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NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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consensus		

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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
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consensus		

NTDB id 1410 SGL RS13950 WP 010873519.1
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consensus

[illegible]

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consensus

[illegible]

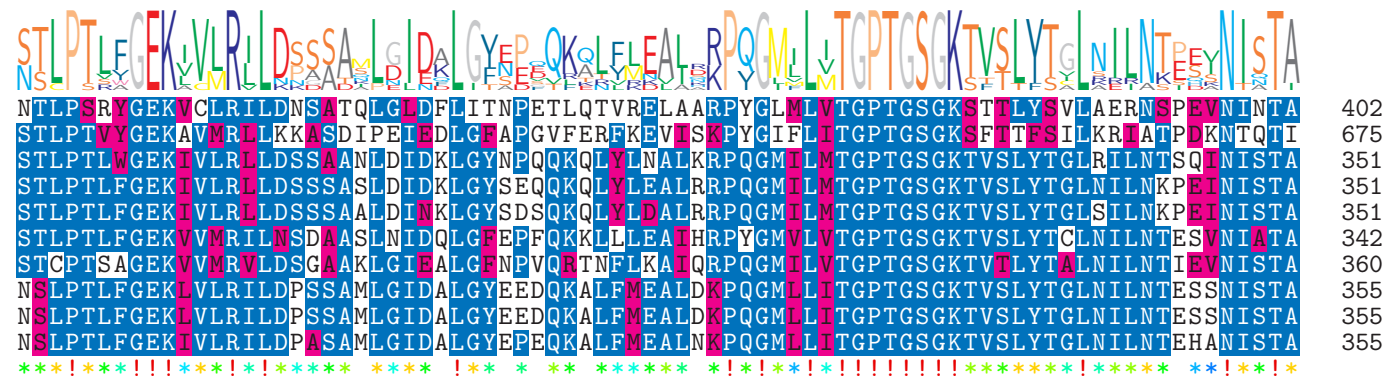
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consensus

Sequence	Position
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ETDLIRYDYAN.....CQQGLGLRELITRYDALPTAKQ.....GNLLLLAVSDPTLLQAEFFRFATG.....LQVELALAD	127
QPELSQYEYAS.....LCQQGLGLRELITRHNALPLHRT.....PSTLLAVADPTNQQAEDDFRFATG.....LQVELVLAD	127
CTSLNQYEYAS.....CQTLLGLRLITRHNALPLNRT.....SSTLLAVADPTNQQAEDDFRFATG.....LQVELVLAD	127
ITDLRHYPRHR.....VLMGVLTEEQVEFHCVPRFR.....GDKVFFAVSDPTQMPQIKQTVSAAQ.....IAVELVIVE	126
MDTINCIDVGT.....IPVNLVNEKLIKRHAMVPLFSR.....GTNLLVATDDPSKQASKEIQFHTG.....LNTHAIVVE	135
YFDLDVYDTSQ.....IPKDLVDQKLLILKHRILPLIQR.....GQILYVATSNPSNIEAIDAIRFNSK.....LLVEPVIVE	135
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consensus
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consensus



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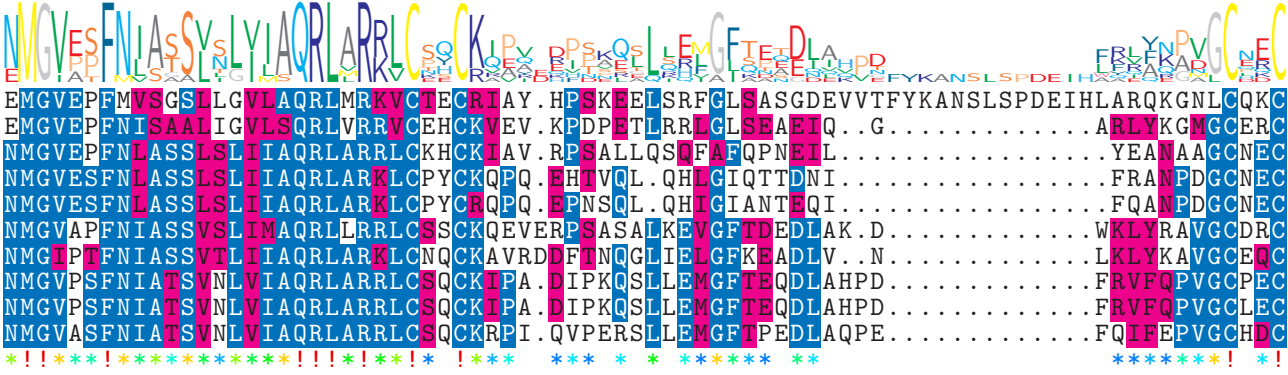
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consensus



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431
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435
435
435

logo

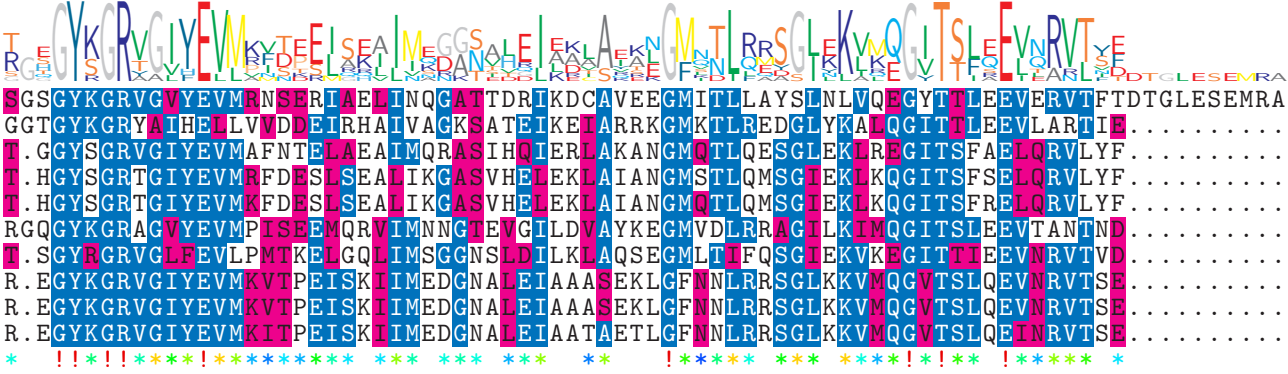
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consensus



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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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved