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NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 76736 H7R67 RS19830 WP 041211787.1		0
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

logo

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	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLL	0
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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consensus

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[illegible]

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consensus

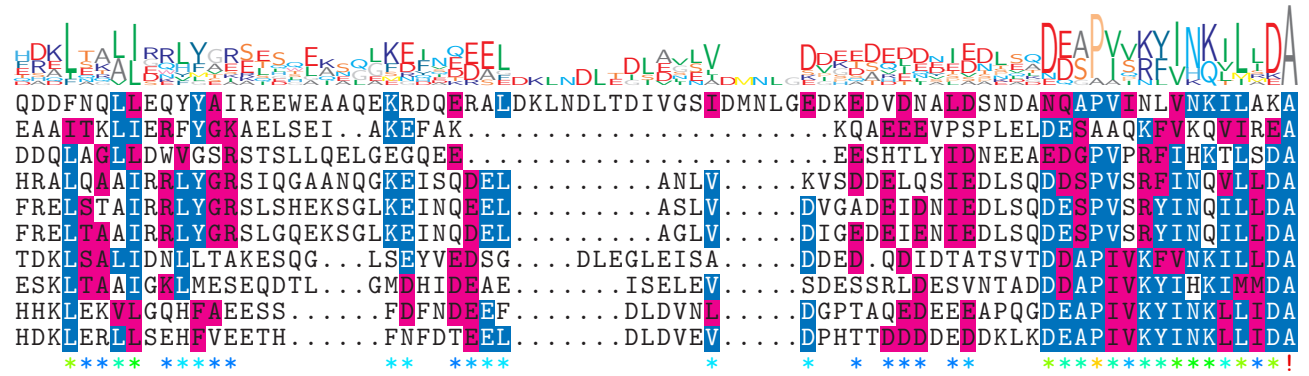
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ILDLRHYPRH....RYLMGVI TEFGVEFHCVPFVRRG...DKYFFAVSDPTQMPQIQKTVSAAG.....I AVLIVE 126
ETDL SRDYDA....NLCCQLGLRELITRYDALPIAKQG...NL LLA VSDPTLLQAEEFRFATG.....LQVELALAD 127
QPESQYEYA....SLCQQLGLRELITRHNAFLHRTP...STLLAVADPTNQQAEDDFRFATG.....LQVELVLAD 127
MTSLNQYEYA....SLCQTGLRLDITRHNAFLPLNRS...STLILAVADPTNQAEDDFRFATG.....LQVELVLAD 127
CLDINCIDVC....TIPVNLVNEKLKRHAMVPLFSRG...TNLYLATDDPSKQASKEIQFHTG.....LNTHAIVVE 135
LLDLAAFOLA....ETPQKYLNQKLEKHVVLPITYTQG...HTLYIAMSDPTNVSALEDGFGSFG.....LHTEALLVE 131
YFDLDVYDTS....QIPKDLVDQKLILKHLRIPLIQRG...QILYVATSNPSNIEADAIRFNASK.....LLVEPVIVE 135
LLDLGVFDPA....LFLKDKIDEKLIQKYIMPLVHRG...HVLYVATSNPTNTEAMDAIRFNASK.....LKVEPIIVE 135

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consensus

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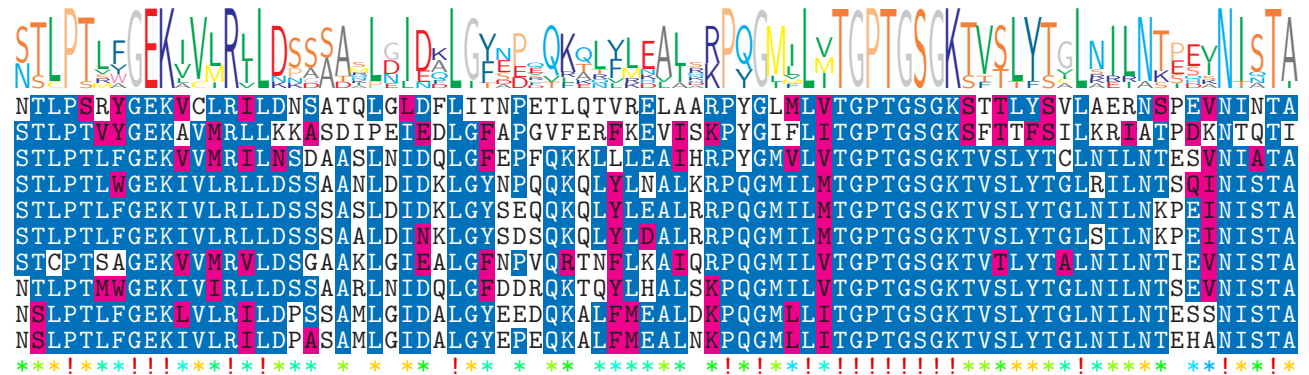
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consensus

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logo

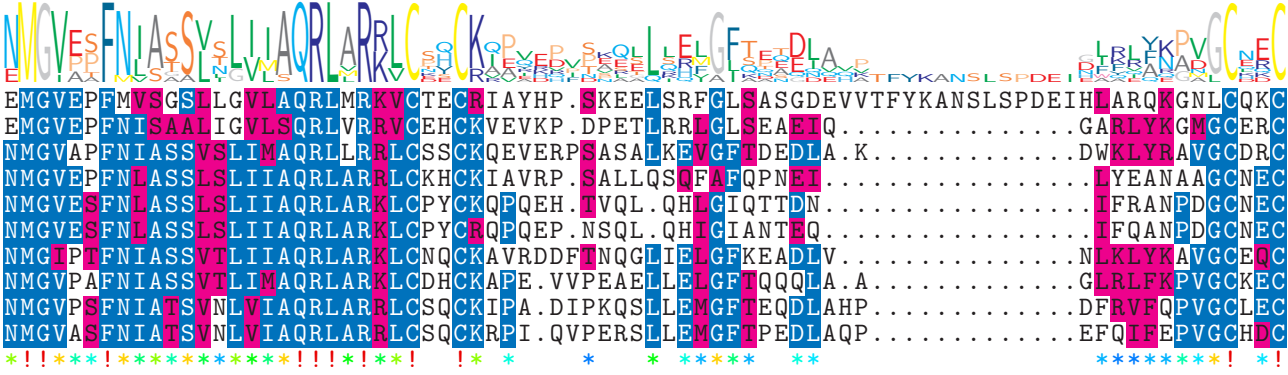
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consensus



482
755
422
431
431
440
434
435
435

logo

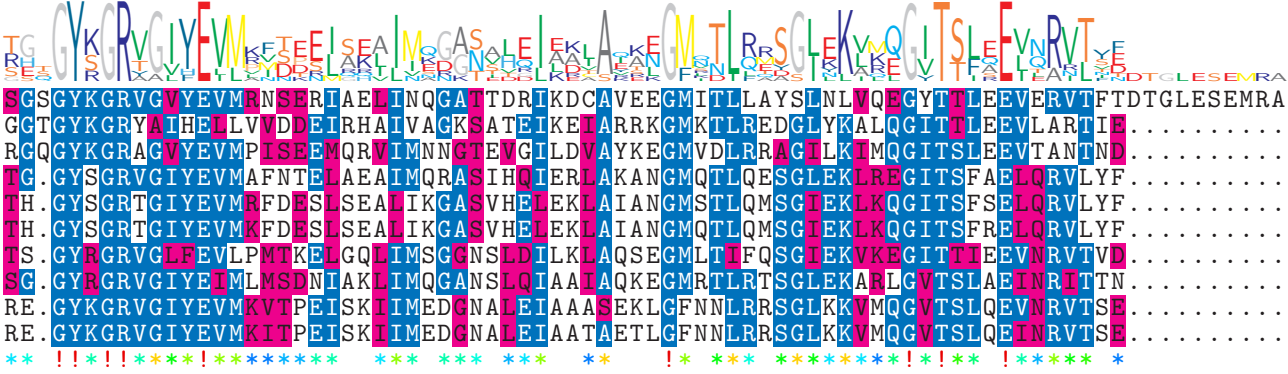
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consensus



561
819
488
493
492
492
505
499
501
501

logo

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consensus



641
889
558
562
561
561
574
568
570
570

logo

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved