

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTIIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
NTDB id 1355 HI 0298 AAC21962.1		0
NTDB id 1063103 ACG4IZ RS02325 WP 000957200.1		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 1410 SGL RS13950 WP 010873519.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
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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 1410	SGL RS13950 WP 010873519.1	LRSDALRYGAVPIIGFQNGEVEVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	6
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NTDB id 1168	A1552VC RS11070 WP 000957200.1		0
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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 1410	SGL RS13950 WP 010873519.1	SKKSRAVALRNYFSPFGNKLIVAGGHVDAEQLRQALIVQVKKTGRSLPEELKAVTGREISPELLRQYKKNQLFELIKVLYGVD	86
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NTDB id 1293	VP RS12240 WP 005479695.1	MHSNLSLILRQKGLLTFQSQEESLLEQVKASGISMPEALLSS..GFFTSSELAEH.....LSSIFGLS	60
NTDB id 1110	NGFG RS09235 WP 003689817.1	MSVGLLRILVQNGVVTVEQAQEHYY.NESQAGKEVLPMLFSD..GVISPKSLAAL.....IARVFSYS	59
NTDB id 1251	GC085 RS07725 WP 011946523.1	..MALATEEYRLQGIQGLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVKN..KILSAEQIALT.....AAQNFQVP	68
NTDB id 1058	ABD1 RS01615 WP 001274986.1	..MSALHTSPKFTIGFFRRLLVEEKHVSAAATMQTALDAAKRAKQDVTAYLIEE..VHLSPSLLAET.....ISAFAEP	68
NTDB id 1018	ACIAD RS01685 WP 004920473.1	..MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDITVPLYIDN..FSISPLTIAEI.....ISLEFGEF	68
consensus			

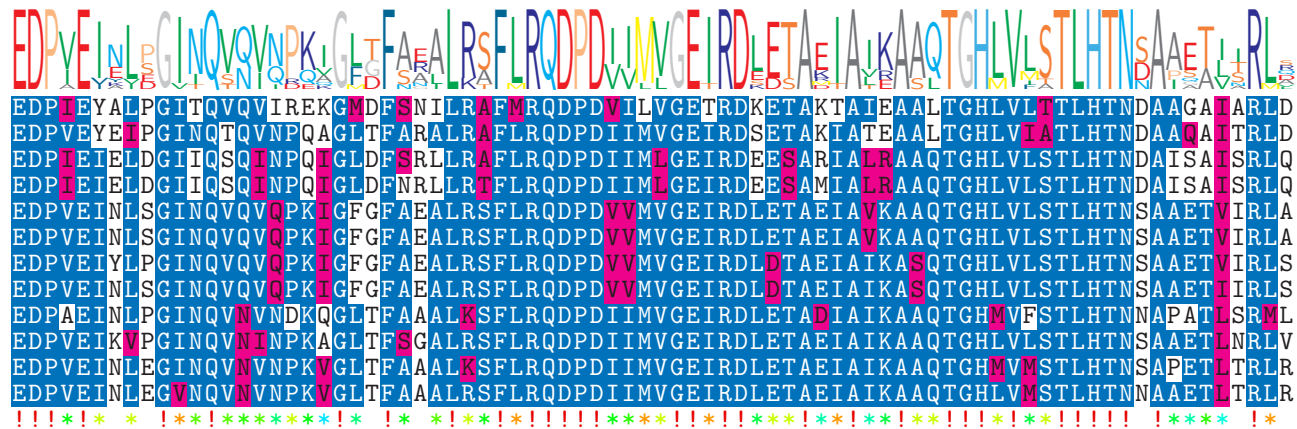
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NTDB id 1168	A1552VC RS11070 WP 000957200.1	ETDLSRYDYAN....LCQQGLRELITRYDALPIAKQG...NLLLLAVSDPTLLQAEEDFRFATG.....LQVELALAD	127
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NTDB id 1110	NGFG RS09235 WP 003689817.1	ILDRLHYPRHR....VLMGVLTEEQMVFEHCVVFRRG...DKVFFAVSDPTQMPQIQKTVSAAG.....IAVELVIVE	126
NTDB id 1251	GC085 RS07725 WP 011946523.1	MLDINCIDVGT....IPVNLVNEKLIKRRHAMVPLFSRG...TNLYLATDDPSKQASLKEIQFHTG.....LNTHAIVVE	135
NTDB id 1058	ABD1 RS01615 WP 001274986.1	YFDLDVYDTSQ....IPKDLVDQKLILKHRILPLIQRG...QILYVATSNPSNIEAIDAIRFNSK.....LLVEPVIVE	135
NTDB id 1018	ACIAD RS01685 WP 004920473.1	LLDLGVFDPAL....FLKDKIDEKLIQKYRIMPLVHRG...HVLVYVATSNPTNIEAMDIAIRFNSK.....LKVEPIIVE	135
consensus			

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NTDB id 1410 SGL RS13950 WP 010873519.1
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NTDB id 1366 NTHI RS01950 WP 011271972.1
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consensus

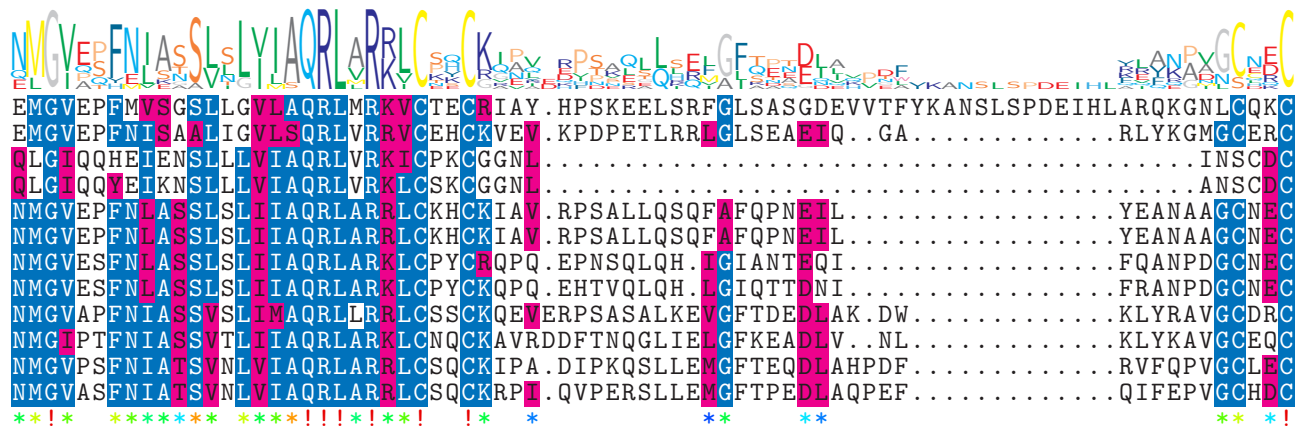


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NTDB id 1410 SGL RS13950 WP 010873519.1
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
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consensus



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NTDB id 1410 SGL RS13950 WP 010873519.1
NTDB id 1028 TT RS08215 WP 011173991.1
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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	KRKSALTCRTCAAELQQEWLDCPYCMTSRFS	672
NTDB id 1028 TT RS08215 WP 011173991.1		889
NTDB id 1366 NTHI RS01950 WP 011271972.1		464
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NTDB id 1063103 ACG4IZ RS02325 WP 000957200.1		562
NTDB id 1168 A1552VC RS11070 WP 000957200.1		562
NTDB id 1401 DSB67 RS12665 WP 010643259.1		561
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved