

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 1293 VP RS12240 WP 005479695.1	.....	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....	0
NTDB id 1110 NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1	.....	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	.....	0
NTDB id 43001 BDGL 003240 YP 004997508.1	.....	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....	0
consensus	.....	0

logo

NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTL SLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTL SLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
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NTDB id 1251 GCO85 RS07725 WP 011946523.1	.....	0
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consensus	.....	0

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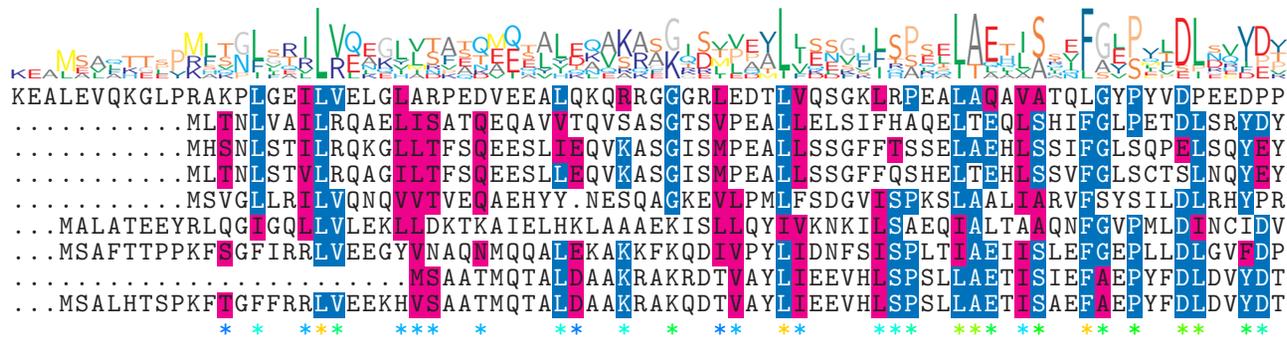
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NTDB id 43001 BDGL 003240 YP 004997508.1	.....	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....	0
consensus	.....	0

logo

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NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....	0
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NTDB id 1251 GCO85 RS07725 WP 011946523.1	.....	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	.....	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....	0
consensus	.....	0

logo

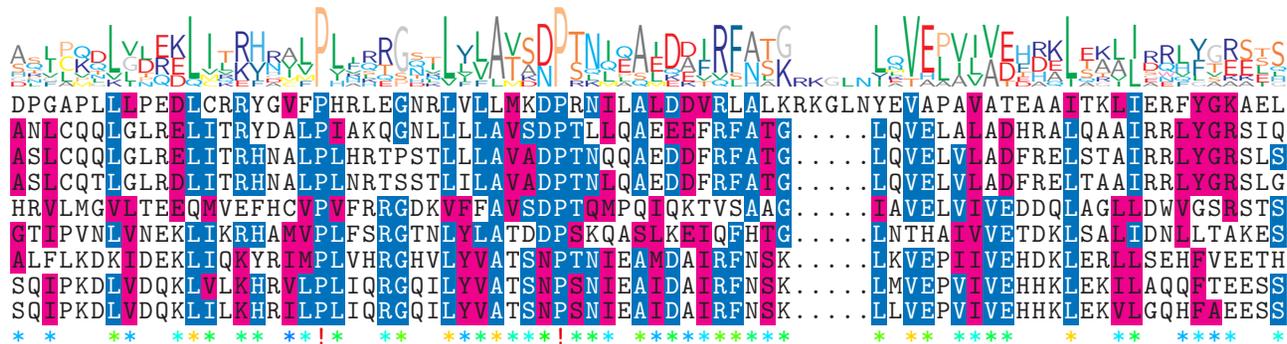
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 consensus



400  
69  
69  
69  
68  
77  
77  
55  
77

logo

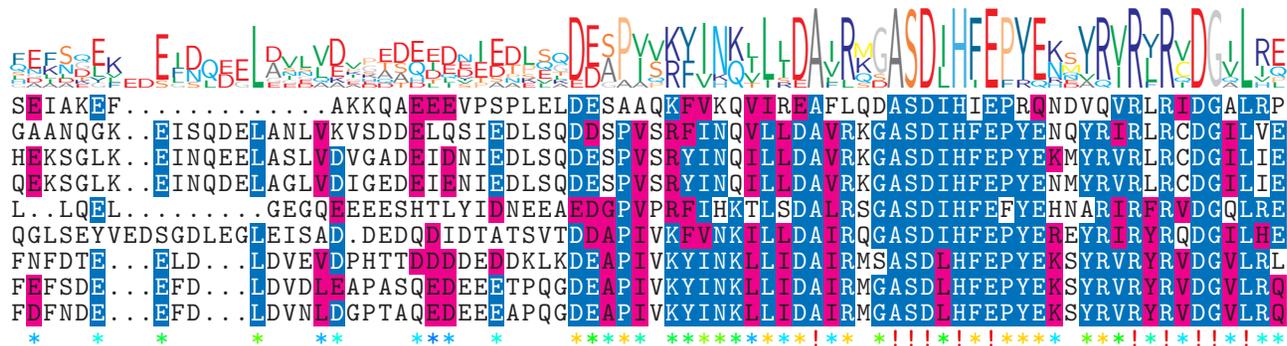
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 NTDB id 43001 BDGL 003240 YP 004997508.1  
 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 consensus



480  
144  
144  
144  
143  
152  
152  
130  
152

logo

NTDB id 1028 TT RS08215 WP 011173991.1  
 NTDB id 1168 A1552VC RS11070 WP 000957200.1  
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 NTDB id 43001 BDGL 003240 YP 004997508.1  
 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 consensus



547  
222  
222  
222  
212  
231  
226  
204  
226

logo

NTDB id 1028 TT RS08215 WP 011173991.1  
 NTDB id 1168 A1552VC RS11070 WP 000957200.1  
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 consensus



625  
301  
301  
301  
292  
310  
305  
283  
305



 non conserved  
 similar  
  $\geq 50\%$  conserved