

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTITGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTITGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
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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	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
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NTDB id 1110 NGFG RS09235 WP 003689817.1		0
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consensus		

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consensus		

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NTDB id 1355 HI 0298 AAC21962.1	MTSY..ALLHTQRTAK.....NGEFT..	21
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NTDB id 1251 GC085 RS07725 WP 011946523.1	..MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVKN..KILSAEQIALT.....AAQNFQVP	68
NTDB id 1058 ABD1 RS01615 WP 001274986.1	..MSALHTSPKFTIGFFRRLLVEEKHVSAAATMQTALDAAKRAKQDVTAYLIEE..VHLSPSLLAET.....ISAFAEP	68
NTDB id 1018 ACIAD RS01685 WP 004920473.1	..MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDITVPLYIDN..FSISPLTIAEI.....ISLEFGEF	68
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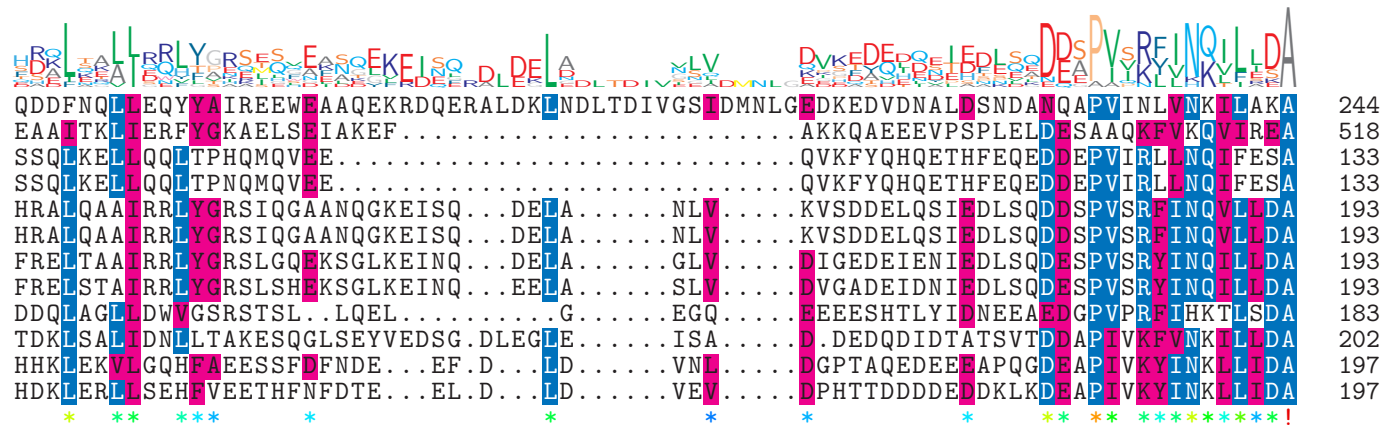
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NTDB id 1168 A1552VC RS11070 WP 000957200.1	ETDLSRYDYAN....LCQQLGLRELITRYDALPIAKQG...NLLLLAVSDPTLLQAEEDFRFATG.....LQVELALAD	127
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	LLDLGVFDPAL....FLKDKIDEKLIQKYRIMPLVHRG...HVLVYVATSNPTNIEAMDIAIRFNSK.....LKVEPIIVE	135
consensus	** * * * * * ! * * * * * * * * * * *	

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NTDB id 1410 SGL RS13950 WP 010873519.1
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consensus



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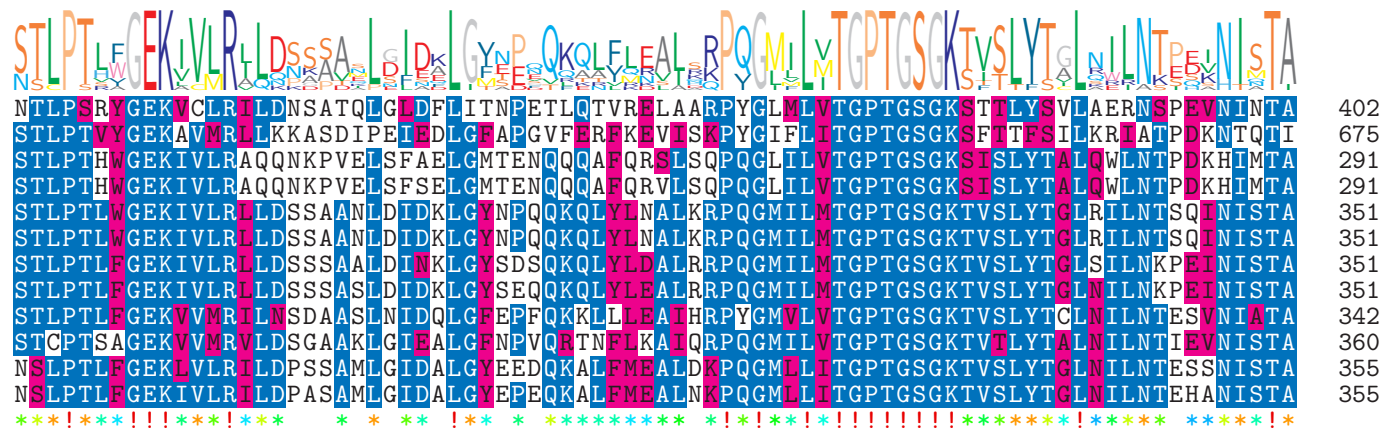
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consensus



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consensus



482
755
371
371
431
431
431
431
422
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435
435

logo

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consensus

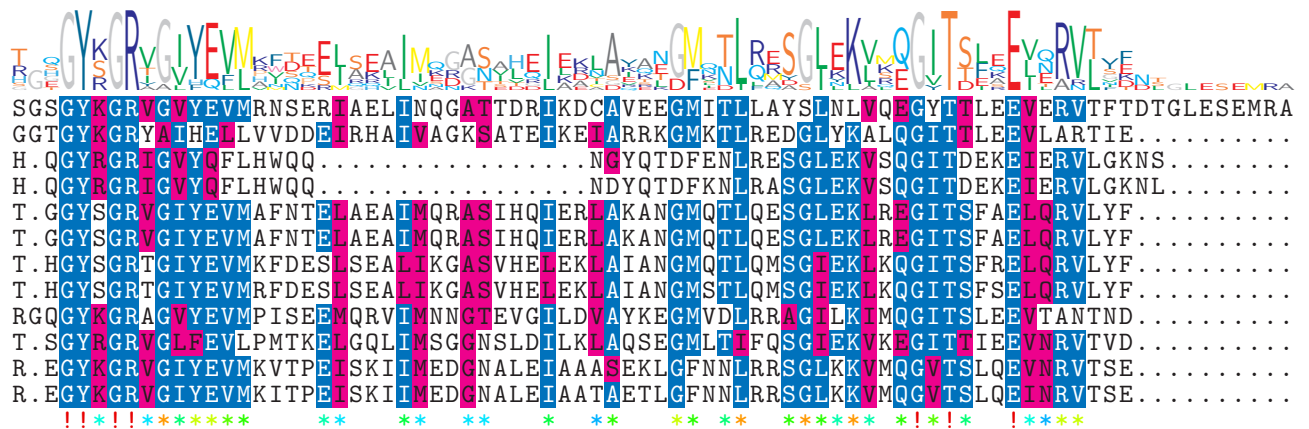


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819
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493
493
492
492
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logo

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consensus



641
889
464
464
562
562
561
561
558
574
570
570

logo

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved