

logo

	MSXLSIGDRLGAALLDAGLLTDEELQRALERHREVEGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEI	75
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVEGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEI	75
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQP	74
NTDB id 1110 NGFG RS09235 WP 003689817.1	0
NTDB id 1168 A1552VC RS11070 WP 000957200.1	0
NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
NTDB id 1251 GC085 RS07725 WP 011946523.1	0
NTDB id 1145261 PHAGEPCYII10 RS23985 WP 003141354.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus		0

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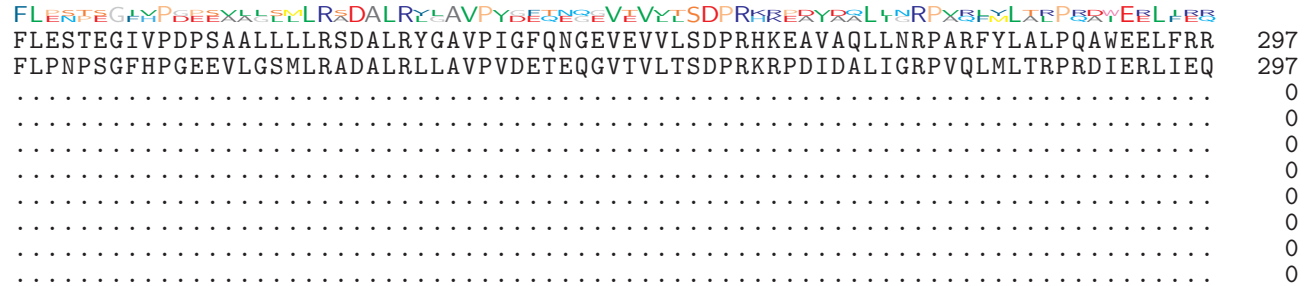
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NTDB id 1028 TT RS08215 WP 011173991.1	PPKVKALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPEL	150
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NTDB id 1293 VP RS12240 WP 005479695.1	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus		0

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	GLEIYVPSGASDAGRTGGKLGELLYKGYIIEKLYALYQERTGSLGRTLYSREKLYEXLAEQEGTT	222
NTDB id 1028 TT RS08215 WP 011173991.1	GLPVPPP..PSGEGQKDLKLGELLQKGWISREALEEALVEQEKTDGDLGRILV.RKGLPEEALYRALAEQKGLE	222
NTDB id 1306 DR RS10060 WP 027479822.1	GLEIIVPSGASDAGRTGGKLGELLYKGYIIEKLYALYQERTGSLGRTLYSREKLYEXLAEQEGTT	222
NTDB id 1110 NGFG RS09235 WP 003689817.1	0
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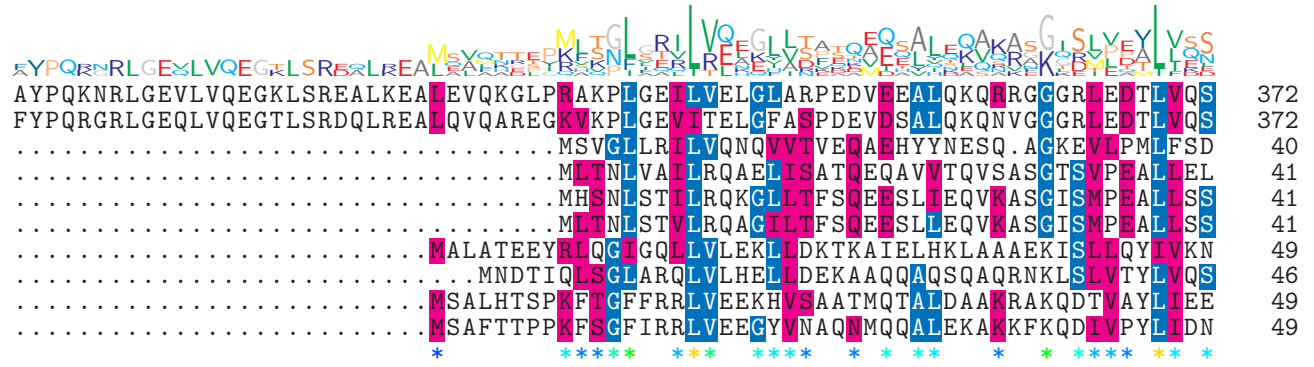
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