

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1		
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1381	A4U84	RS02010	WP 021113424.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...FVAAEEVQSNLSKSIDFRTNKDKDAVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 111192	ACMCNG	RS08505	WP 226691996.1		0
NTDB id 1347	HI 0435	AAC22094.1			0
NTDB id 1373	NTHI	RS02645	WP 011272046.1		0
consensus					0

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NTDB id 1186	GTF74	RS01320	WP 000788426.1		
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGKYLEVVISKLKADEKPKPKSVLEKEG	156
NTDB id 1408	DSB67	RS13980	WP 010643500.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGKYLEVVISKLKADEKPKPKSVLEKEG	156
NTDB id 1381	A4U84	RS02010	WP 021113424.1	DLNNTSVKDDQLYLLDVKDFSTVVESVEVFRDSTTRTLVAYIDEDYSHDYRLTGKYLEVVKVGLKANEPDKSILDKEG	157
NTDB id 111192	ACMCNG	RS08505	WP 226691996.1	MRYLFLLFF.....ATFPVL.....AN	17
NTDB id 1347	HI 0435	AAC22094.1		MKGEFKKCGLFFLCFC.....CAFSSMAEF..KEN	28
NTDB id 1373	NTHI	RS02645	WP 011272046.1	MKKYFLKCGYFLVCF..LPLIVFANPKTDN	29
consensus					29
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NTDB id 1186	GTF74	RS01320	WP 000788426.1		
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPVQQVLDITLQVKGLDKRVDGNVILVAPKEELDRE	236
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPVQQVLDITLQVKGLDKRVDGNVILVAPKEELDRE	236
NTDB id 1381	A4U84	RS02010	WP 021113424.1	KLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPVQQVLDITLQVKGLDKRVDGNVILVAPKEELDRE	237
NTDB id 111192	ACMCNG	RS08505	WP 226691996.1	QQISISIRNAPTAEIISYLAETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKENGYYIHQAQEHKQHT	97
NTDB id 1347	HI 0435	AAC22094.1		QVFSIRLKAQPMVSSLLQELALVCGTNLIDDEIEGSLSLQLDKVDFDRILRSVAKNKGSLRVEDDYYLCKNSTGTFA	108
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ERFFIRLSQAPLAQTLEQLAFQCDVNLVIGDIEENKISLKLNNIDMPRLQITAKSKHITLNKDDGYIYNGSQSGKGQV	109
consensus					109
					***** * **!* ***** *!* ** !***** ***** * * *!* *****! ***** ***** * * *

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NTDB id 1186	GTF74	RS01320	WP 000788426.1		
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQAL.....EKARLAEELGDLKSEIIKNFAKASDIAAMIGGECNVNMLSERGSISIDERTNSLLIRELPDNI	307
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQAL.....EKARLAEELGDLKSEIIKNFAKASDIAAMIGGECNVNMLSERGSISIDERTNSLLIRELPDNI	307
NTDB id 1381	A4U84	RS02010	WP 021113424.1	KQAL.....EKQKLEKEIGELSSSEIIKVNFAKASDIAEMINGECNISMLSDRGSMIDERTNSLLIRELPENIEVI	308
NTDB id 111192	ACMCNG	RS08505	WP 226691996.1	TAQLVNALPKLITKTIKLHYSKASEVIESLT.KGGCNLLSESGYLHFDDRSNSIIVKDSAAASVKNF	162
NTDB id 1347	HI 0435	AAC22094.1		TAYVETSQAVANEAFKAMPSEPMLISRTVKLHFAKAEVNMKSLT.TGSGSLLSAQGSIAFDDRSNLLIQDEAKSVKM	187
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ACNLTITNEPHLVSHTVKLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRSNLLVIQDEPRSVQNI	174
consensus					174
					* * * *! * * * *! * * * *! * * * *! * * * *! * * * *! * * * *! * * * *! * * * *

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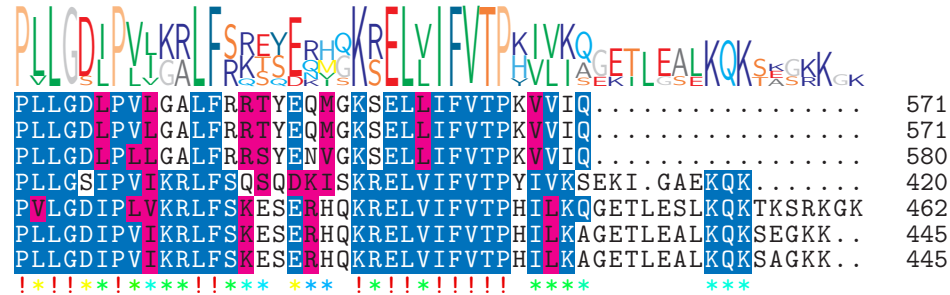
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NTDB id 1347 HI 0435 AAC22094.1
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consensus

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- non conserved
- similar
- $\geq 50\%$ conserved