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consensus		

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consensus		

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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		0

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NTDB id 1410 SGL RS13950 WP 010873519.1
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consensus

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[illegible]

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consensus

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[illegible]

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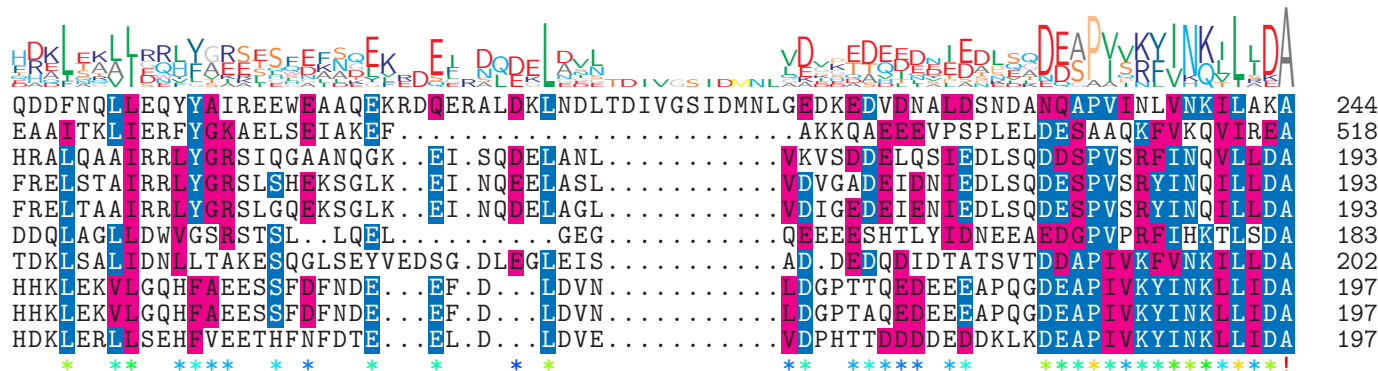
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consensus

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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.....VLMGVLTEEQMVFEHCVPLVFR...GDKVFFAVSDPTQMPQIQKTVSAAG.....LAVELVIVE	126
MLDINCIIVGT.....IPVNLVNEKLIKRRHAMVPLFSR...GTNLVLATDDPSKQASLIKELVPHGTG.....LNTHAIVIVE	135
YFDLDVYDTSQ.....IPKDLVDQKLIKLRHILPLIQR...GQILYVATSNPSNTEADATRFNSK.....LLVEPVIVE	135
YFDLDVYDTSQ.....IPKDLVDQKLIKLRHILPLIQR...GQILYVATSNPSNTEADATRFNSK.....LLVEPVIVE	135
LLDLGVFDPAL.....FLDKDIDELLIKRYRIMPLVHR...GHVLYVATSNPTNTEADATRFNSK.....LKVEPIVE	135

logo

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NTDB id 1168 A1552VC RS11070 WP 000957200.1
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consensus



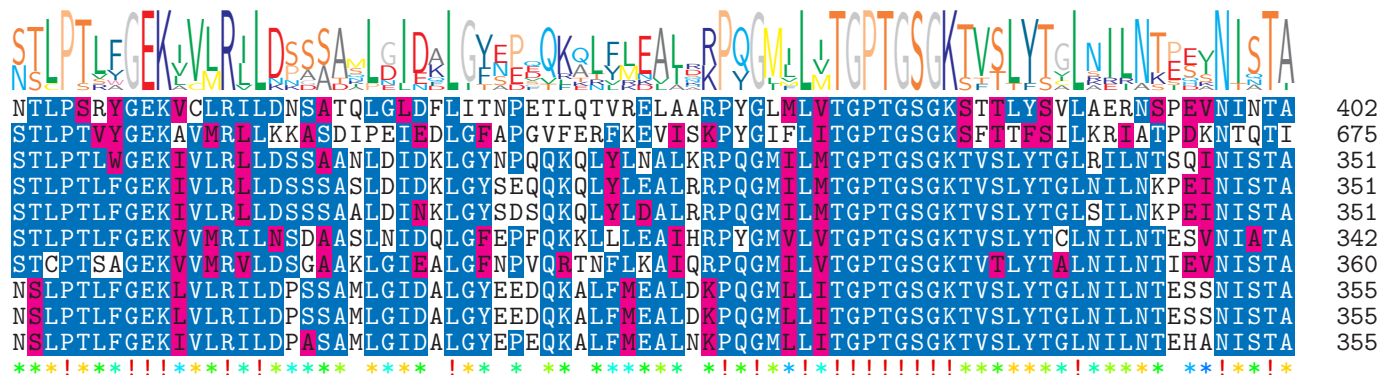
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consensus



logo

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consensus



logo

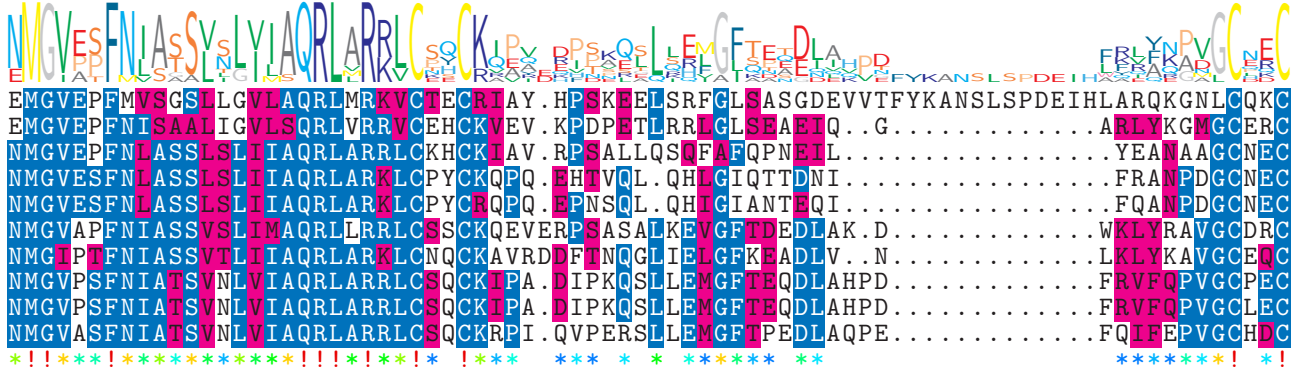
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consensus



482
755
431
431
431
422
440
435
435
435

logo

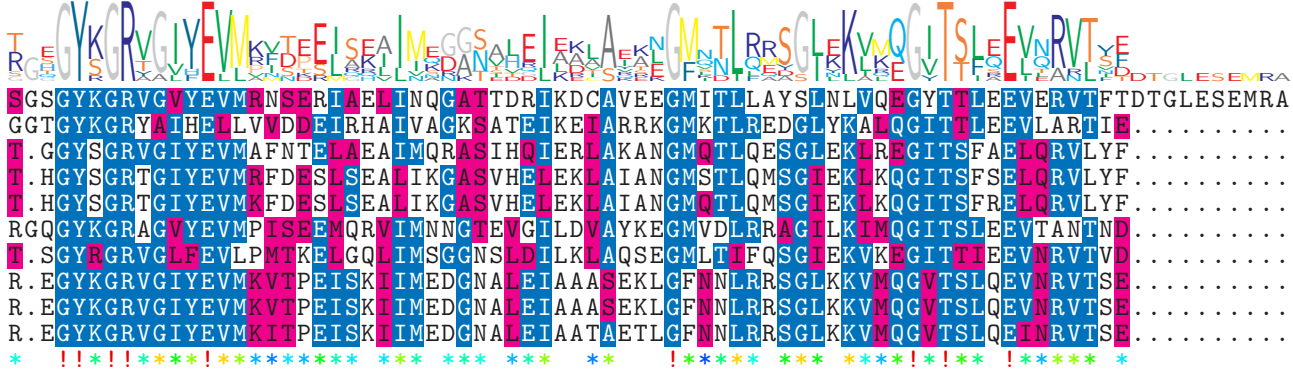
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus



561
819
493
492
492
488
505
501
501
501

logo

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NTDB id 1006090 ABI198 RS16600 WP 001274990.1
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
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