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NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus	0

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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTL SLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus	0

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NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVPDPSAALLL	240
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus	0

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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
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consensus	0

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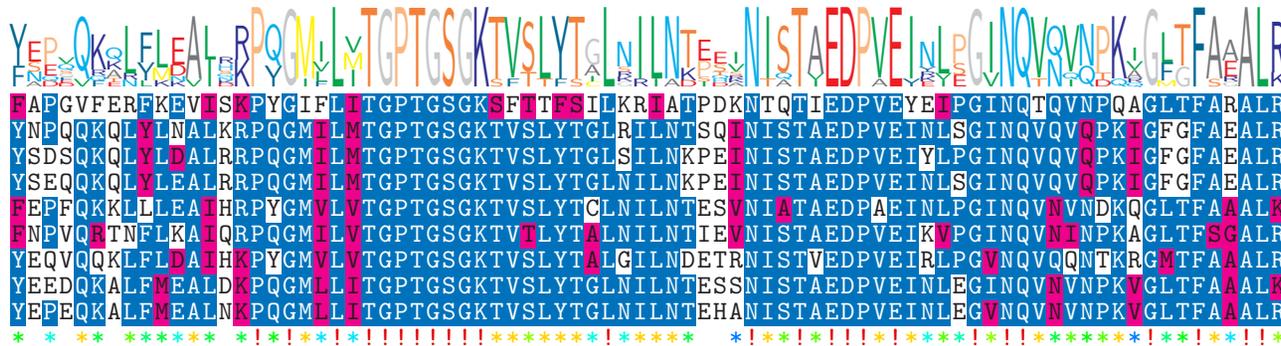
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 NTDB id 1251 GC085 RS07725 WP 011946523.1 QGLSEYVEDSG.DLEGLEISADEDDQDID.TATSVTDDAPIVFKVVKVLLDAIRKGDASDIHFEPYEREYRIRYRQDGILH 230
 NTDB id 367748 FHQ07 RS05755 WP 139715908.1 LADA...DND.GLEGLDVSGGDEDLSNDSGVDAAGDDTPVVKVVKVLLDAIRKGDASDIHFEPYETEYRVRRLRIDGILK 229
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 NTDB id 367748 FHQ07 RS05755 WP 139715908.1 TVKRVPVKLHQRIAAVVKVMAQLDIAEKRVQDGRIKLNLKSK.TKQIDFRVSTLPTLWGEKIVMRLDGSAAKLGIDKLG 308
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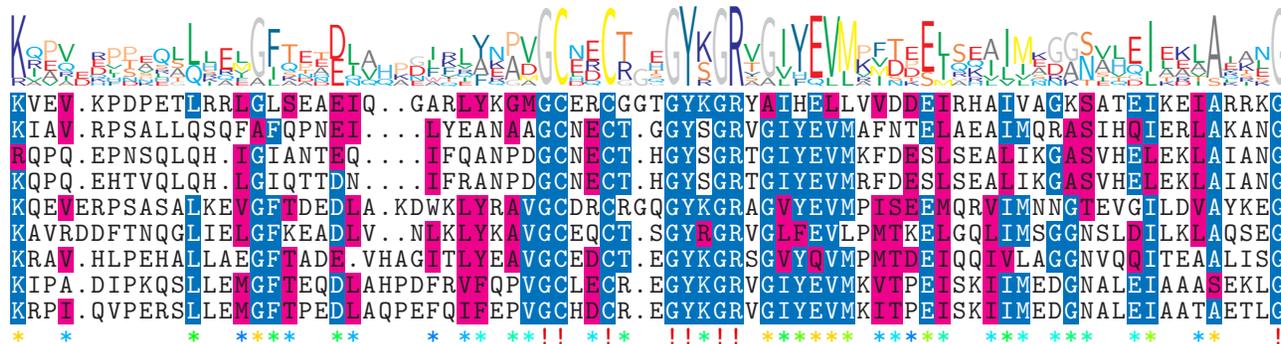
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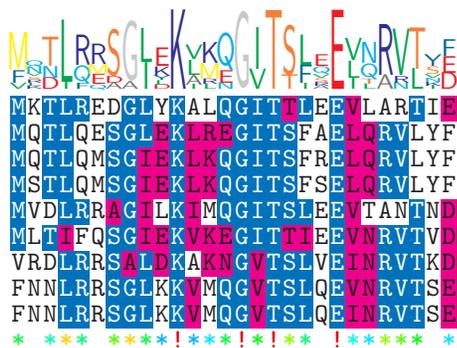
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 non conserved
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