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	MSVLTICDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 998000 ABET68 RS03385 WP 014616069.1	0
NTDB id 1110 NGFG RS09235 WP 003689817.1	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus		

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	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPVPPPSG	
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPVPPPSG	160
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NTDB id 998000 ABET68 RS03385 WP 014616069.1	0
NTDB id 1110 NGFG RS09235 WP 003689817.1	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus		

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	EGQKDLKLGELLQKGI SREALEEALVEQEKTCDLLGRILVRKGLPEEALYRALAEQKLEFLESTEGIVDPDPSAALLL	
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NTDB id 998000 ABET68 RS03385 WP 014616069.1	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus		

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	LRSDALRYGAVPIGFQNGEVEVWLSDP RHKEAVAQLLNRP ARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	
NTDB id 1028 TT RS08215 WP 011173991.1	LRSDALRYGAVPIGFQNGEVEVWLSDP RHKEAVAQLLNRP ARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	320
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consensus		

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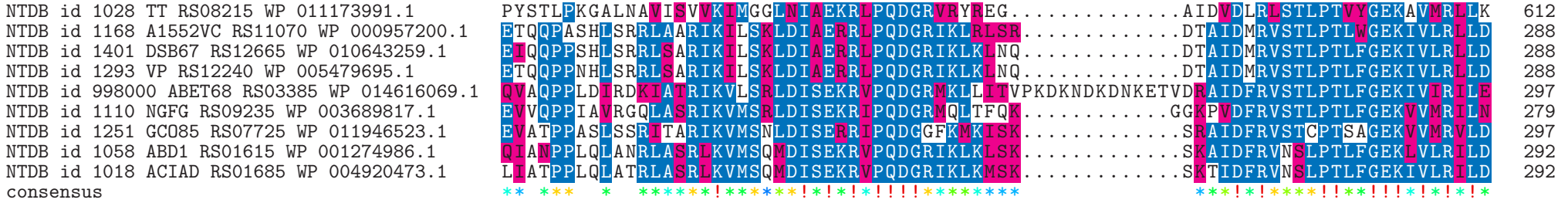
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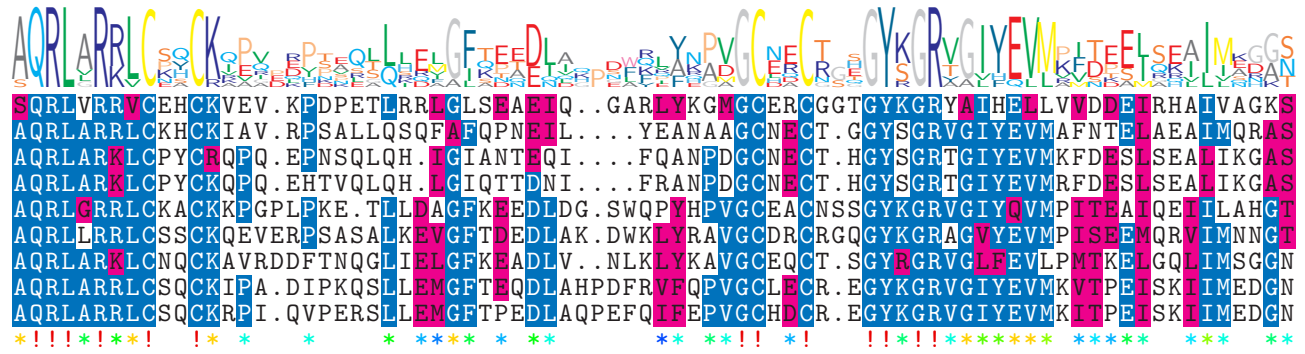
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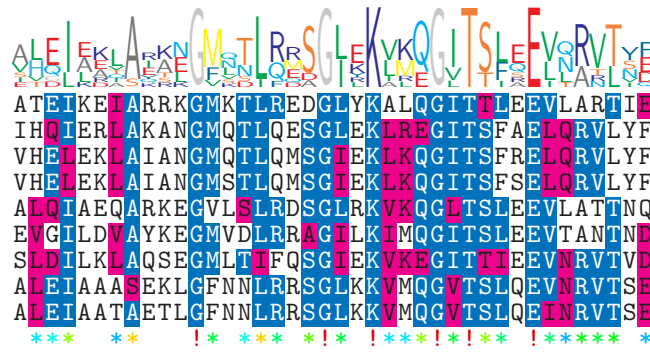
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 non conserved
 similar
 $\geq 50\%$ conserved