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NTDB id 1009 ACIAD RS15175 WP 004923730.1	MNLYEYRRVAIAIMQ...VASAQVAITNVVPMNIPGQGTEIRVMFNGLPP.QPQAYQLEQPARLILDFDKA	71
NTDB id 1052 ABD1 RS15850 WP 001017033.1	MNHVFRQFSMGAVAIAIMQ...AASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKA	71
NTDB id 1039 HON27 RS01460 WP 031993779.1	MNHVFRQFSMGAVAIAIMQ...AASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKA	71
NTDB id 1284 ACE17W RS12370 WP 010946666.1	...MR...KIIAILILISASLSMAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFIST	70
NTDB id 1257 GC085 RS04855 WP 163621000.1	...MR...KIIAILILISASLSLAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFIST	70
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NTDB id 1408 DSB67 RS13980 WP 010643500.1		0
NTDB id 392949 GDV60 RS02850 WP 092232908.1		0
NTDB id 1193 PAKAF RS26665 WP 003161249.1	MNSGLSR....LGIALLAAMF...APALLAADLEKLDVAALPGDRVELKLQFDEPVAA.PRGYTIEQPARTALDLPGV	70
consensus	**** * * * * *	

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NTDB id 1009 ACIAD RS15175 WP 004923730.1	QQQLPSS...ITTYTNEASSVDVSDPMRLTYNLKPAASYTTRVEGNTLWLYNNAQSSSNLP	133
NTDB id 1052 ABD1 RS15850 WP 001017033.1	GQKLAKS..LIPVTTNEASSIDVSSDDKRARVVVNLDKAGAFTRVEGNVFILKINNIQSTATP.....	133
NTDB id 1039 HON27 RS01460 WP 031993779.1	QQGLKQS..KISVATNEASSVDVTSDDQRSRLTVNLDKAGAFTRVEGNVFILKINSAQTSNKP.....	133
NTDB id 1284 ACE17W RS12370 WP 010946666.1	QQGLKQS..KISVATNEASSVDVTSDDQRSRLTVNLDKAGAFTRVEGNVFILKINSAQTSNKP.....	137
NTDB id 1257 GC085 RS04855 WP 163621000.1	DMQIPSDQKTKTITQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSIYSL.VLNGKSNDF.....ENSK	137
NTDB id 1186 GTF74 RS01320 WP 000788426.1	DMQIPSDQKTKTITQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSVYSL.VLNGKSNDF.....ENSK	26
NTDB id 1175 A1552VC RS12105 WP 000788426.1	MKNGLKT..YVAQTW.....LTLWVGLALCA.....SS..	26
NTDB id 1408 DSB67 RS13980 WP 010643500.1	MKNGLKT..YVAQTW.....LTLWVGLALCA.....SS..	28
NTDB id 392949 GDV60 RS02850 WP 092232908.1	MKMRHGLTLTPRKLTGL.....FSLWFSLLLAP.....	0
NTDB id 1193 PAKAF RS26665 WP 003161249.1	QNKL..GTKNRELISVGNTRSVTVVEAKDRTRLIINLTTLSYTTTRVEGNNLFVVVGNSPAGASVASTAPVKASPAPASYA	148
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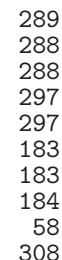
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NTDB id 1175 A1552VC RS12105 WP 000788426.1	..MVFSAESATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTADVADDKLYLLDVKDFSTPVESVE	105
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NTDB id 1193 PAKAF RS26665 WP 003161249.1	QPIKPKPYVPAGRAIRNIDFQRGEKGEGNVVIDLSDPTLSPDIQEQQGKIRLIDFAKTQLPDA LRVRLDVKDFATPVQFVN	
consensus	* * * * *	

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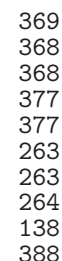
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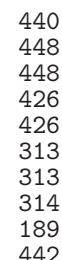
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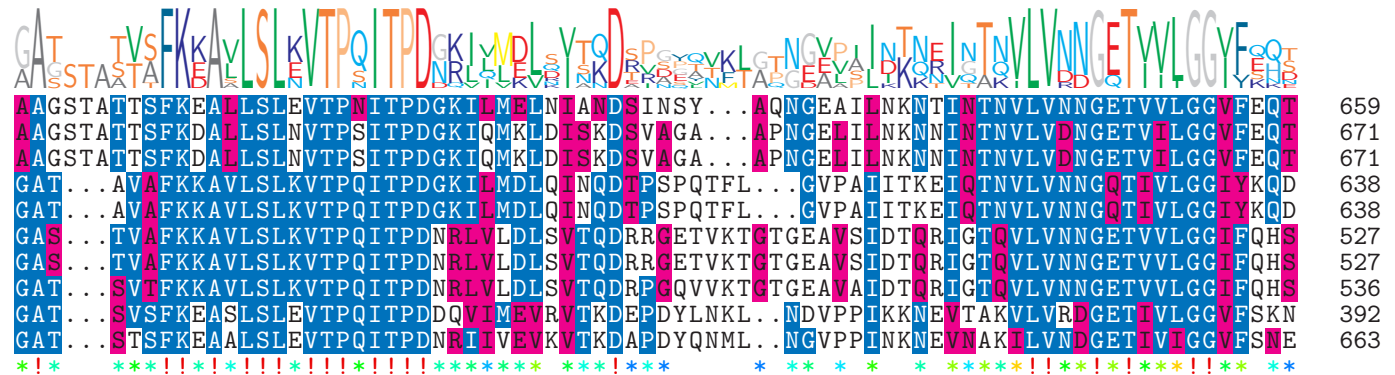
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consensus

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NTDB id 1186 GTF74 RS01320 WP 000788426.1	INNSVDKVPLLGDLPLVGLALFRRTYEQMGKSELLIFVTPKVVIIQ.....	571
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NTDB id 392949 GDV60 RS02850 WP 092232908.1	QSKVVDKVPFLGDPYLLGRLFRRDVVAQRKSELLVFLTPRIMNNQAIAVSR.....	443
NTDB id 1193 PAKAF RS26665 WP 003161249.1	QSKSVEKVPFLGELPYLLGRLFRRDVTVTDRKNELLVFLTPRIMNNQAIAIGR.....	714
consensus	*****!***!!**!*!!** * * !!!*!!***** *	

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved