

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTITCDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1110 NGFG RS09235 WP 003689817.1	0
NTDB id 1168 A1552VC RS11070 WP 000957200.1	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1	0
NTDB id 103063 AACH48 RS19380 WP 201900045.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus		

logo

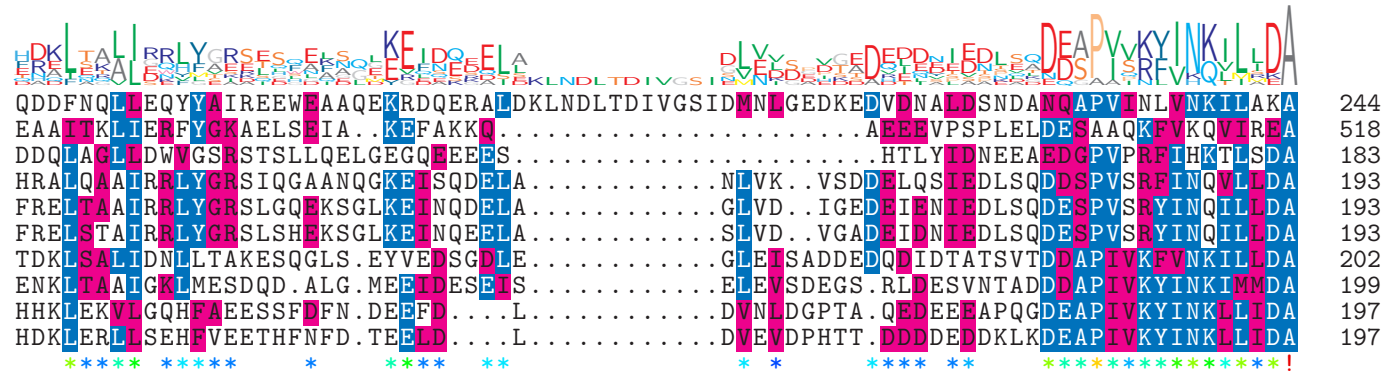
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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPVPPPSG	160
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLQKGIWISREALEEALVEQEKTGDLLGRILVVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
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NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1	0
NTDB id 103063 AACH48 RS19380 WP 201900045.1	0
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus		

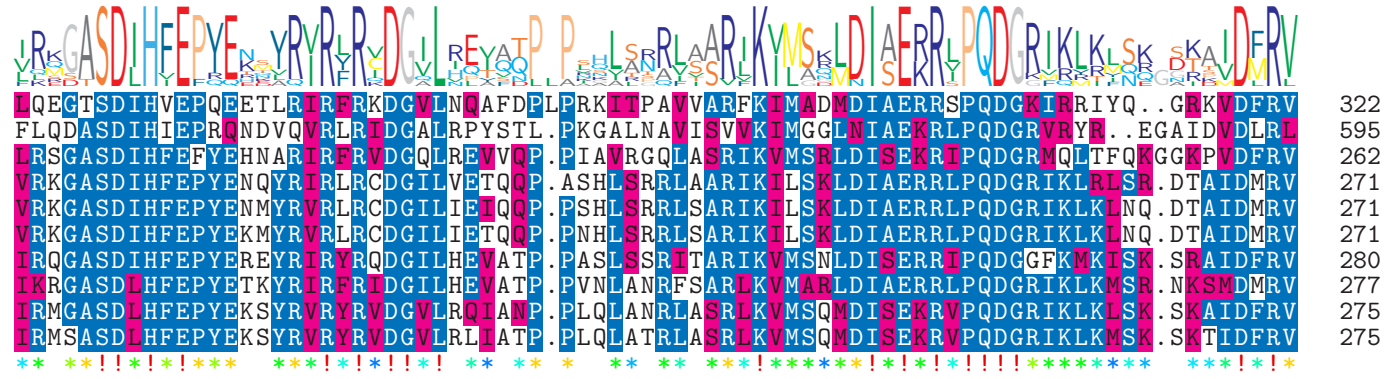
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 consensus



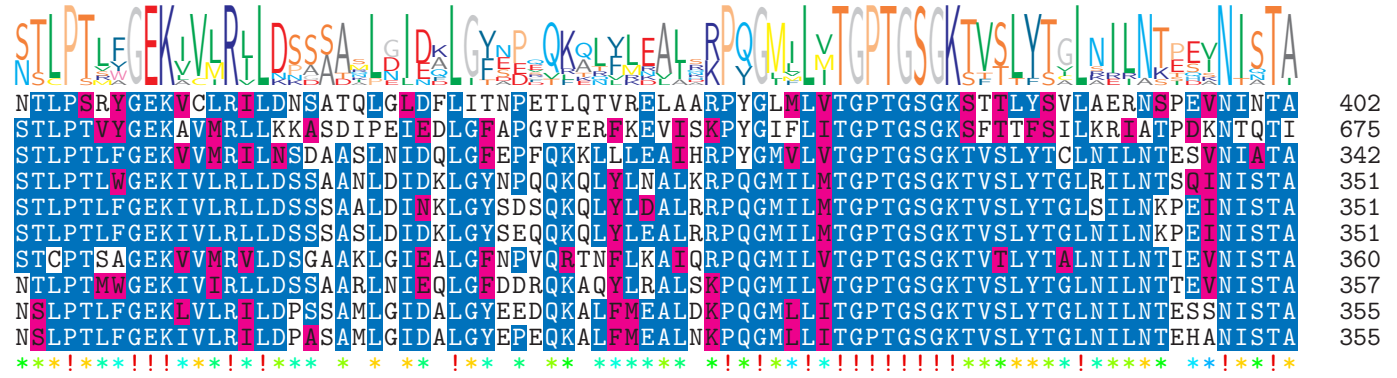
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 consensus



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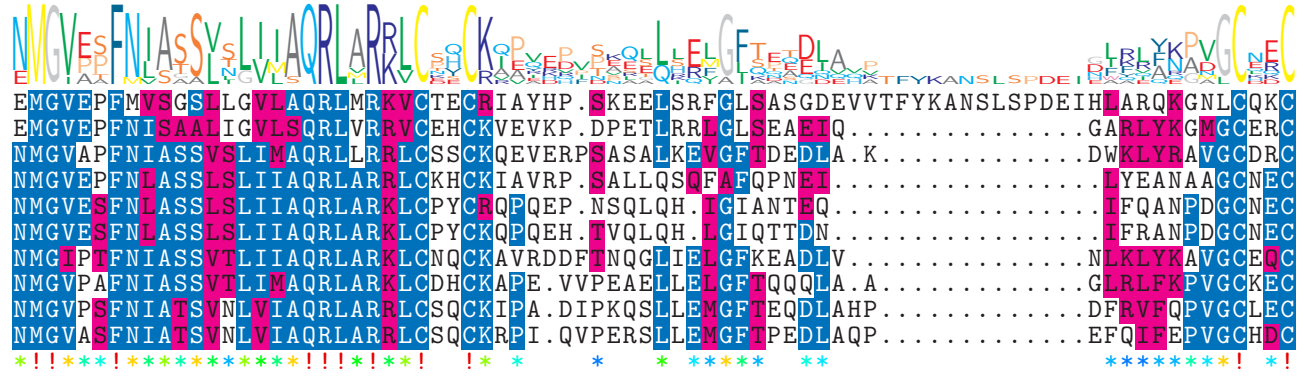
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



482
755
422
431
431
431
440
437
435
435

logo

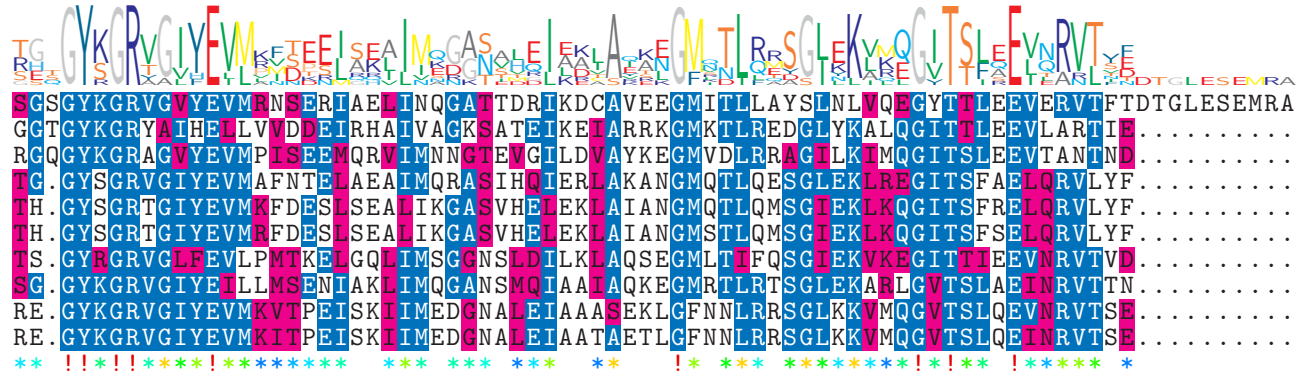
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 consensus



561
819
488
493
492
492
505
502
501
501

logo




NTDB id 1410 SGL RS13950 WP 010873519.1
 NTDB id 1028 TT RS08215 WP 011173991.1
 NTDB id 1110 NGFG RS09235 WP 003689817.1
 NTDB id 1168 A1552VC RS11070 WP 000957200.1
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



641
889
558
562
561
561
574
571
570
570

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	KRKSALTCRTCAAELQQEWLDCPYCMTSRFS	672
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NTDB id 103063 AACH48 RS19380 WP 201900045.1	571
NTDB id 1058 ABD1 RS01615 WP 001274986.1	570
NTDB id 1018 ACIAD RS01685 WP 004920473.1	570
consensus		

 non conserved
 similar
 ≥ 50% conserved