQITIAKSKHLTLNKDDGVVYVNGSQSG....	105
consensus					

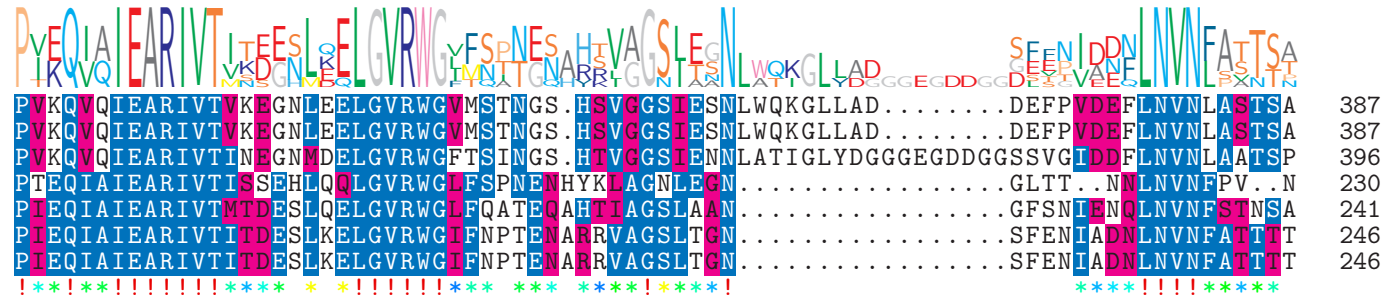
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KQALEKARLAEEIGDLKSEIKNFPAKASDIAAMIGGECNVNMLSERGSISIDERTNSLLIRELPDNIAVIRETIESLDI	316
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEEIGDLKSEIKNFPAKASDIAAMIGGECNVNMLSERGSISIDERTNSLLIRELPDNIAVIRETIESLDI	316
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKQKLEKEIGELSSSEIKNFPAKASDIAEMINGECNISMLSDRGSMTIDERTNSLLIRELPENIEVIRETIESLDI	317
NTDB id 1381	A4U84	RS02010	WP 021113424.1	.KQHTTAQLVNALPKLITTKIKLHYSKASEVIESLT.KGCGNLLSESGLYHFDERSNSITVKDSAAASVKNFTQLIESLDK	171
NTDB id 993527	AAIB12	RS11240	WP 010907082.1	...VAANDTHPTSLMTSSIKLHFAKAAEVVKSILT.SCGSLLSVGSSLSFDERTNLLIQDEPQSIQRKALVAEMDK	178
NTDB id 1347	HI 0435	AAC22094.1		.KQVAGNLTNNEPHLVSHTVKLHFAKASELMKSLT.TCGSLLSPAGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK	183
NTDB id 1373	NTHI	RS02645	WP 011272046.1	.KQVAGNLTNNEPHLVSHTVKLHFAKASELMKSLT.TCGSLLSPAGSITFDDRSNLLVIQDEPRFVQNIKKLIAEMDK	183
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- non conserved
- similar
- $\geq 50\%$ conserved