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NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1014011 ABWW73 RS17235 WP 001274990.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
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consensus		

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consensus			0

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NTDB id 1014011 ABWW73 RS17235 WP 001274990.1		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
NTDB id 1028 TT RS08215 WP 011173991.1
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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[illegible]

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consensus

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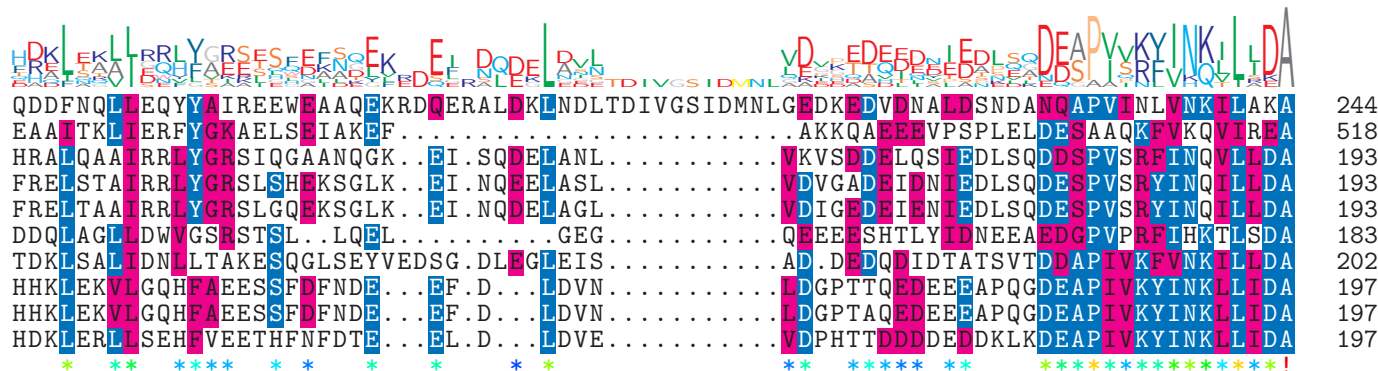
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
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Sequence	Position
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ETDLSRYDYAN.....LCQQGLGLRELITRYDALPIAKQ...GNLLLVASDPTLLQAEERFRFATG.....LQVELALAD	127
QPELSQYFYAS.....LCQQGLGLRELITRHNALPLHRT...PSTLLLVADPTNQQAEDDFRFATG.....LQVELVLAD	127
CTSLNQYFYAS.....LCQTLGLRLITRHNALPLNRT...SSTLLLVADPTNLQABDDFRFATG.....LQVELVLAD	127
ILDRLHYPRHR.....VLMGVLTEEQMVFEHCVPLVFRR...GDKVFFAVSDPTQMPQIQKTVSAAG.....LAVELVIVE	126
MLDINCIIVGT.....IPVNLVNEKLIKRRHAMVPLFSR...GTNLVLATDDPSKQASLKEIQFHTG.....LNTHAIVVE	135
YFDLDVYDTSQ.....IPKDLVDQKLIKLRHILPLIQR...GQILYVATSNPSNIEATDAIFRNSK.....LLVEPVIVE	135
YFDLDVYDTSQ.....IPKDLVDQKLIKLRHILPLIQR...GQILYVATSNPSNIEATDAIFRNSK.....LLVEPVIVE	135
LLDLGVFDPAL.....FLDKDIDEXLIKRYRIMPLVHR...GHVLYVATSNPTNIEATDAIFRNSK.....LKVEPIVE	135

logo

NTDB id 1410 SGL RS13950 WP 010873519.1
NTDB id 1028 TT RS08215 WP 011173991.1
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consensus



logo

NTDB id 1410 SGL RS13950 WP 010873519.1
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consensus



logo

NTDB id 1410 SGL RS13950 WP 010873519.1
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consensus



logo

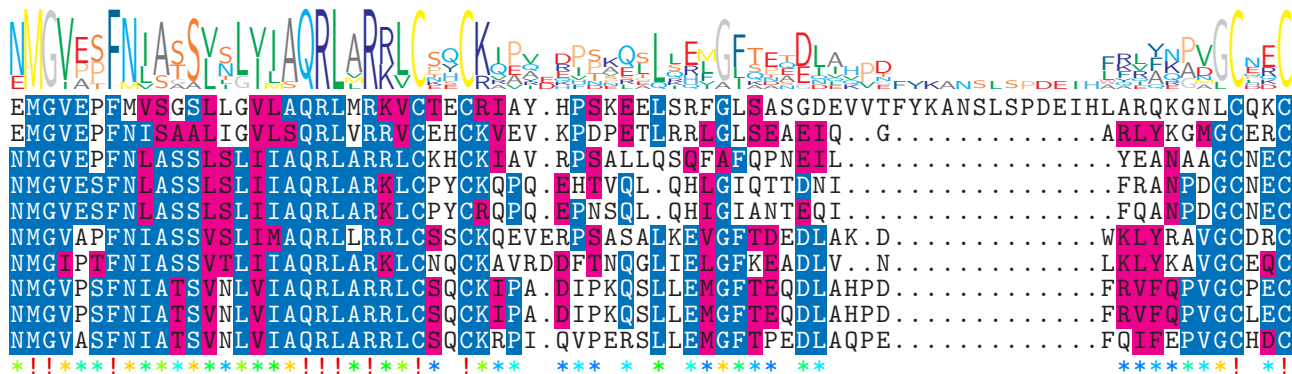
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consensus



482
755
431
431
431
422
440
435
435
435

logo

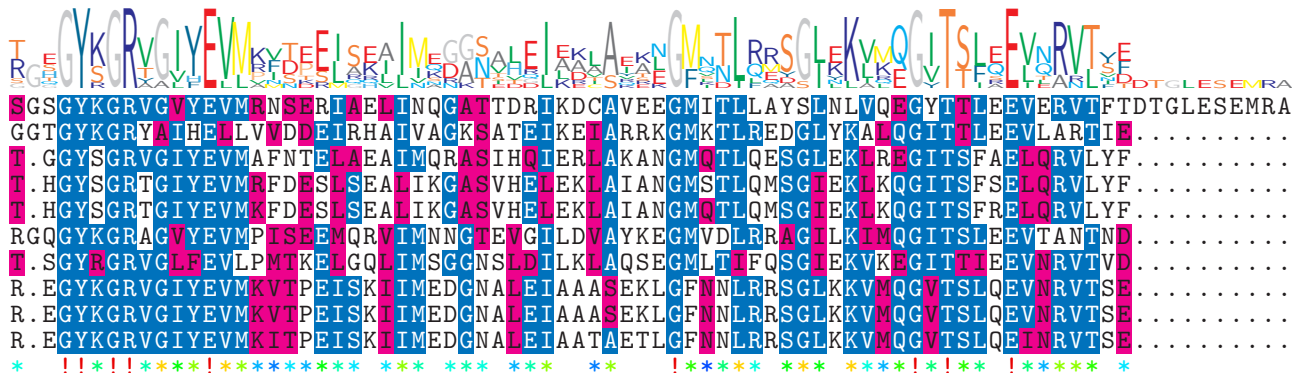
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consensus



561
819
493
492
492
488
505
501
501
501

logo

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NTDB id 1014011 ABWW73 RS17235 WP 001274990.1
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consensus



641
889
562
561
561
558
574
570
570
570

logo

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NTDB id 1014011 ABWW73 RS17235 WP 001274990.1		570
NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved