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NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 306066 DV160 RS04030 WP 002220278.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
consensus

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[illegible]

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NTDB id 1402 DSB67 RS12670 WP 010643257.1
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NTDB id 306066 DV160 RS04030 WP 002220278.1
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consensus

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DRLATYKEKTEALKKAKIKKALYTPVSVIAVAIGLVFVMMIFVLP	EFKVFANFGAELPAETQVMSWQYVWVYLL	
EKVLTIRISQKAKIKKALYTPVSVIAVAIGLVFVMMIFVLP	EFKVFANFGAELPAETQVMSWQYVWVYLL	
ERLATYREKSEQLRAKVIKALYTPSMVVLVALGVSYLMLTMVLP	EFKVFANFGAELPAETQVMSWQYVWVYLL	232
ERLATYREKSEQLRAKVIKALYTPAMVVLVALGVSYLMLTRVLP	EFKVFANFGAELPAETQVMSWQYVWVYLL	231
DKLATYKEKTQAIRKKVKKTALYTPVSVIAVAIGLVFVMMIFVLP	EFKVFANFGAELPAETQVMSWQYVWVYLL	234
DKLATYKEKTQAIRKKVKKTALYTPVSVIAVAIGLVFVMMIFVLP	EFKVFANFGAELPAETQVMSWQYVWVYLL	234
DKLATYKEKTQAIRKKVKKTALYTPVSVIAVAIGLVFVMMIFVLP	EFKVFANFGAELPAETQVMSWQYVWVYLL	234
DKVATYKEKIEIKKKIKKALYTPAVMVVALLVITAGLLIYVVP	EFKVFANFGAELPAETQVMSWQYVWVYLL	231
DRVATYKEKTEALKKAKIKKAMNYP	IAVVLVAIVITATLLIKVVP	230
DRVATYKEKSELLKKIKKAMNYP	IAVVLVAIVITATLLIKVVP	233
DRVAVYKEKSEVLKKIKKAMKYP	ATVIVVAIVVITILLIKVVP	233
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consensus

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A L G F A I Y E L K L K R S K F R D R L P I C D I V K A I R T L A T F A A G Y P L V D L S T A G A G N L I Y E A T	
S F S Y E A C K L R K A N S F Q I R L K T S R L G L K F P I G V L S K A A I A K F S R T L A T S F A A G I P I L A S L K T T A K T S G N V H F E T A I N	
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SFVSLISGKIMSQRSDSFRLLAARSSLRVPILGPVLSKAAIAKFSRTLATSFSAGIPTILTALKTTSKTSGNMHYQLAIE	311
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NTDB id 1016 ACIAD RS01680 WP 004920476.1
consensus
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392
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logo

NTDB id 1169	A1552VC RS11075 WP 000648511.1	VVAMYLPIFNLMSV	LG.	408
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NTDB id 1198	PSJM300 03950 AFN76868.1	IIAMYLPIFQMG	SVV..	405
NTDB id 1059	ABD1 RS01610 WP 000279215.1	VIAMYLPIFQMG	SVV..	408
NTDB id 1016	ACIAD RS01680 WP 004920476.1	VIAMYLPIFQMG	SVV..	408
consensus		*****	*****	

VVAMYLPIFNLGSA
VVAMYLPINLMSVLG
VTAMYLPINFNLSVLG
LVAMYLPILFNLCNVVA
LVAMYLPILFNLCNVVA
LVAMYLPILFNLCNVVA
VVAMYLPILFKLGSAV
ITAMYLPILFQMGSSV.
VIAMYLPILFQMGSSV.
VIAMYLPILFQMGSSV.

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