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NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
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NTDB id 361071 EA667 RS17300 WP 001274988.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
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	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
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NTDB id 1110 NGFG RS09235 WP 003689817.1		0
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NTDB id 361071 EA667 RS17300 WP 001274988.1		0
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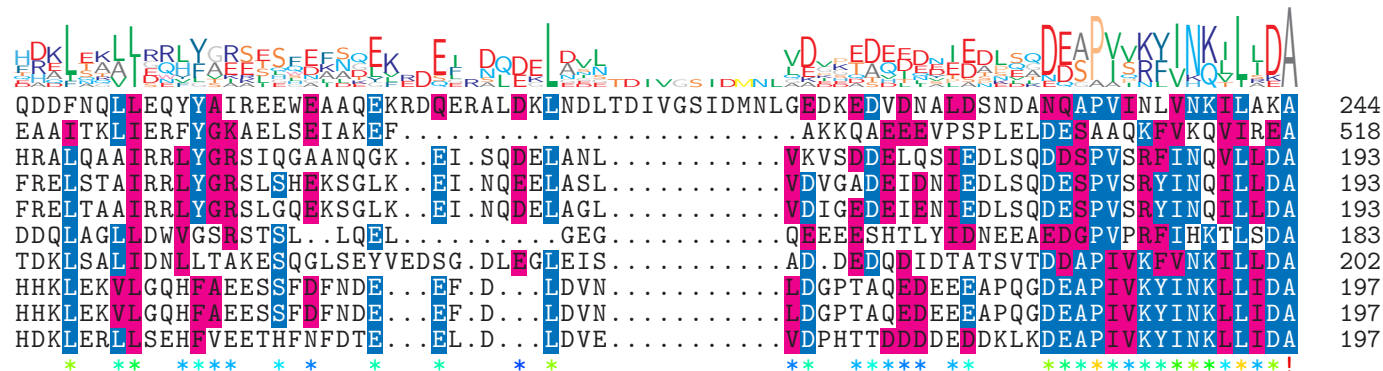
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consensus		

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NTDB id 1168	A1552VC RS11070 WP 000957200.1	MLTNLVALLRQAELISATQEQAQVVTQVSASGTSVPPEALLE..LSIFHAQELTEQ.....LSHTFGIP	60
NTDB id 1293	VP RS12240 WP 005479695.1	MHSNLSTILRRQKGLTFSSQESLIEQVKASGISMPALLS..SGFFTSSELAEH.....LSSIFGLS	60
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NTDB id 1251	GCO85 RS07725 WP 011946523.1	...MALATEEYRLQGTGLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVK..NKILSAEQIALT.....AAQNFQVP	68
NTDB id 361071	EAE67 RS17300 WP 001274988.1	...MSALHTSPKFTGFFRRLVEEKHVSAATMQTALDAAKRAKQDTVAYLIE..EVHLSPSLLAET.....ISAEFAEP	68
NTDB id 1058	ABD1 RS01615 WP 001274986.1	...MSALHTSPKFTGFFRRLVEEKHVSAATMQTALDAAKRAKQDTVAYLIE..EVHLSPSLLAET.....ISAEFAEP	68
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consensus		* * * * *	

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consensus



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consensus



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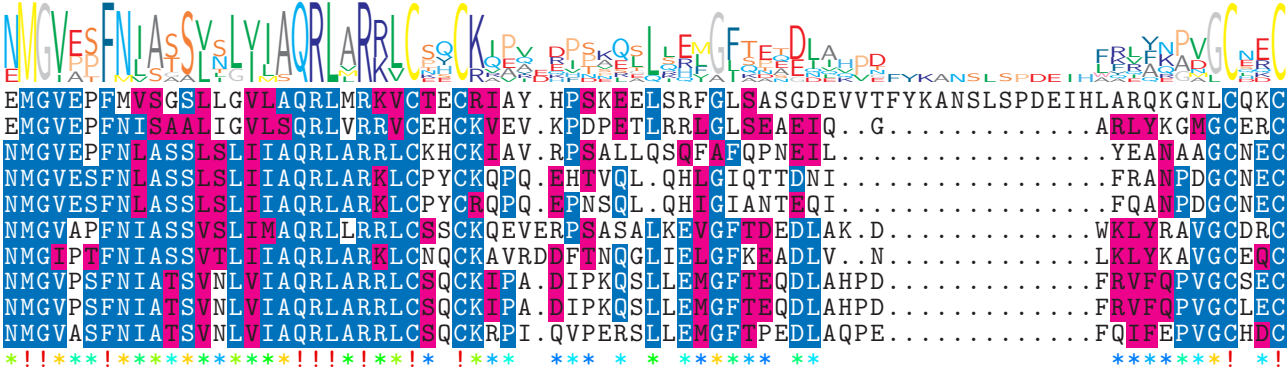
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consensus		



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logo

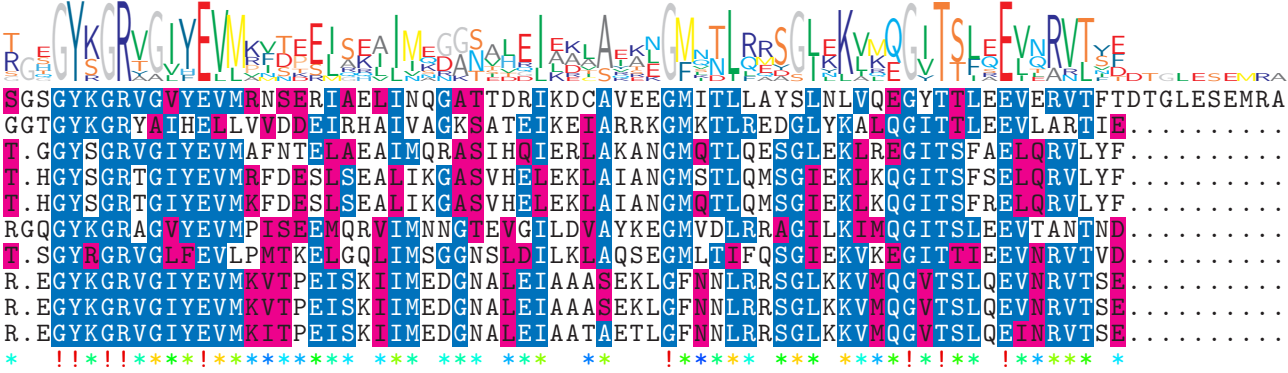
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consensus		



561
819
493
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492
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consensus		



641
889
562
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved