

logo

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NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1168	A1552VC RS11070 WP 000957200.1	0
NTDB id 1293	VP RS12240 WP 005479695.1	0
NTDB id 1401	DSB67 RS12665 WP 010643259.1	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
NTDB id 79060	PotMrB4 RS05680 WP 172432750.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

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		ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	
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NTDB id 1293	VP RS12240 WP 005479695.1	0
NTDB id 1401	DSB67 RS12665 WP 010643259.1	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
NTDB id 79060	PotMrB4 RS05680 WP 172432750.1	0
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NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

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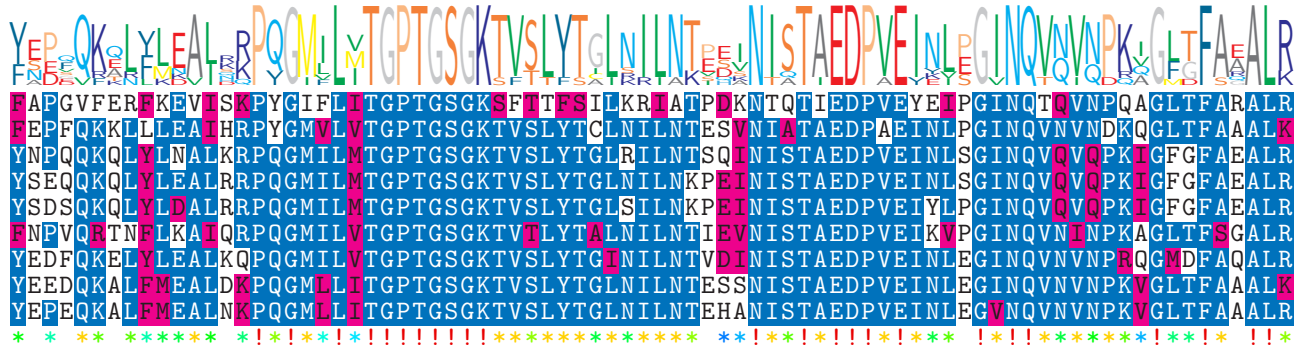
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NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
NTDB id 79060	PotMrB4 RS05680 WP 172432750.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

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		LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRNLGEVLVQEGKLSREAL	
NTDB id 1028	TT RS08215 WP 011173991.1	LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRNLGEVLVQEGKLSREAL	320
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NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

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NTDB id 1028 TT RS08215 WP 011173991.1
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 NTDB id 79060 PotMrB4 RS05680 WP 172432750.1
 NTDB id 1058 ABD1 RS01615 WP 001274986.1
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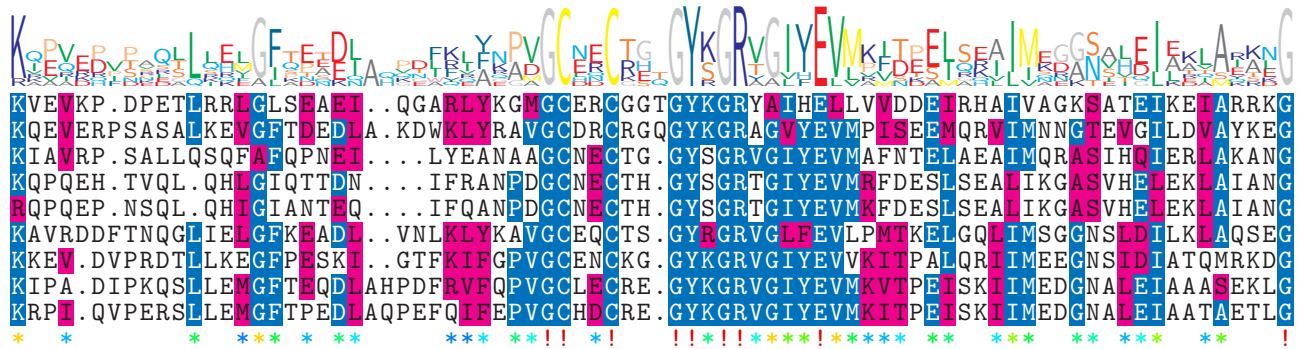
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1



889
558
562
561
561
574
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570

 non conserved
 similar
 $\geq 50\%$ conserved