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	MSXLSLIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
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NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAILLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1293 VP RS12240 WP 005479695.1	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus	0

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	RSIRPQTALNLQAFPPFALE..GDRRLVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGA	157
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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLISLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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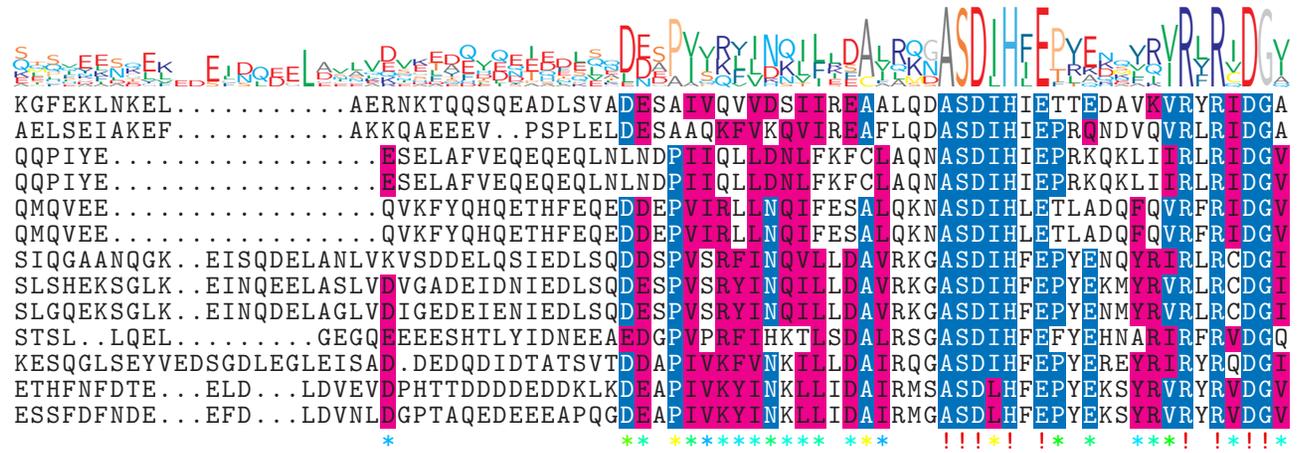
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consensus*.....**.....*.....*.....*.....***.....*	

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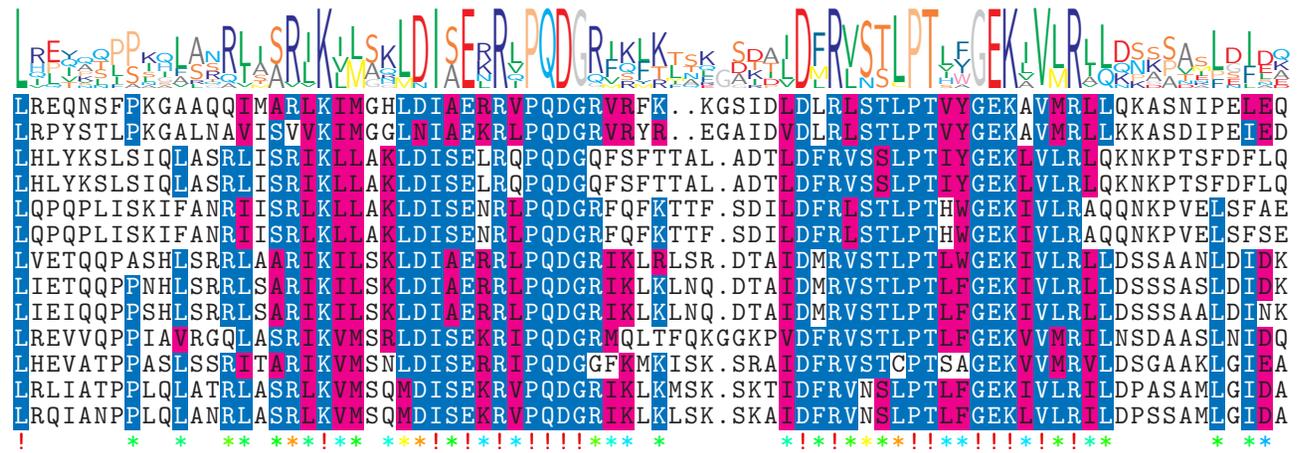
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 consensus



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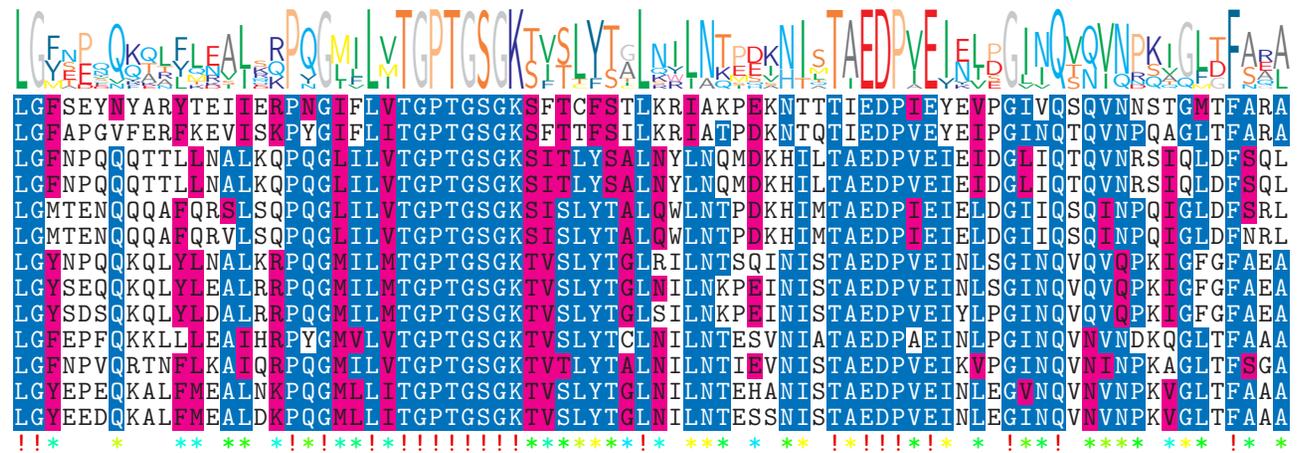
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 consensus



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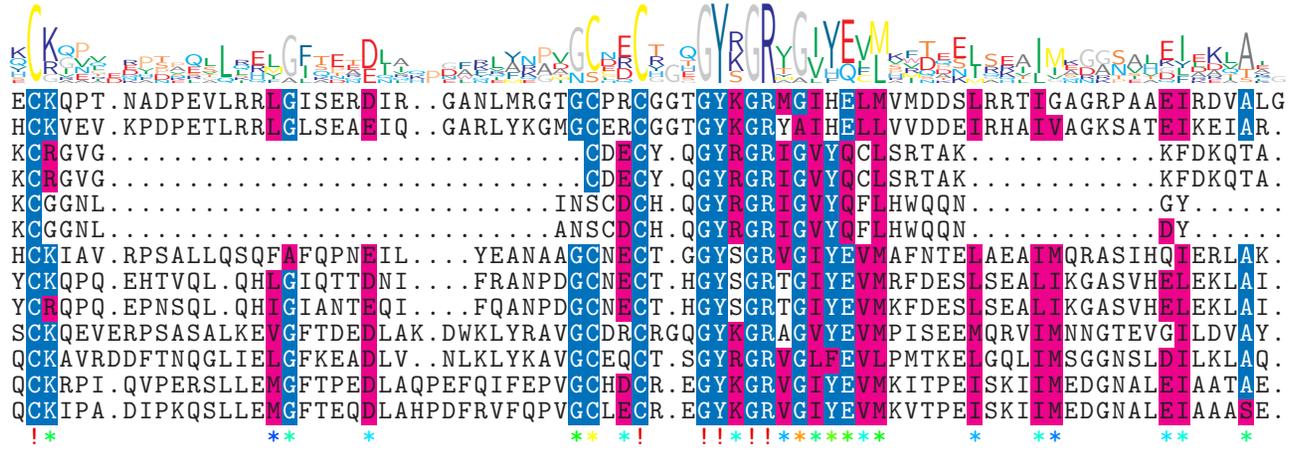
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 consensus



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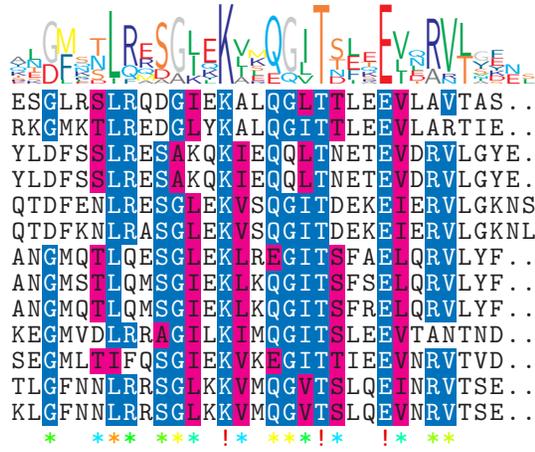
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 consensus



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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 consensus



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461
464
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561
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574
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570

 non conserved
 similar
 $\geq 50\%$ conserved