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NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTI ^{CD} KRLGAALLDAGLLTDEEL ^{QR} ALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRF ^{GI} PLVELHRVEIPPKVK	80
NTDB id 1054276 BCV29 RS07470 WP 016794572.1	MSVLTI ^{CD} KRLGAALLDAGLLTDEEL ^{QR} ALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRF ^{GI} PLVELHRVEIPPKVK	0
NTDB id 1168 A1552VC RS11070 WP 000957200.1	MSVLTI ^{CD} KRLGAALLDAGLLTDEEL ^{QR} ALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRF ^{GI} PLVELHRVEIPPKVK	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	MSVLTI ^{CD} KRLGAALLDAGLLTDEEL ^{QR} ALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRF ^{GI} PLVELHRVEIPPKVK	0
NTDB id 1293 VP RS12240 WP 005479695.1	MSVLTI ^{CD} KRLGAALLDAGLLTDEEL ^{QR} ALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRF ^{GI} PLVELHRVEIPPKVK	0
NTDB id 1110 NGFG RS09235 WP 003689817.1	MSVLTI ^{CD} KRLGAALLDAGLLTDEEL ^{QR} ALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRF ^{GI} PLVELHRVEIPPKVK	0
NTDB id 1251 GC085 RS07725 WP 011946523.1	MSVLTI ^{CD} KRLGAALLDAGLLTDEEL ^{QR} ALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRF ^{GI} PLVELHRVEIPPKVK	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MSVLTI ^{CD} KRLGAALLDAGLLTDEEL ^{QR} ALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRF ^{GI} PLVELHRVEIPPKVK	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MSVLTI ^{CD} KRLGAALLDAGLLTDEEL ^{QR} ALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRF ^{GI} PLVELHRVEIPPKVK	0
consensus	MSVLTI ^{CD} KRLGAALLDAGLLTDEEL ^{QR} ALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRF ^{GI} PLVELHRVEIPPKVK	0

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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLD ^{TL} SL ^{EE} VEDLTGLVVEPYQT ^{TK} SAFLYALAKHYPELGLPVP ^{PP} PSG	160
NTDB id 1054276 BCV29 RS07470 WP 016794572.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLD ^{TL} SL ^{EE} VEDLTGLVVEPYQT ^{TK} SAFLYALAKHYPELGLPVP ^{PP} PSG	0
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLD ^{TL} SL ^{EE} VEDLTGLVVEPYQT ^{TK} SAFLYALAKHYPELGLPVP ^{PP} PSG	0
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NTDB id 1293 VP RS12240 WP 005479695.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLD ^{TL} SL ^{EE} VEDLTGLVVEPYQT ^{TK} SAFLYALAKHYPELGLPVP ^{PP} PSG	0
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consensus	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLD ^{TL} SL ^{EE} VEDLTGLVVEPYQT ^{TK} SAFLYALAKHYPELGLPVP ^{PP} PSG	0

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NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELL ^{LL} QKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGL ^{EF} LESTEGIVPDPSAALLL	240
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NTDB id 1168 A1552VC RS11070 WP 000957200.1	EGQKDLKLGELL ^{LL} QKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGL ^{EF} LESTEGIVPDPSAALLL	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	EGQKDLKLGELL ^{LL} QKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGL ^{EF} LESTEGIVPDPSAALLL	0
NTDB id 1293 VP RS12240 WP 005479695.1	EGQKDLKLGELL ^{LL} QKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGL ^{EF} LESTEGIVPDPSAALLL	0
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NTDB id 1028 TT RS08215 WP 011173991.1	LRSDALRYGAVPIGFQNGEVEV ^{VL} SDPRHKEAVAQLLNRPARFYALPQAW ^{EEL} FRRAYPQKNRLGEVLVQEGKLSREAL	320
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
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consensus
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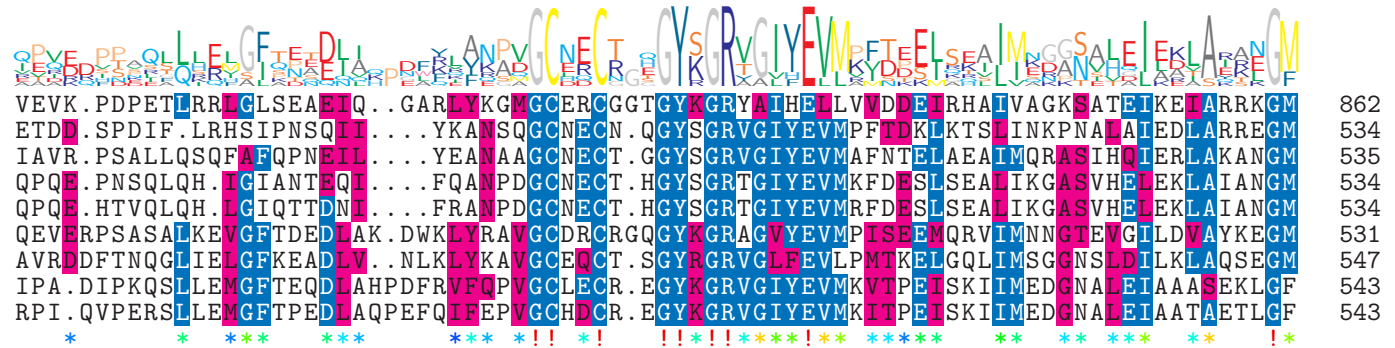
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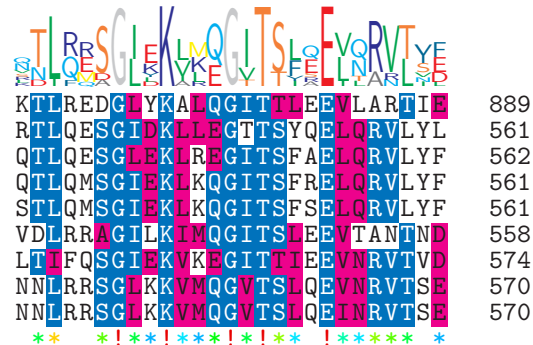
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
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consensus
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 non conserved
 similar
 $\geq 50\%$ conserved