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NTDB id 1284 ACE17W RS12370 WP 010946666.1	MRKIIAILIL..ISASISMAYSKDNSLISVKVLPPEPVRVIRDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 1257 GCO85 RS04855 WP 163621000.1	MRKIIAILIL..ISASISLAYSKDNSLISVKVLPPEPVRVIRDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id 1186 GTF74 RS01320 WP 000788426.1		0
NTDB id 1175 A1552VC RS12105 WP 000788426.1		0
NTDB id 1054242 BCV31 RS07765 WP 016787694.1		0
NTDB id 1408 DSB67 RS13980 WP 010643500.1		0
NTDB id 1193 PAKAF RS26665 WP 003161249.1	MNSGLSRLGIALIAA.MFAPALLAADIEKIDVAALPGDRVEIKLQFDEPVA.APRGYTIEQPARIALDLPGVQNKIGT..	76
NTDB id 1009 ACIAD RS15175 WP 004923730.1	MKQIIRQFSIGTVAYAIMQVASAQVAITNVVPMNIPGGTEIRVMFENGLPP.QPQAYQLEQPARLILDFDKAGQKIAK..	77
NTDB id 1039 HON27 RS01460 WP 031993779.1	MNHVFRQFSMGAVAIAIMQAAASQVSMTNIVPMQIAGGGTEIRVMFENGLPP.QPQAYQLENPSRLILDFDKAQQGKQ..	77
NTDB id 1052 ABD1 RS15850 WP 001017033.1	MNHVFRQFSMGAVAIAIMQAAASQVSMTNIVPMQIAGGGTEIRVMFENGLPP.QPQAYQLENPSRLILDFDKAQQGKQ..	77
consensus	* *	

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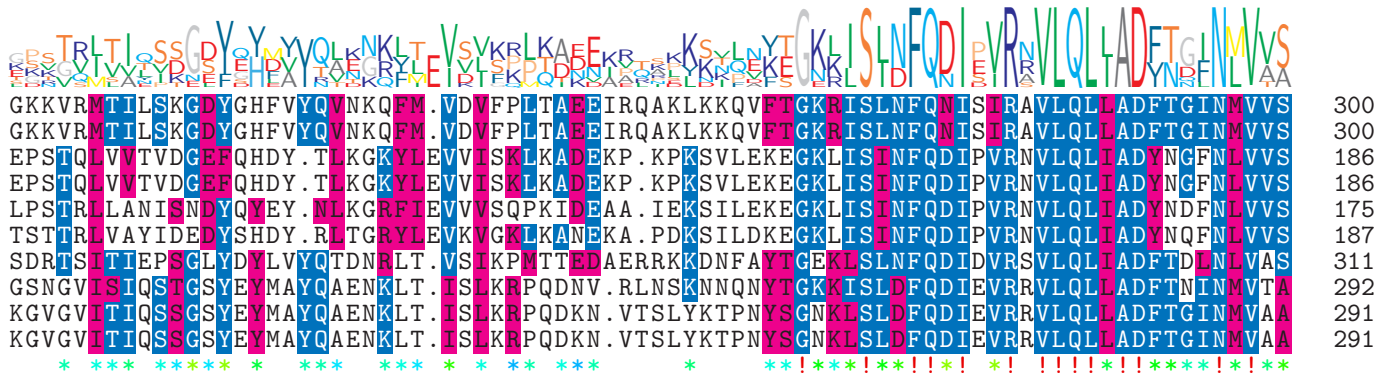
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NTDB id 1186 GTF74 RS01320 WP 000788426.1	MKNGLKT..YVAQTWLTLWVGLALCAS.....SMVF	29
NTDB id 1175 A1552VC RS12105 WP 000788426.1	MKNGLKT..YVAQTWLTLWVGLALCAS.....SMVF	29
NTDB id 1054242 BCV31 RS07765 WP 016787694.1	MNRGLQGIVLSVVLFF.....TV	18
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NTDB id 1193 PAKAF RS26665 WP 003161249.1	KNRELISVGNTRSVTVVEAKDRTRLIINLTTLSSYTTRVEGN...NLFVVVGNSPAGASVASTAPVKASPAPASYAQPIK	152
NTDB id 1009 ACIAD RS15175 WP 004923730.1	SLIPVTTNEASSIDVSSDDKRARVVVNLKDA GAF TTRVEGN...VFIKLINNIQSTAT.....PVAT..SA...	138
NTDB id 1039 HON27 RS01460 WP 031993779.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDTGAF TTRVEGN...TFILKINSAQTSNK.....PLPV..VS...	138
NTDB id 1052 ABD1 RS15850 WP 001017033.1	SKISVATNEASSVDVTSDDQRSRLTVNLKDA GAF TTRVEGN...TFILKINSAQTSNK.....PLPV..VS...	138
consensus	* *	

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NTDB id 1284 ACE17W RS12370 WP 010946666.1	TNQVVNAKHEIKRIDFRGIEKQGGRVLIDVSDIGIPIDVTQVGKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIITMQQE	221
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NTDB id 1175 A1552VC RS12105 WP 000788426.1	.SAESATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRK	108
NTDB id 1054242 BCV31 RS07765 WP 016787694.1	.MSHAETSNKLENIIDFRVNKDK EAVIIIE LATITAVVDVQKVQEGLSIELINTEVDDDKLYLLDVKDFATLVEGIEVEFRE	97
NTDB id 1408 DSB67 RS13980 WP 010643500.1	.AAEEVQSNLSKSIDFRITNKDKDAVIVVELVSPAATVDIKRVQEGLSIDLNTSVKDDQLYLLDVKDFSTPVESVEVFRD	109
NTDB id 1193 PAKAF RS26665 WP 003161249.1	PKPYVPAGRAIRNIDFQRGEKGEKNVVIDLSDPTLSPDIQEQQGKIRLDFAKTQLPDALRVRLDVKDFATPVQFVNASAQ	232
NTDB id 1009 ACIAD RS15175 WP 004923730.1	PAPQCGIGNIGFQRGAQGEGLVVVDLQGNTPVDVQQQCSKVVRFPGIIKIPTHLARRLNTTDFATPVASIDSYND	214
NTDB id 1039 HON27 RS01460 WP 031993779.1	AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQCSKVVRTITGKIPTHLARRLNVNDFATPVSSIDAYND	213
NTDB id 1052 ABD1 RS15850 WP 001017033.1	AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQCSKVVRTITGKIPTHLARRLNVNDFATPVSSIDAYND	213
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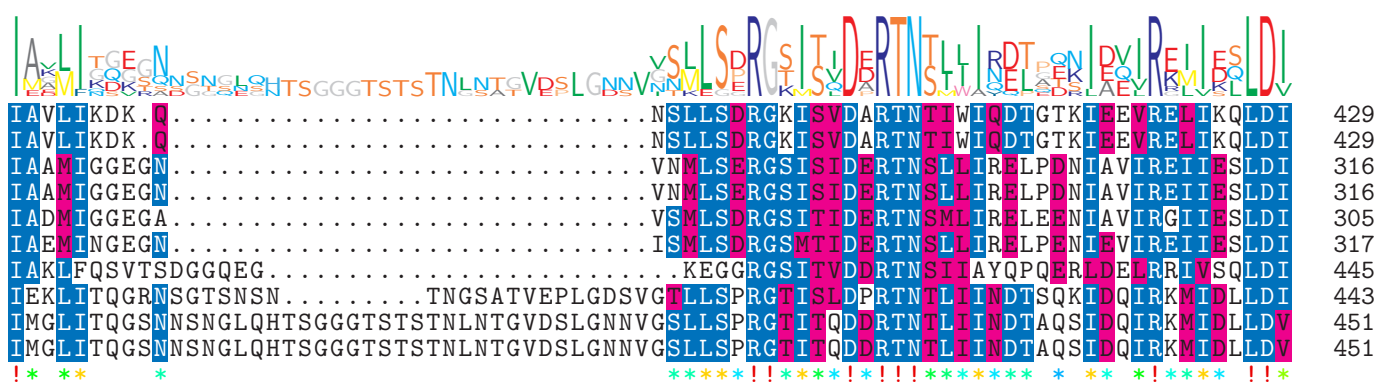
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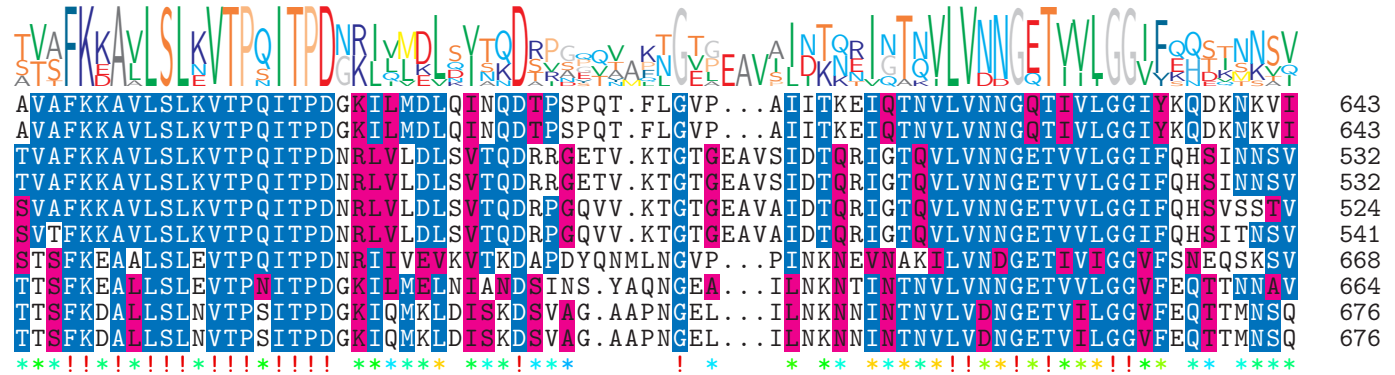
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X non conserved
X similar
X ≥ 50% conserved