

logo

NTDB id 1009 ACIAD RS15175 WP 004923730.1	MNLYEYRQFSMGAKIYLLILLLISASLSMAYLEKDNMLESYKPLAETAEENGTRLEDEM SKLLKQL ASFFITQK AATYXLDLST	71
NTDB id 1039 HON27 RS01460 WP 031993779.1	MKQIIRQFSIGTVAYAIMQV.....ASAQVAITNVVPMNIPGGGTEIRVMFNGLPP.QPQAYQLEQPARLILDFDKA	71
NTDB id 1052 ABD1 RS15850 WP 001017033.1	MNHVFRQFSMGAVAIAIMQA.....ASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKA	71
NTDB id 1186 GTF74 RS01320 WP 000788426.1	MNHVFRQFSMGAVAIAIMQA.....ASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKA	0
NTDB id 1175 A1552VC RS12105 WP 000788426.1		0
NTDB id 1408 DSB67 RS13980 WP 010643500.1		0
NTDB id 1257 GC085 RS04855 WP 163621000.1	M.....RKIITAILILISASLSLAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFIST	70
NTDB id 1284 ACE17W RS12370 WP 010946666.1	M.....RKIITAILILISASLSMAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFIST	70
NTDB id 153764 AKN93 RS01540 WP 414820372.1		0
NTDB id 1193 PAKAF RS26665 WP 003161249.1	MNSGLS.....RLGITALLAAMFA....PALLAADLEKLDVAALPGDRVELKLQFDEPVAA.PRGYTIEQPARTALDLPGV	70
consensus	**** * * * * *	

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NTDB id 1009 ACIAD RS15175 WP 004923730.1	QQQLPSSITLYTNEASSVDVSDDDRRRLIYNLKRPSLSYTRVEGNELVWVSKQTSNKP	133
NTDB id 1039 HON27 RS01460 WP 031993779.1	GQKLAKS..LIPVTTNEASSIDVSSDDKRARVVVN LKDA GAFTTRVEGNVFILKINNIQSTATP.....	133
NTDB id 1052 ABD1 RS15850 WP 001017033.1	QQGLKQS..KISVATNEASSVDVTSDDQRSRLTVNLKDTGAFTTRVEGNTFILKINSAQTSNKP.....	133
NTDB id 1186 GTF74 RS01320 WP 000788426.1	QQGLKQS..KISVATNEASSVDVTSDDQRSRLTVNLKDA GAFTTRVEGNTFILKINSAQTSNKP.....	24
NTDB id 1175 A1552VC RS12105 WP 000788426.1	MKNGLKT..YVAQTWLTWL.....VGLALCA	24
NTDB id 1408 DSB67 RS13980 WP 010643500.1	MKNGLKT..YVAQTWLTWL.....VGLALCA	28
NTDB id 1257 GC085 RS04855 WP 163621000.1	MKMRHGLTLTPRKLTLGLFSLW.....FSLLLAP	137
NTDB id 1284 ACE17W RS12370 WP 010946666.1	DMQIPSDQKTKTITQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSVYSLVLNGKSNDL.F.....ENSK	137
NTDB id 153764 AKN93 RS01540 WP 414820372.1	DMQIPSDQKTKTITQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSIYSLVLNGKSNDL.F.....ENSK	8
NTDB id 1193 PAKAF RS26665 WP 003161249.1	QNKLL..GTKNREL SVGNTRSVTVVEAKDRTRLIINTLTLSSYTTRVEGNLFFVVGNSPAGASVASTAPVKASPAPASYA	148
consensus	* * * * *	

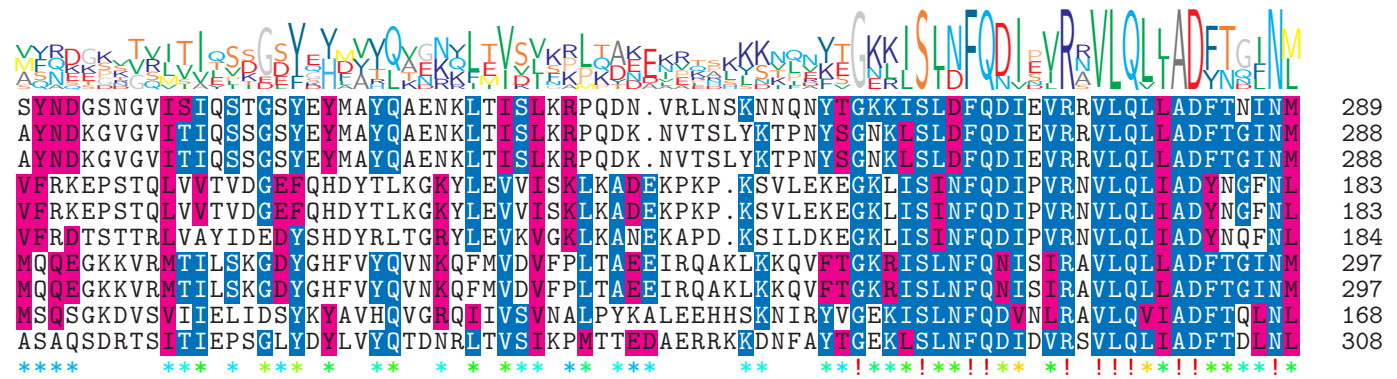
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NTDB id 1186 GTF74 RS01320 WP 000788426.1	SSMVFSAESATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVE	104
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NTDB id 1257 GC085 RS04855 WP 163621000.1	EVFITNQVVNAKHEIKRIDFRGIEKQGGRLIDVSDTGIPIDVTQVGKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIIT	217
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NTDB id 153764 AKN93 RS01540 WP 414820372.1	SELTIIYSKVVSNQTVENLYFKRGQGEGNVILTLTEPDVRFVKEESDKIHLVLLNTSLSKQINTHLKTKEFATPVQLIS	88
NTDB id 1193 PAKAF RS26665 WP 003161249.1	QPIKPKPYVPAGRAIRNIDFQRGEKGEGNVIDLSPTLSPDIQEQQGKIRLIDFAKTQLPDALRVRLDVKDFATPVQFVN	228
consensus	* * * * *	

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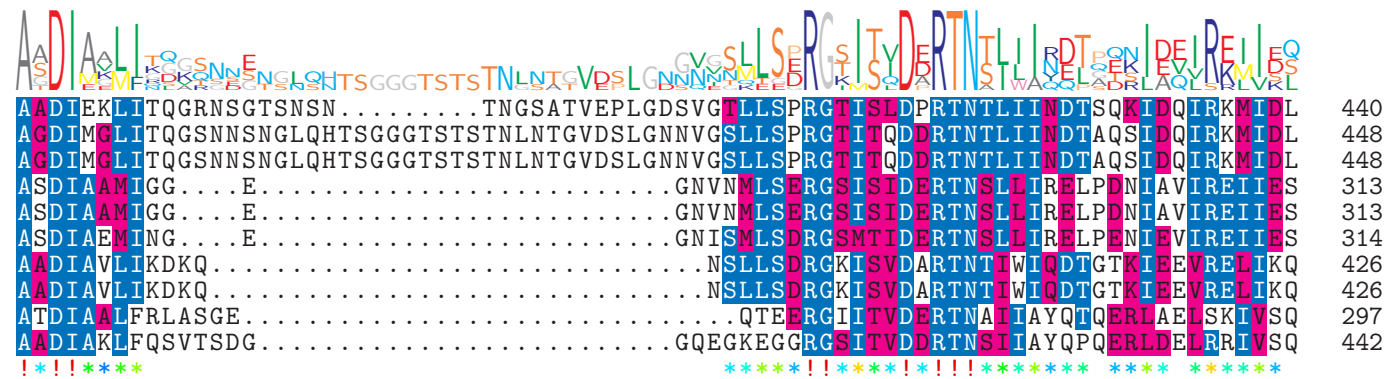
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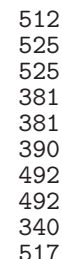
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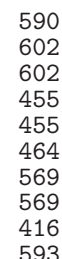
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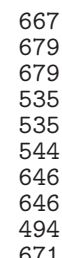
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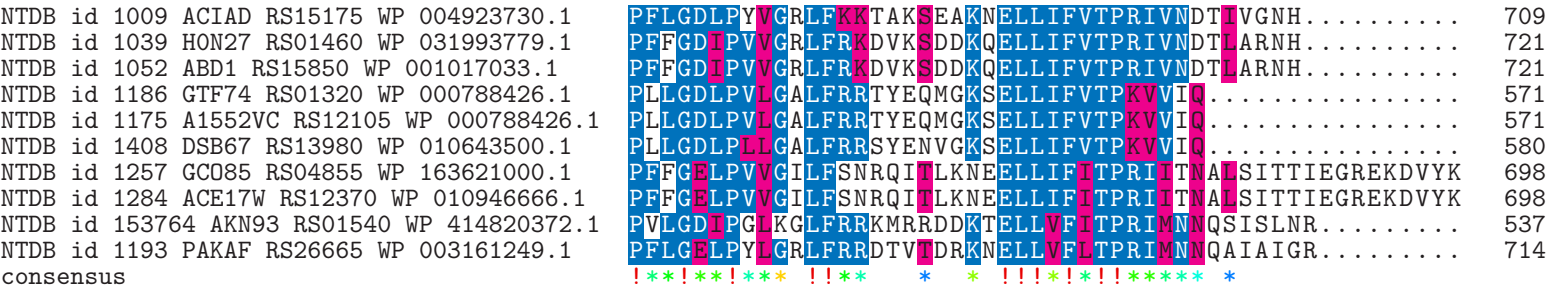
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consensus

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logo



☐ non conserved
☒ similar
☒ ≥ 50% conserved