

NTDB id	Accession	Sequence	Score
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NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1401	DSB67 RS12665 WP 010643259.1		0
NTDB id 1110	NGFG RS09235 WP 003689817.1		0
NTDB id 1251	GC085 RS07725 WP 011946523.1		0
NTDB id 1014839	ABWW75 RS17235 WP 001274990.1		0
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consensus			

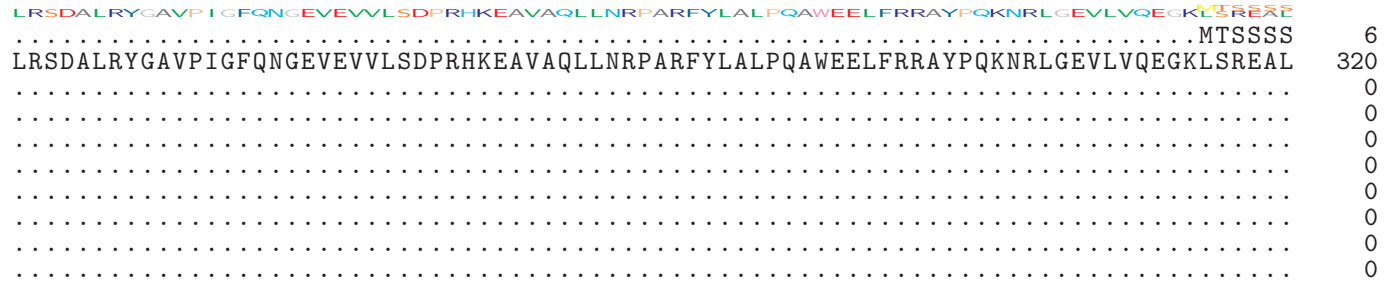
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consensus	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLD	

NTDB id	Accession	Sequence	Score
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NTDB id 1110	NGFG RS09235 WP 003689817.1		0
NTDB id 1251	GC085 RS07725 WP 011946523.1		0
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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 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

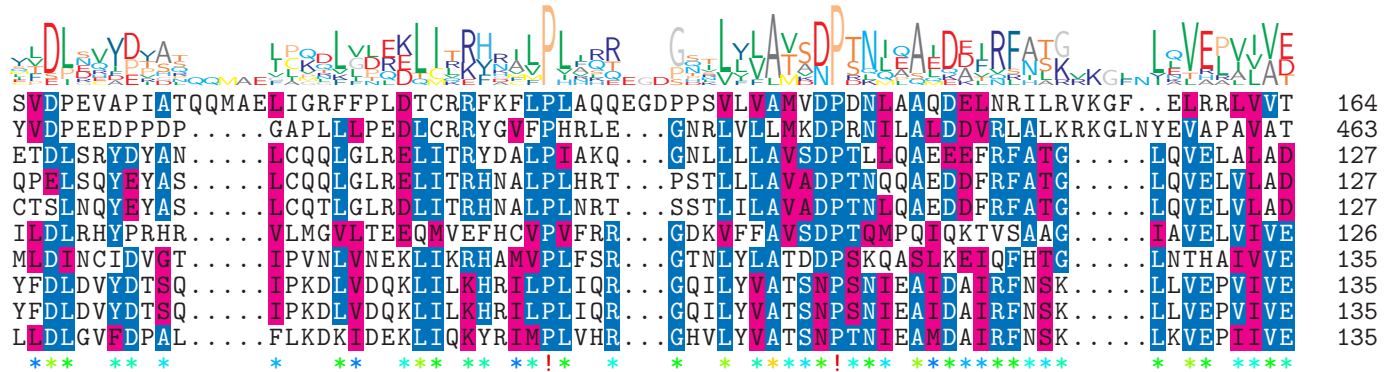
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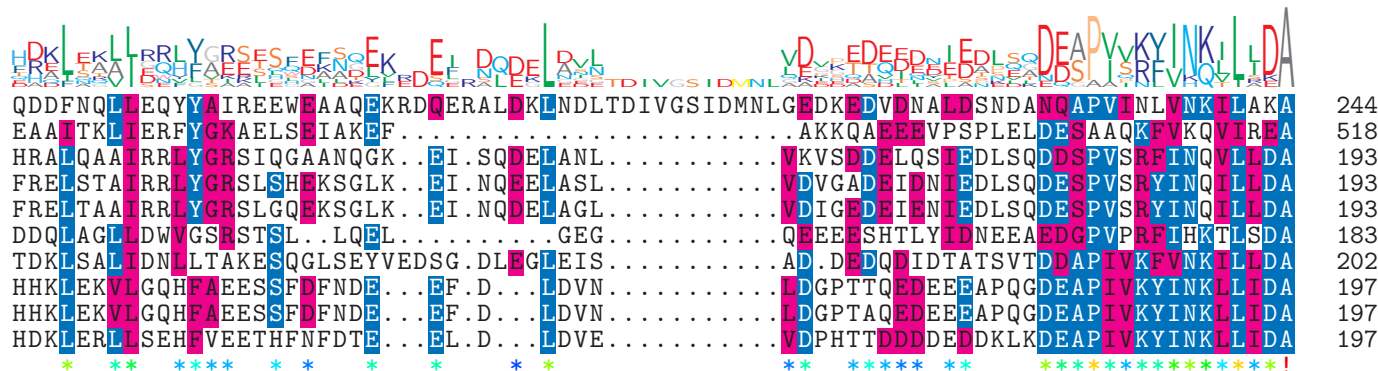
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consensus

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logo

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NTDB id 1168 A1552VC RS11070 WP 000957200.1
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consensus



logo

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consensus



logo

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consensus



logo

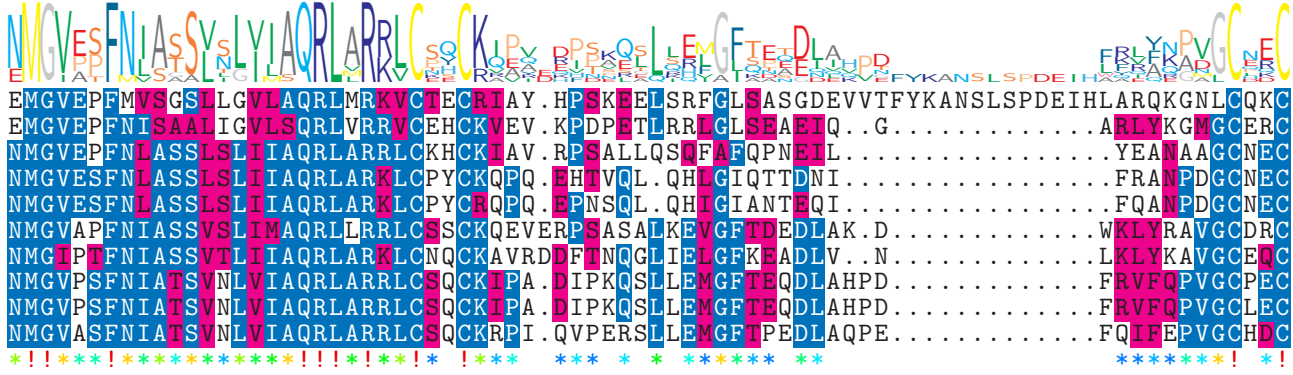
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NTDB id 1014839 ABWW75 RS17235 WP 001274990.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1
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consensus



482
755
431
431
431
422
440
435
435
435

logo

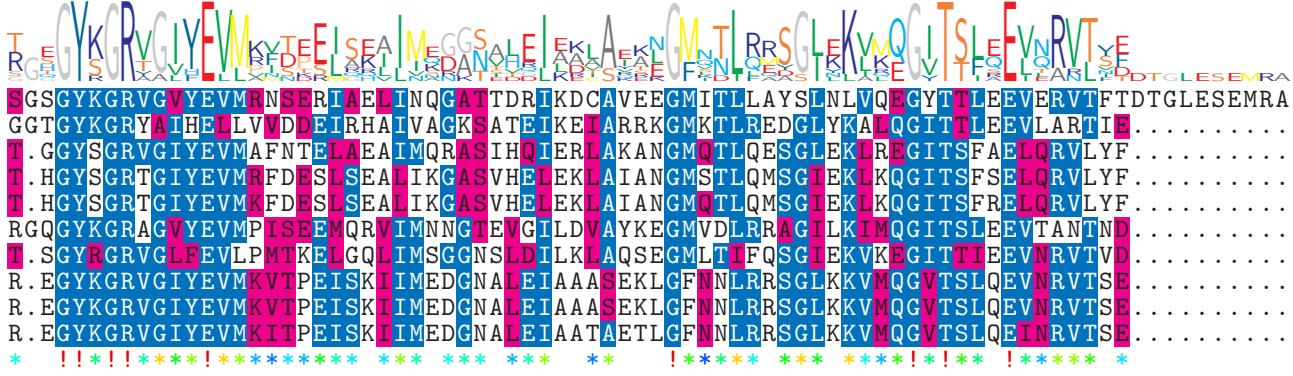
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus



561
819
493
492
492
488
505
501
501
501

logo

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consensus



641
889
562
561
561
558
574
570
570
570

logo

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NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved