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NTDB id 1293	VP RS12240 WP 005479695.1		0
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NTDB id 1058	ABD1 RS01615 WP 001274986.1		0
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consensus			

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consensus		

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NTDB id 1110050 AC0CIC RS17055 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
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NTDB id 1401 DSB67 RS12665 WP 010643259.1
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NTDB id 1110050 ACQCIC RS17055 WP 001274990.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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[illegible]

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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus
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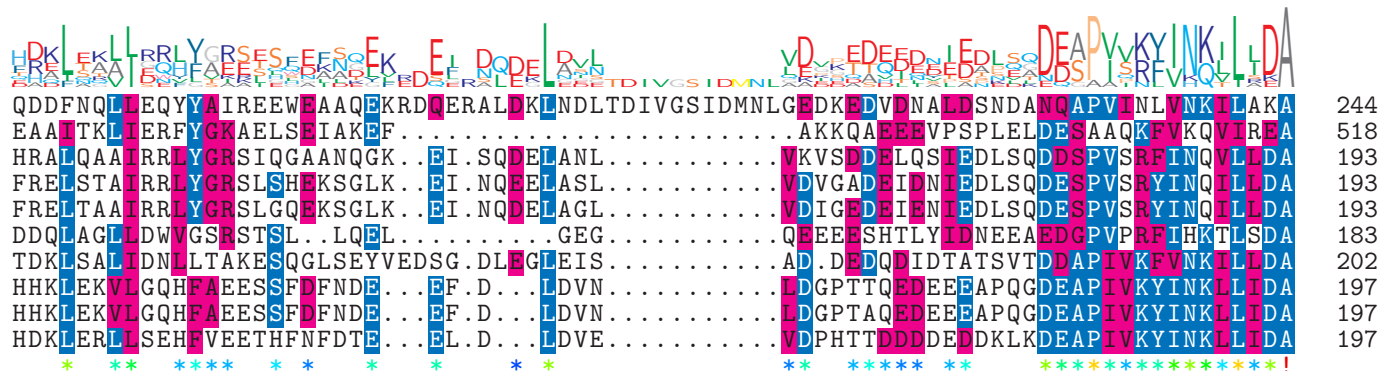
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consensus
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Sequence	Position
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YVDPEEDPPDP.....GAPLLLPEDLCRRYGVFPHRLE...GNRLVLLMKDPRNIALDDVRLALKRKGLNVEVAPAVAT	463
ETDLSRYDYAN.....LCQQGLGLRELITRYDALPIAKQ...GNLLLVASDPTLLQAEERFRFATG.....LQVELALAD	127
QPELSQYEYAS.....LCQQGLGLRELITRHNALPLHRT...PSTLLLVADPTNQQAEDDFRFATG.....LQVELVLAD	127
CTSLNQYEYAS.....LCQTLGLRLITRHNALPLNRT...SSTLLLVADPTNLQABDDFRFATG.....LQVELVLAD	127
ILDRLHYPRHR.....VLMGVLTEEQMVFEHCVPVFRR...GDKVFFAVSDPTQMPQIQKTVSAAG.....IAVELVIVE	126
MLDINCIIVGT.....IPVNLVNEKLIKRRHAMVPLFSR...GTNLVLATDDPSKQASLKEIVFHTG.....LNTHAIVVE	135
YFDLDVYDTSQ.....IPKDLVDQKLIKLRHILPLIQR...GQILYVATSNPSNIEATDAIFRNSK.....LLVEPVIVE	135
YFDLDVYDTSQ.....IPKDLVDQKLIKLRHILPLIQR...GQILYVATSNPSNIEATDAIFRNSK.....LLVEPVIVE	135
LLDLGVFDPAL.....FLDKDIDEXLIKRYRIMPLVHR...GHVLYVATSNPTNIEAMDAIFRNSK.....LKVEPIVE	135

logo

NTDB id 1410 SGL RS13950 WP 010873519.1
NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1168 A1552VC RS11070 WP 000957200.1
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consensus



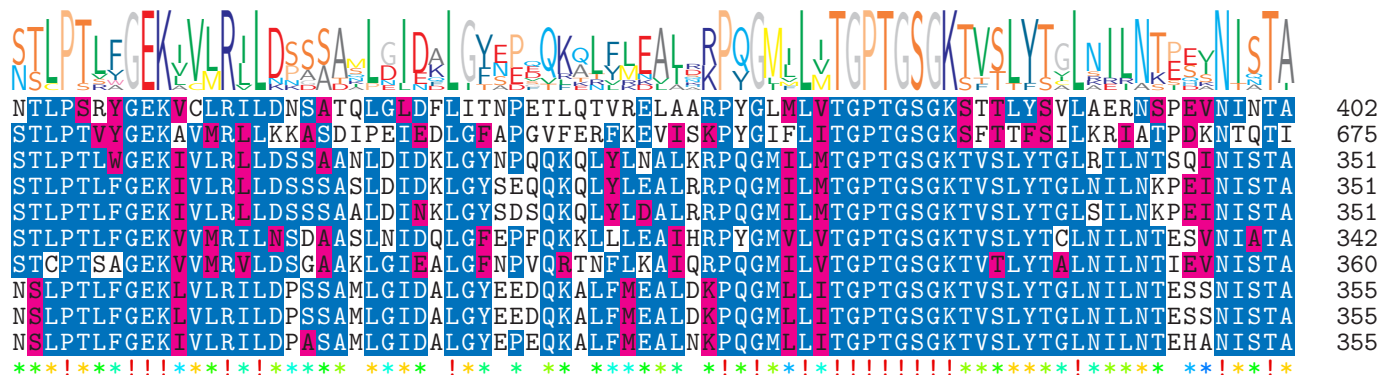
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consensus



logo

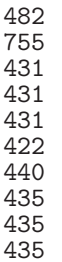
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consensus



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NTDB id 1410 SGL RS13950 WP 010873519.1
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consensus

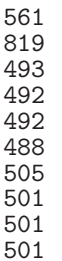
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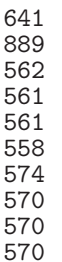
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NTDB id 1110 NGFG RS09235 WP 003689817.1
NTDB id 1251 GC085 RS07725 WP 011946523.1
NTDB id 1110050 ACQCIC RS17055 WP 001274990.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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NTDB id 1410 SGL RS13950 WP 010873519.1
NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1168 A1552VC RS11070 WP 000957200.1
NTDB id 1293 VP RS12240 WP 005479695.1
NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 1110 NGFG RS09235 WP 003689817.1
NTDB id 1251 GC085 RS07725 WP 011946523.1
NTDB id 1110050 ACQCIC RS17055 WP 001274990.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus
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logo

NTDB id 1410 SGL RS13950 WP 010873519.1	KRKSALTCRTCAAELQQEWLDCPYCMTSRFS	672
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NTDB id 1110050 AC0CIC RS17055 WP 001274990.1		570
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved