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NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
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NTDB id 1293 VP RS12240 WP 005479695.1		0
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NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
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consensus		

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consensus		

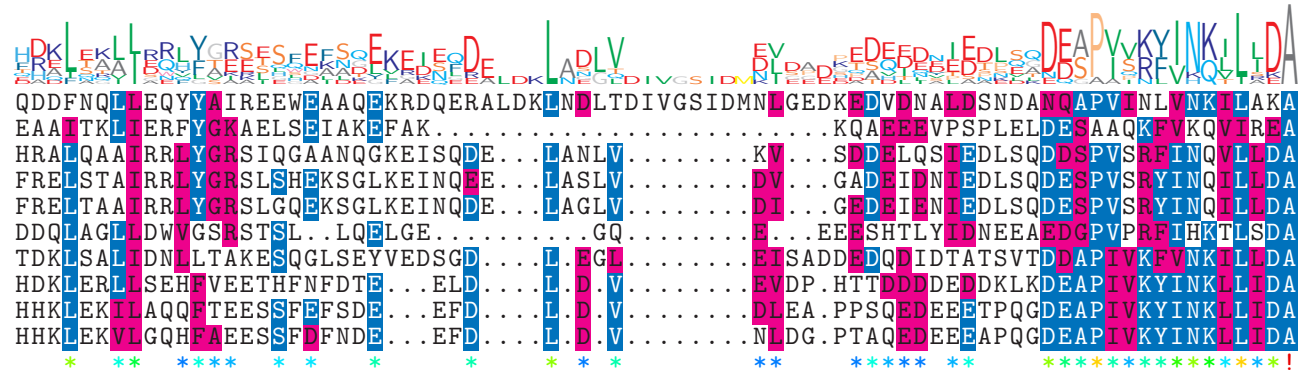
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NTDB id 1251	GCO85 RS07725 WP 011946523.1	...MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLA AA EKI SLLQYIVK..NKILSAEQIALT.....AAQNFQVP	68
NTDB id 1018	ACIAD RS01685 WP 004920473.1	...MSAFTTPPKFSGFIRRLVEEGYVNANMQQALEKAKKFQDIVPYLID..NFSISPPLTAEI.....ISLEFGEP	68
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consensus		* * * * *	

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NTDB id 1293	VP RS12240 WP 005479695.1	QPELSQQEYAS.....LCQQLGLRELITRNHALPLHRT...PSTLLILAVADPTNQQAEDDFRFATG.....LQVELVLAD	127
NTDB id 1401	DSB67 RS12665 WP 010643259.1	CTSLNQHYEYAS.....LCQTGLGRLDLITRHNALPLNRT...SSTILLAVADPTNLQAEDDFRFATG.....LQVELVLAD	127
NTDB id 1110	NGFG RS09235 WP 003689817.1	ILDLRHYPRHR.....VLMGVLTTEEQMVEFHCVPVFRR...GDKVFFAVSDPTQMPQIQKTVSAG.....I AVELVIVE	126
NTDB id 1251	GCO85 RS07725 WP 011946523.1	MLDINCIDVGT.....IPVNLVNEKLIKRHAMVPLFSR...GTNLYLATDDPSKQASLK EIQFHTG.....LNTHA IVE	135
NTDB id 1018	ACIAD RS01685 WP 004920473.1	LLDLGVEDPAL.....FLKDKEDEKLIQKYRIMPLVHR...GHVLYVATSNPNTIEAMD AIRFN SK.....LKVEPIIVE	135
NTDB id 377454	FR838 RS12145 WP 032042313.1	YFDLDVYDTSQ.....TPKDLVDQKLV LKH RVLPLIQR...GGILYVATSNSPN IEAIDAIRFN SK.....LMVEPVIVE	135
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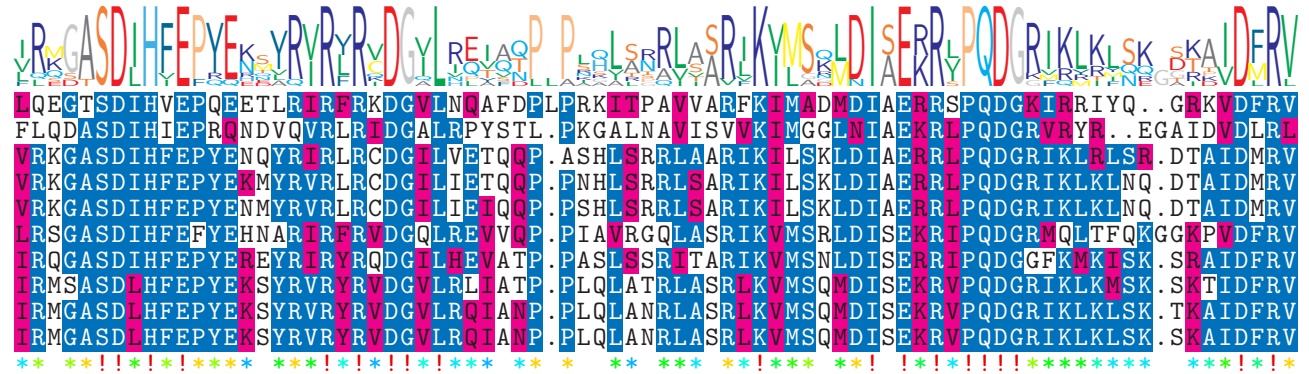
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consensus

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consensus
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consensus

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logo

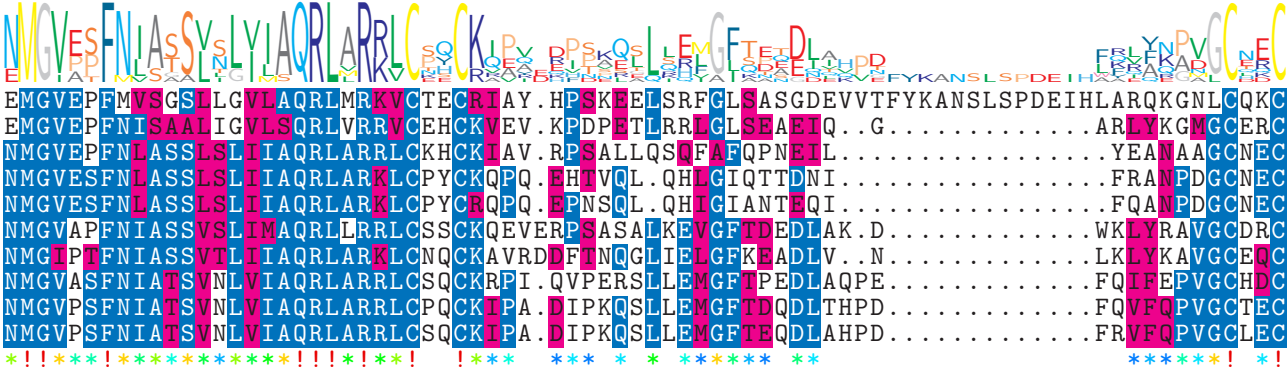
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consensus



482
755
431
431
431
422
440
435
435
435

logo

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consensus



561
819
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492
492
488
505
501
501
501

logo

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NTDB id 1058 ABD1 RS01615 WP 001274986.1
consensus



641
889
562
561
561
558
574
570
570
570

logo

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved