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NTDB id 1347 HI 0435 AAC22094.1	MNSGLSRLGIALLAAMFAPALLAADLEKLDVAALPGDRVELKLQFDEPVAAPRGYTTIEQPARIALDLPGVQNKLGTKNRE	0
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NTDB id 1193 PA5040 NP 253727.1	MNSGLSRLGIALLAAMFAPALLAADLEKLDVAALPGDRVELKLQFDEPVAAPRGYTTIEQPARIALDLPGVQNKLGTKNRE	80
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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 1347 HI 0435 AAC22094.1	LSVGNTRSVTVWEAKDKIRITNLRASISLTTTRVEGNNLFVVGNSPAGASVASAAPVKASPAPASYAQPIKPKPYVPAG	0
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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	MKMRHGLTLTPRKLTGL.....FSLWFSLLLAP.....FVAAEEVQS	37
consensus		

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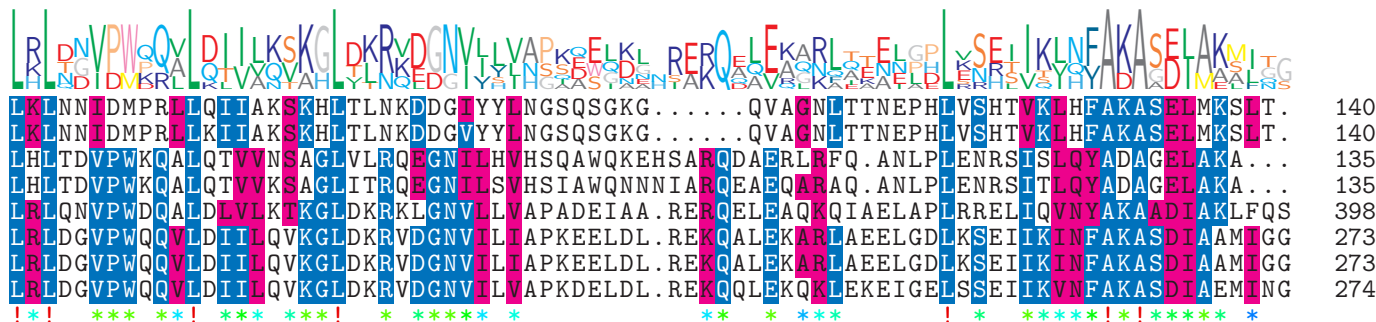
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NTDB id 1175 A1552VC RS12105 WP 000788426.1	NQLENIDFRVNKEKAAYVVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLV	116
NTDB id 1408 DSB67 RS13980 WP 010643500.1	NSLKSIDFRTNKDKDAVIVVELVSPAIVDIKRVQEGLSIDLLNTSVKDDQLYLLDVKDFSTVVESVEVFRDTSTTRLVA	117
consensus		

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NTDB id 1347 HI 0435 AAC22094.1	MKKYFLKCGYFLVCFCLPLIVF.....ANPKTDNERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKIS	67
NTDB id 1373 NTHI RS02645 WP 011272046.1	MKKYFLKCGYFLVCFCLPLIVF.....ANPKTDNECFIRLSQAPLAQTLEQLAFQQDVNLVIGERLEGNIS	67
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NTDB id 1186 GTF74 RS01320 WP 000788426.1	TVDGEFQHDYTLKG.KYLEVVISKLK.ADEKPKPKSVLEKEGKLISINFDIPVRNVQLQIADYNGFNLVVSDDSVVGNLT	194
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NTDB id 1408 DSB67 RS13980 WP 010643500.1	YIDEDYSHDYRLTG.RYLEVVKGLK.ANEKAPDKSILDKEGKLISINFDIPVRNVQLQIADYNGFNLVVSDDSVVGNLT	195
consensus	* * ** * * * ** *** **** **!* *!* * !!!*** * ****	

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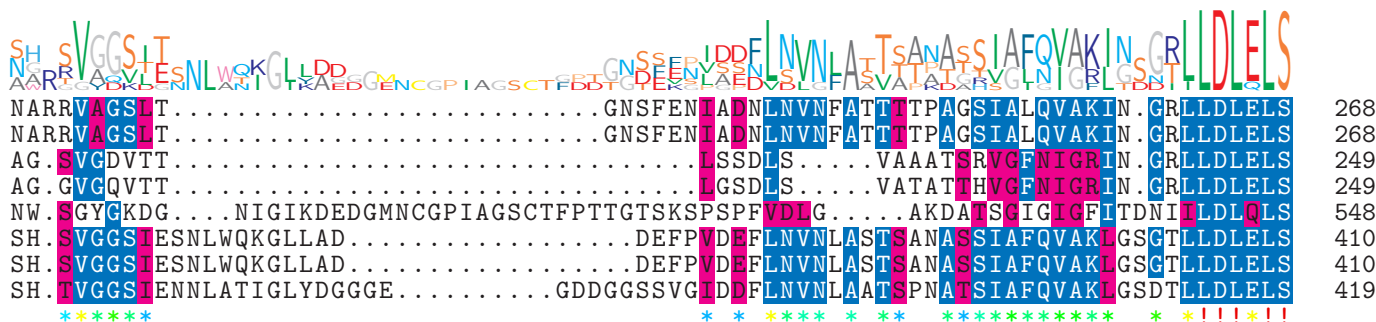
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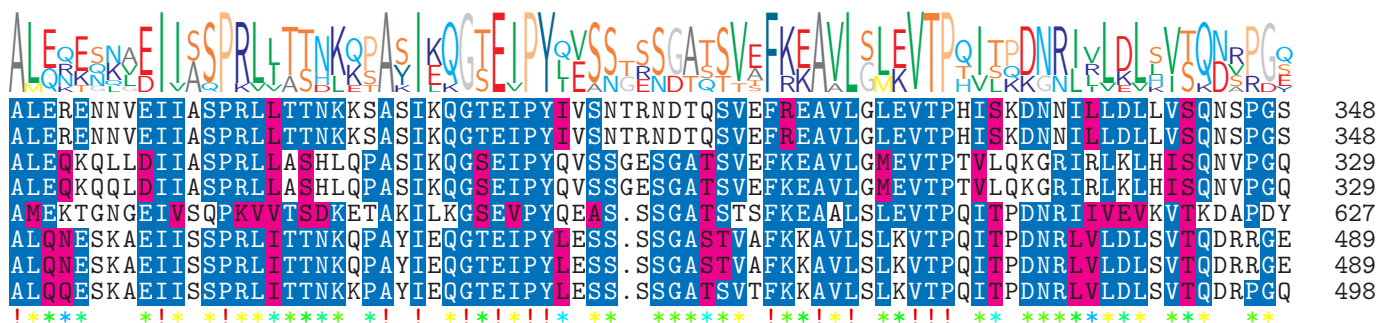
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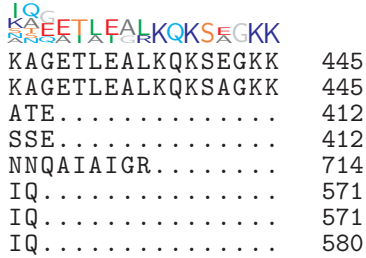
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NTDB id 1408 DSB67 RS13980 WP 010643500.1
consensus



X non conserved
X similar
X ≥50% conserved