

logo

NTDB id 119301 AU66 RS18550 WP 000016747.1	MSGRLLDVAVPLNSLTGVGAAQSSSKLAKIGLHTVQ.....DLLLHLPLRYEDRTH...LY.....	51
NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
NTDB id 1433 BW25113 RS17570 WP 000815987.1		0
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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 119301 AU66 RS18550 WP 000016747.1	.PIGELLPGIYATVEGEVLNLCNITFGGRRMMTC.QISDGSGLT..MRFFNFNAAMK.NSLATGRRVLAYGEAKRGKYGA	126
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NTDB id 1186 GTF74 RS01320 WP 000788426.1	MKNGLKT..YVAQTW.....LTLW.....VGLALCA.....	24
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NTDB id 1408 DSB67 RS13980 WP 010643500.1	MKMRHGLTLTPRKLTGL.....FSLW.....FSLLLAP.....	28
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NTDB id 1175 A1552VC RS12105 WP 000788426.1	SS....MVFSAESATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSIELLKTD	82
NTDB id 1408 DSB67 RS13980 WP 010643500.1	FVAEEVQSNSLKSIDFRTNKDKDAVIVVELVSPAAIVDIKRVQEGLSIDLLNTS	83
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NTDB id 1373 NTHI RS02645 WP 011272046.1	MKKYFLKCGYFLVCFCLPLIV.....FANPKT	27
NTDB id 1433 BW25113 RS17570 WP 000815987.1	MKQWIAALLML.....IPGVQAA	19
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NTDB id 1186 GTF74 RS01320 WP 000788426.1	VADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEGFQHDYTLKGKYLEVVISKLK.ADEKPK.....PKSVLEK	154
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NTDB id 1408 DSB67 RS13980 WP 010643500.1	VKDDQLYLLDVKDFSTPVESVEVFRDTSSTRIVAYIDEDYSHDYRLTGRIYLEVVGKILK.ANEKAP.....DKSILDK	155
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NTDB id 1175 A1552VC RS12105 WP 000788426.1	EGKLSISINFQDIPVRNVLQLIA....DYN	228
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consensus	*****	

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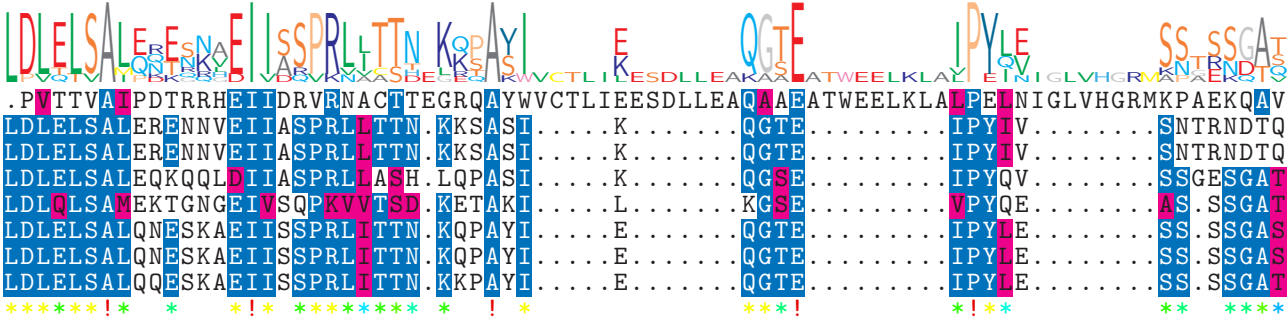
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NTDB id 119301 AU66 RS18550 WP 000016747.1	ENARRVAGSL	456
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NTDB id 1175 A1552VC RS12105 WP 000788426.1	GSH.SVGGSIENSLWQKGLLAD	404
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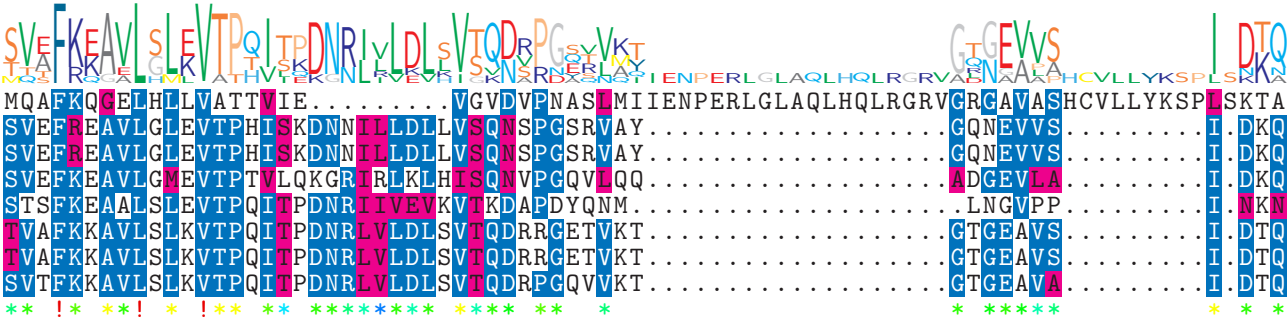
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NTDB id 1408 DSB67 RS13980 WP 010643500.1
consensus



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NTDB id 119301 AU66 RS18550 WP 000016747.1
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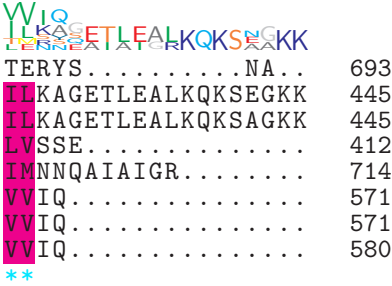
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consensus



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NTDB id 1347 HI 0435 AAC22094.1
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consensus



693
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445
412
714
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571
580

 non conserved
 similar
 $\geq 50\%$ conserved