

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MSXLSIGDRRLGAALLDQGYLDNBLQRALERHSEVGGRLA5VLYD5GYSERRIA5AIE55GIPLVNLLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1410 SGL RS13950 WP 010873519.1		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 390819 F8A90 RS03700 WP 200019034.1		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 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
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consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	LSMLRSDALRYLAVPYGESNVEVYTSDPKRKRERYRLLNRPXSLMTRPSTYEELEHSEYPQRNRLGEXLVQEGGLSR	317
NTDB id 1028 TT RS08215 WP 011173991.1	GSMLRADALRLLAVPVDETEQQVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLLGEQLVQEGTLSR	317
NTDB id 1410 SGL RS13950 WP 010873519.1	LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR	3
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	DQLREALQVQAREGKVKPLGEVITELGFASPDVEVDSALQKQNVGGRIEDTLVQSG..KLSPEMPLA.....RSLAAQL	388
NTDB id 1028 TT RS08215 WP 011173991.1	EALKEALEVQKGLPRAKPLGEILVELGLIARPEVVEEALQKQRGGRIEDTLVQSG..KLRPEALA.....QAVATQL	388
NTDB id 1410 SGL RS13950 WP 010873519.1	SSSSKKSRAVALRNYFSPFGNKLVLVAGGHVDAEQLRQALVQVVKKTGRSLPEELKAVTGRELSPPELLRQYKKNQLFELKVLV	83
NTDB id 1168 A1552VC RS11070 WP 000957200.1	MLTNLVAITLRQAEILSATQEAVVTQVSASGTSVPEALLELS..IFHAQELT.....EQLSHIF	57
NTDB id 1401 DSB67 RS12665 WP 010643259.1	MLTNLSTVLRQAGILTFSCHEESLLEQVKASGISMPEALLSSG..FFQSHELT.....EHLSSVF	57
NTDB id 1293 VP RS12240 WP 005479695.1	MHSNLSITLRQKGLLTFSCHEESLLEQVKASGISMPEALLSSG..FFTSSELA.....EHLSSIF	57
NTDB id 1110 NGFG RS09235 WP 003689817.1	MSVGLLRILVQNQVVTVEQAHEHY..NESQAGKEVLPMLFSDG..VISPKSLA.....ALIARVF	56
NTDB id 1251 GC085 RS07725 WP 011946523.1	MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVKNK..ILSAEQIA.....LTAAQNF	65
NTDB id 390819 F8A90 RS03700 WP 200019034.1	MPQSEHMLKGFAARLVDSGLLEPQAAQLAEQEANEAGESLLHHIIAHG..LVTARDAT.....LAAAWAY	63
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MSALHTSPKFTGFFRRLVEEKHVSAAATMQTALDAAKRAKQDTVAYLIEEV..HLSPSLLA.....ETISAEF	65
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MSAFTTPPKFSGFIRRLVEEGYVNAQNMQALEKAKKFKQDTPVYLIIDNF..SISPLTIA.....EIIISLEF	65
consensus	* * * * *	

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NTDB id 1306 DR RS10060 WP 027479822.1	GYEFLDPIQNPPDPKV....ALMIPAEATARRVVPVRLQG...NSLVVAMKDPRNVFALDDKLITG.....KEILPA	455
NTDB id 1028 TT RS08215 WP 011173991.1	GYFYVDPEEDPPDPGA....PLLIPEDLCRRYGVFPHRLEG...NRLVLLMKDPRNIALALDDVRLALKRKG...GLNVEVAPA	460
NTDB id 1410 SGL RS13950 WP 010873519.1	GVDSVDPEVAPIATQQMAELIGRFFPLDTCRRFKFLPLAQEGDPPSVLVAMVDPDNLAQDELNRILRVKG..FELRRL	161
NTDB id 1168 A1552VC RS11070 WP 000957200.1	GLPETDLSRYDYANLC....QQGLGRELITRYDALPIAKQG...NLLLLAVSDPTLLQAEFEF...RFATG..LQVELA	124
NTDB id 1401 DSB67 RS12665 WP 010643259.1	GLSCTSLNQYEYASLC....QTLGLRLDITRHNAALPLNRTS...STLILAVADPTNLQAEDDF...RFATG..LQVELV	124
NTDB id 1293 VP RS12240 WP 005479695.1	GLSQPELSQYEYASLC....QQGLGRELITRHNAALPLHRT...STLLAVADPTNQQAEDDF...RFATG..LQVELV	124
NTDB id 1110 NGFG RS09235 WP 003689817.1	SYSILDLRHYPRHRVL....MGVLTTEEQMVVEFHCVPVFRRG...DKVFFAVSDPTQMPQIQKT...VSAAG..IAVELV	123
NTDB id 1251 GC085 RS07725 WP 011946523.1	GVPMLDINCIIVGTIP....VNLVNEKLIKRRHAMVPLFSRG...TNLYLATDDPSKQASLKEI...QFHTG..LNTHAI	132
NTDB id 390819 F8A90 RS03700 WP 200019034.1	GLPYVDLDALRIDS LPS....IDELPEKVLRLKGLVPLPLYRRE...HRLTVGVSPSTLTHLDEL...QFATG..LSIESV	131
NTDB id 1058 ABD1 RS01615 WP 001274986.1	AEPYFDLDVYDTSQIP....KDLVDQKLILKHRILPLIQRG...QILYVATSNPSNIEAIDAI...RFNSK..LLVEPV	132
NTDB id 1018 ACIAD RS01685 WP 004920473.1	GEPLLDLGVFDPALFL....KDKIDIEKLIQKYRIMPLVHRG...HVLVYVATSNPTNIEAMDAI...RFNSK..LKVEPI	132
consensus	*** **	

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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 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

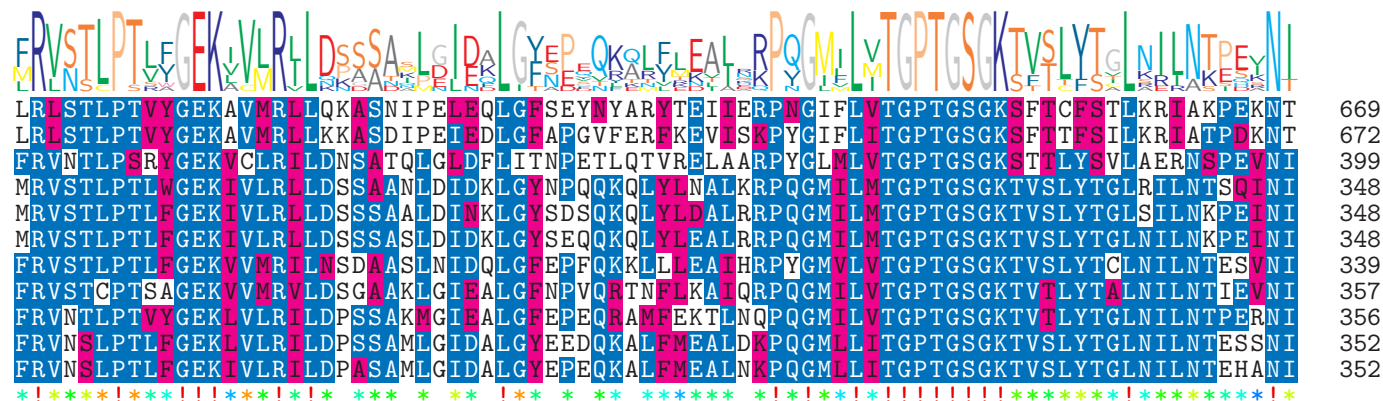
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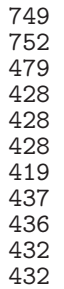
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

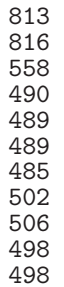
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NTDB id 390819 F8A90 RS03700 WP 200019034.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus
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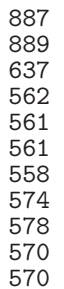
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NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 1293 VP RS12240 WP 005479695.1
NTDB id 1110 NGFG RS09235 WP 003689817.1
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NTDB id 390819 F8A90 RS03700 WP 200019034.1
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consensus
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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 1306 DR RS10060 WP 027479822.1	EMRAKRKSALTCRTCAAELQQEWLDCPYQMTSRFS	887
NTDB id 1028 TT RS08215 WP 011173991.1		889
NTDB id 1410 SGL RS13950 WP 010873519.1	EMRAKRKSALTCRTCAAELQQEWLDCPYQMTSRFS	672
NTDB id 1168 A1552VC RS11070 WP 000957200.1		562
NTDB id 1401 DSB67 RS12665 WP 010643259.1		561
NTDB id 1293 VP RS12240 WP 005479695.1		561
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NTDB id 1251 GC085 RS07725 WP 011946523.1		574
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

- X non conserved
- X similar
- X ≥ 50% conserved