

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1355 HI 0298 AAC21962.1		0
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
NTDB id 422400 G3U99 RS06590 WP 165579506.1		0
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		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 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
NTDB id 1355 HI 0298 AAC21962.1		0
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
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NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	240
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[illegible]

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Sequence	Position
DPGAPILLPEDI	1-10
LCRRYG	11-15
VFP	16-18
PHRLE	19-23
GNRLVLLMK	24-29
DP	30
NRN	31-33
ILALDD	34-37
VR	38
LAL	39-41
LR	42
LR	43
LR	44
LR	45
LR	46
LR	47
LR	48
LR	49
LR	50
LR	51
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LR	95
LR	96
LR	97
LR	98
LR	99
LR	100

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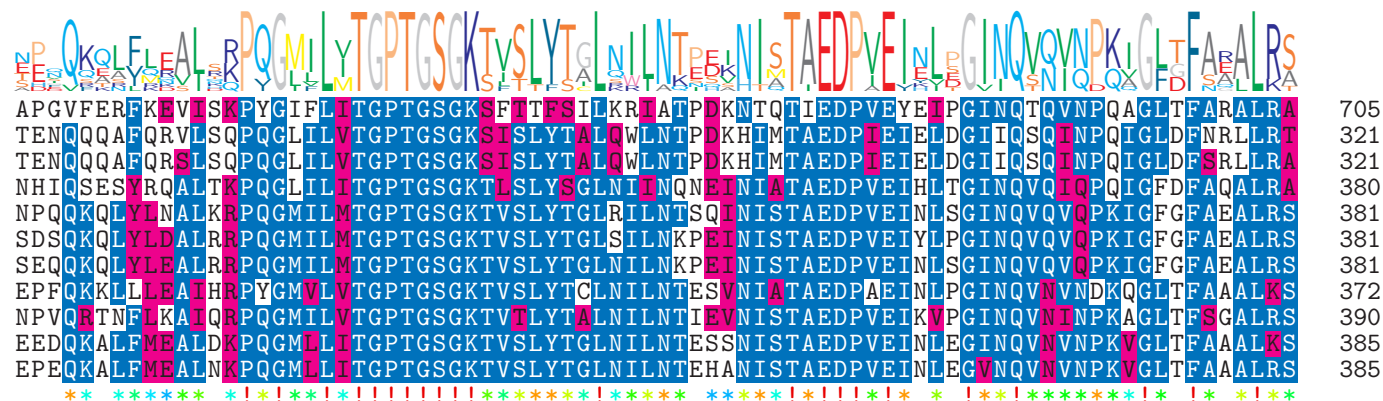
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- ☐ non conserved
- ☒ similar
- ☒ $\geq 50\%$ conserved