

Accession	Sequence	Length
NTDB id 1186 GTF74 RS01320 WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA ¹ AVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175 A1552VC RS12105 WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA ² AVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408 DSB67 RS13980 WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...FVAAEEVQSNLSKSIDFRTNKDKDAVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 1381 A4U84 RS02010 WP 021113424.1		0
NTDB id 307125 DXH82 RS03715 WP 115911560.1		0
NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
consensus		

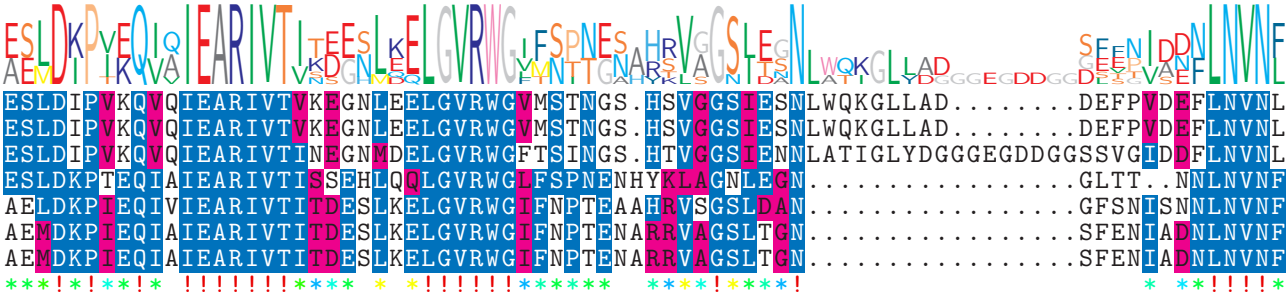
NTDB id 1186	GTF74 RS01320 WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKG.KYLEVVISKLKADEKPKPKSVLEKE	155
NTDB id 1175	A1552VC RS12105 WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKG.KYLEVVISKLKADEKPKPKSVLEKE	155
NTDB id 1408	DSB67 RS13980 WP 010643500.1	DLLNTSVKDDQLYLLDVKDFSTVVESVEVFRDTSTTRLVAYIDEDYSHDYRLTG.RYLEVKVGKLIKANEKAPDKSILDKE	156
NTDB id 1381	A4U84 RS02010 WP 021113424.1	MRYLFLFF.....ATFPVLA	16
NTDB id 307125	DXH82 RS03715 WP 115911560.1	MIKQKIKTKCGQFLMCFLILWTT.....YS..AAE	28
NTDB id 1347	HI 0435 AAC22094.1	MKKYFLKCGYFLVCFCLPLIV.....FANPKTD	28
NTDB id 1373	NTHI RS02645 WP 011272046.1	MKKYFLKCGYFLVCFCLPLIV.....FANPKTD	28
consensus		* * *	*

[illegible]

NTDB id 1186	GTF74	RS01320	WP 000788426.1	EKQAL E ... KARLAEELGDLKSEIKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREII	311
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	EKQAL E ... KARLAEELGDLKSEIKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREII	311
NTDB id 1408	DSB67	RS13980	WP 010643500.1	EKKQL E ... KQKLEKEIGELSSEI IKVNFAKASDIAEMINGECNISMLSDRGSMITIDERTNSLLIRELPENIEVIREII	312
NTDB id 1381	A4U84	RS02010	WP 021113424.1	T.....TAQLVNALPKLITKTIKLHYSKASEVTESLT.KGQGNNLSESGLYLFHDDRNSNIIVKDSAASVKNFQTQLI	166
NTDB id 307125	DXX82	RS03715	WP 115911560.1	AEKMA E PMAISGESLPSETPLVSATVTKLHFAKASDVMMKSLT.TGSGSLLSPSGTTITFDDRNSNVLLIQDDARSVKNIKLI	187
NTDB id 1347	HI 0435	AAC22094.1		GGVAGNLTTNEPHLVSHTVKLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRSNLLVIQDEPRSVQNIKLI	178
NTDB id 1373	NTHI	RS02645	WP 011272046.1	GGVAGNLTTNEPHLVSHTVKLHFAKASELMKSLT.TGSGSLLSSAGSITFDDRSNLLVIQDEPRFVQNIKLI	178
consensus				* * * * *	

logo

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NTDB id 1347 HI 0435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus



382
382
391
227
250
241
241

logo

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NTDB id 1175 A1552VC RS12105 WP 000788426.1
NTDB id 1408 DSB67 RS13980 WP 010643500.1
NTDB id 1381 A4U84 RS02010 WP 021113424.1
NTDB id 307125 DXX82 RS03715 WP 115911560.1
NTDB id 1347 HI 0435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus



461
461
470
304
329
320
320

logo

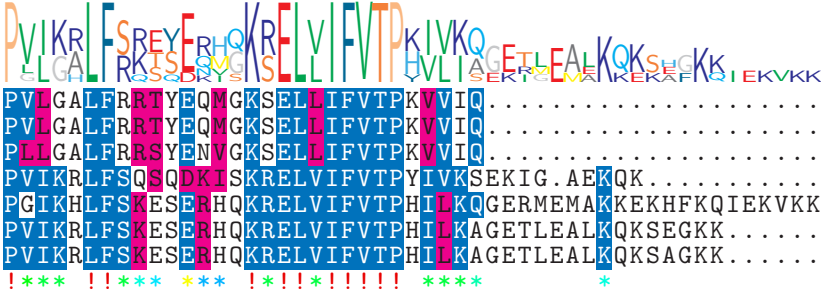
NTDB id 1186 GTF74 RS01320 WP 000788426.1
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NTDB id 1347 HI 0435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus



541
541
550
381
409
400
400

logo

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NTDB id 1175 A1552VC RS12105 WP 000788426.1
NTDB id 1408 DSB67 RS13980 WP 010643500.1
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NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus



571
571
580
420
460
445
445

☒ non conserved
☒ similar
☒ ≥ 50% conserved