

logo

NTDB id 1381 A4U84 RS02010 WP 021113424.1	MNSGLSRLGIALLAAMFAPALLAADLEKLDVAALPGDRVELKLQFDEPVAAAPRCYTI EQPARIALDLPGVQNKLGTKNRE	0
NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
NTDB id 73947 NMEC018 RS14430 WP 000816011.1		0
NTDB id 1433 BW25113 RS17570 WP 000815987.1		0
NTDB id 1193 PA5040 NP 253727.1	MNSGLSRLGIALLAAMFAPALLAADLEKLDVAALPGDRVELKLQFDEPVAAAPRCYTI EQPARIALDLPGVQNKLGTKNRE	80
NTDB id 1186 GTF74 RS01320 WP 000788426.1		0
NTDB id 1175 A1552VC RS12105 WP 000788426.1		0
NTDB id 1408 DSB67 RS13980 WP 010643500.1		0
consensus		

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NTDB id 1381 A4U84 RS02010 WP 021113424.1	LSVGNTRSVTVWEAKDRTRLIINLTALSSYTTRVEGNNLFVVVGNSPAGASVASAAPVKASPAPASYAQPIKPKPYVPAG	0
NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
NTDB id 73947 NMEC018 RS14430 WP 000816011.1		0
NTDB id 1433 BW25113 RS17570 WP 000815987.1		0
NTDB id 1193 PA5040 NP 253727.1	LSVGNTRSVTVVEAKDRTRLIINLTALSSYTTRVEGNNLFVVVGNSPAGASVASAAPVKASPAPASYAQPIKPKPYVPAG	160
NTDB id 1186 GTF74 RS01320 WP 000788426.1	MKNGLKT..YVAQTW.....LTLWVGLALCAS.....S....MVFSAESATA	36
NTDB id 1175 A1552VC RS12105 WP 000788426.1	MKNGLKT..YVAQTW.....LTLWVGLALCAS.....S....MVFSAESATA	36
NTDB id 1408 DSB67 RS13980 WP 010643500.1	MKMRHGLTLTPRKLTLGL.....FSLWFSLLLAP.....FVAAEEVQS	37
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NTDB id 1381 A4U84 RS02010 WP 021113424.1	NQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	0
NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
NTDB id 73947 NMEC018 RS14430 WP 000816011.1		0
NTDB id 1433 BW25113 RS17570 WP 000815987.1		0
NTDB id 1193 PA5040 NP 253727.1	RAIRNIDFQRGEKGEGNVVIDLSDP TLSPDIQEQGGKIRLDFAKTQLPDALRVRLDVKDFATPVQFVNASAQSDRTSITI	240
NTDB id 1186 GTF74 RS01320 WP 000788426.1	NQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	116
NTDB id 1175 A1552VC RS12105 WP 000788426.1	NQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	116
NTDB id 1408 DSB67 RS13980 WP 010643500.1	NSLKSIDFR TNKDKDAVIVVELVSPAAIVDIKRVQEGLSIDL LNTSVKDDQLYLLDVKDFSTVVESVEVFRDTS TTRLVA	117
consensus		

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NTDB id 1381 A4U84 RS02010 WP 021113424.1	MRYLFLLFFAT.....FPVLANQQISUSIRNAPTAEIISYLAZETGKNITISDEIKDTKN	55
NTDB id 1347 HI 0435 AAC22094.1	MKKYFLKCGYFLVCFCLPLI.....VFANPKTDNERFFIRLSQAPLAQTLQQLAFQQDVNLVIGDILENKIS	67
NTDB id 1373 NTHI RS02645 WP 011272046.1	MKKYFLKCGYFLVCFCLPLI.....VFANPKTDNECFIRLSQAPLAQTLQQLAFQQDVNLVIGERLEGNIS	67
NTDB id 73947 NMEC018 RS14430 WP 000816011.1	MKQWIAADLL.ML.....IPGVAAAKPQKVITLMDVDPVAVLQVLAQEKLNLVVSPDVSGTIVS	59
NTDB id 1433 BW25113 RS17570 WP 000815987.1	MKQWIAADLL.ML.....IPGVAAAKPQKVITLMDVDPVAVLQVLAQEKLNLVVSPDVSGTIVS	59
NTDB id 1193 PA5040 NP 253727.1	EPISGLYDYL VYQTDNRLTVSILK.PMTTEDAERRKKDNFAYTGEKLSINFDQIDVRSVLQQLADFTDLNLVASD TVQGNIT	319
NTDB id 1186 GTF74 RS01320 WP 000788426.1	TVDGEFQHDYTLKGKYLEVVVIS.KLK.ADEKPKPKSVLEKEGKLSINFDQIPVRNVLQLIADYNGFNLVVSDSVVGNLT	194
NTDB id 1175 A1552VC RS12105 WP 000788426.1	TVDGEFQHDYTLKGKYLEVVVIS.KLK.ADEKPKPKSVLEKEGKLSINFDQIPVRNVLQLIADYNGFNLVVSDSVVGNLT	194
NTDB id 1408 DSB67 RS13980 WP 010643500.1	YIDEDYSHDYRLTGRYLEVKG.KLK.ANEKAPDKSLDKEGKLSINFDQIPVRNVLQLIADYNGFNLVVSDSVVGNLT	195
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consensus



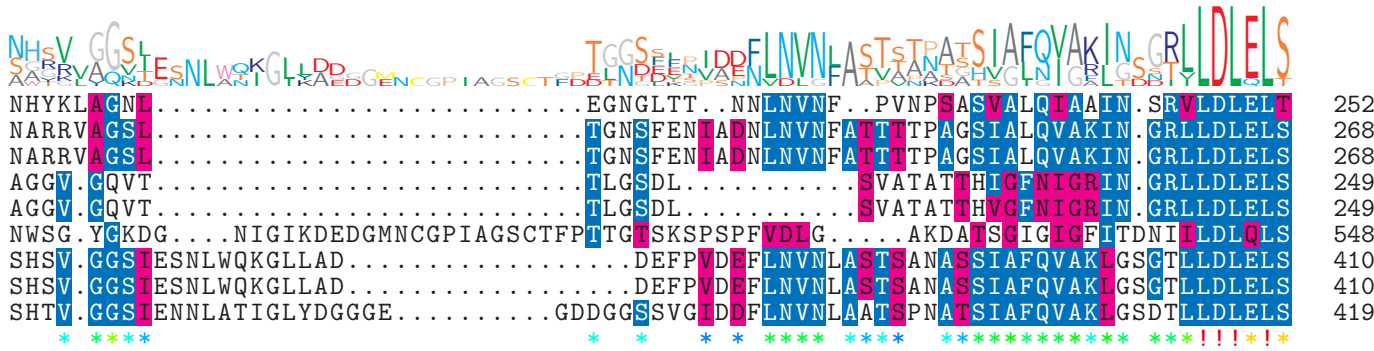
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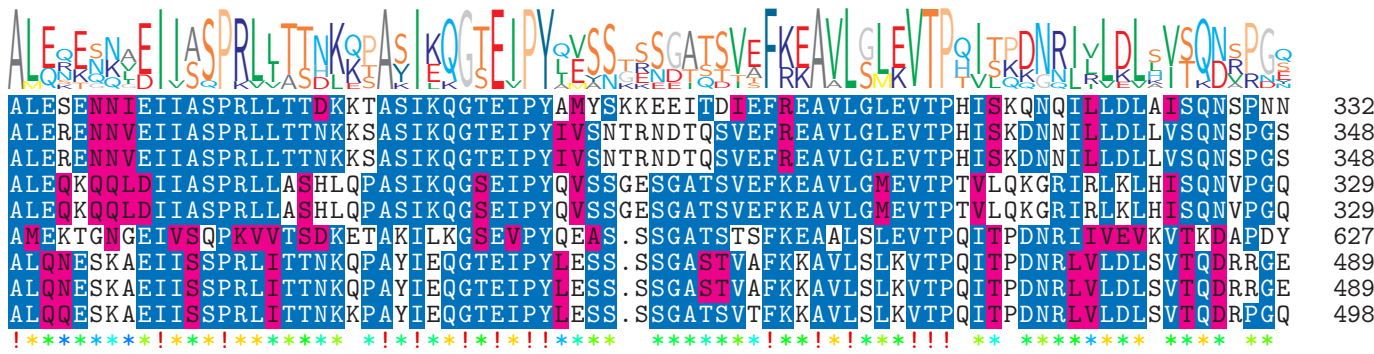
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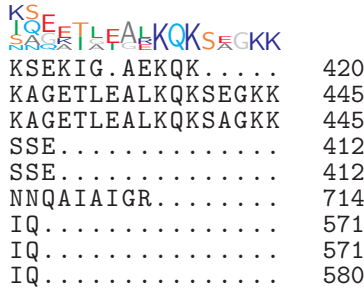
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consensus



X non conserved
X similar
X ≥ 50% conserved