

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1384 A4U84 RS08525 WP 010787006.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 240700 CF389 RS02830 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSEEEVDTGLWVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
NTDB id 1384 A4U84 RS08525 WP 010787006.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSEEEVDTGLWVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
NTDB id 240700 CF389 RS02830 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	240
NTDB id 1384 A4U84 RS08525 WP 010787006.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
NTDB id 240700 CF389 RS02830 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	LRS DALRYGAVPIGFQNGEVEVWLS D PRHKEA VAQLLN RPARFYLA LPQAW EELFR RAYPQKNRLGEVLVQEGKLSREAL	320
NTDB id 1384 A4U84 RS08525 WP 010787006.1	LRS DALRYGAVPIGFQNGEVEVWLS D PRHKEA VAQLLN RPARFYLA LPQAW EELFR RAYPQKNRLGEVLVQEGKLSREAL	0
NTDB id 240700 CF389 RS02830 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPYCDPEEDPI	399
NTDB id 1384 A4U84 RS08525 WP 010787006.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPPYDPEEDP.	14
NTDB id 240700 CF389 RS02830 WP 003153083.1	MSQYTICEQQTERI	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	*	

logo

NTDB id 1028 TT RS08215 WP 011173991.1	FDISPSRWQPNLLPKQLRRYLVYPVBLTGRLWLVKDPRLNTALEIVRIMTRVQYEPVWITSDELTLLNALS	471
NTDB id 1384 A4U84 RS08525 WP 010787006.1	PDPGAPLLLLPEDLCRRYGVPFHRLEGNRIVLLMKDPRNITLALDDVRLALKRKGLNVEVAPAVATEAAITKLLI....	89
NTDB id 240700 CF389 RS02830 WP 003153083.1	FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDDENNLTACEIFAFM.TH....KQIEPVVIASDELKYLNNALS	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	*	

logo

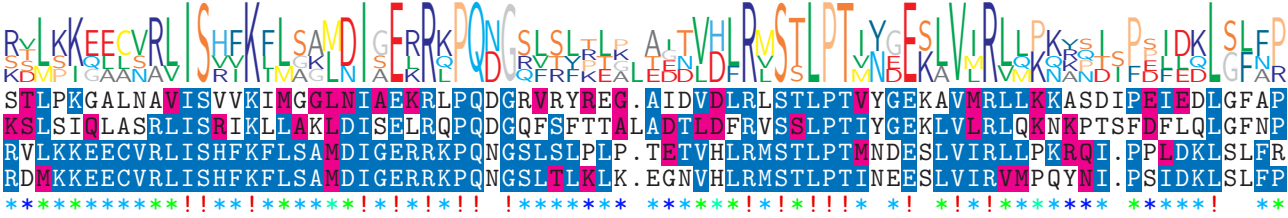
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NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 240700 CF389 RS02830 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus



548
158
45
45

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 240700 CF389 RS02830 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus



627
238
123
123

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 240700 CF389 RS02830 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus



706
317
203
203

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 240700 CF389 RS02830 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus



786
397
283
283

logo

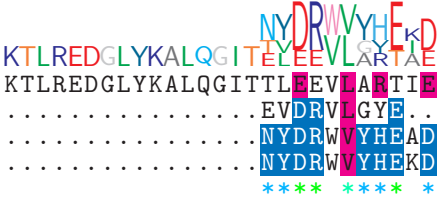
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NTDB id 96 BSU 24730 NP 390353.1
consensus



862
452
345
345

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 240700 CF389 RS02830 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus



889
461
356
356

 non conserved
 similar
 $\geq 50\%$ conserved