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consensus		

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NTDB id 1401 DSB67 RS12665 WP 010643259.1	MLTNLSTVIRQAGILTFSSQEESSLLEQVKASGISMPEALLSS..GFFQ.SHEITEH.....L	53
NTDB id 1293 VP RS12240 WP 005479695.1	MHSNLSTIRQKGLLTFSQEESSLIEQVKASGISMPEALLSS..GFFT.SSELAEH.....L	53
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	MSAFTTPPKFSGFIRRIVEEGYVNAQNMQQALEKAKKFKQDIVPYLIDN..FSIS.PLTIAEI.....I	61
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MSALHTSPKFTGFFRRIVEEKHVSAAATMQTALDAAKRAKQDTVAYLIEE..VHLS.PSLIAET.....I	61
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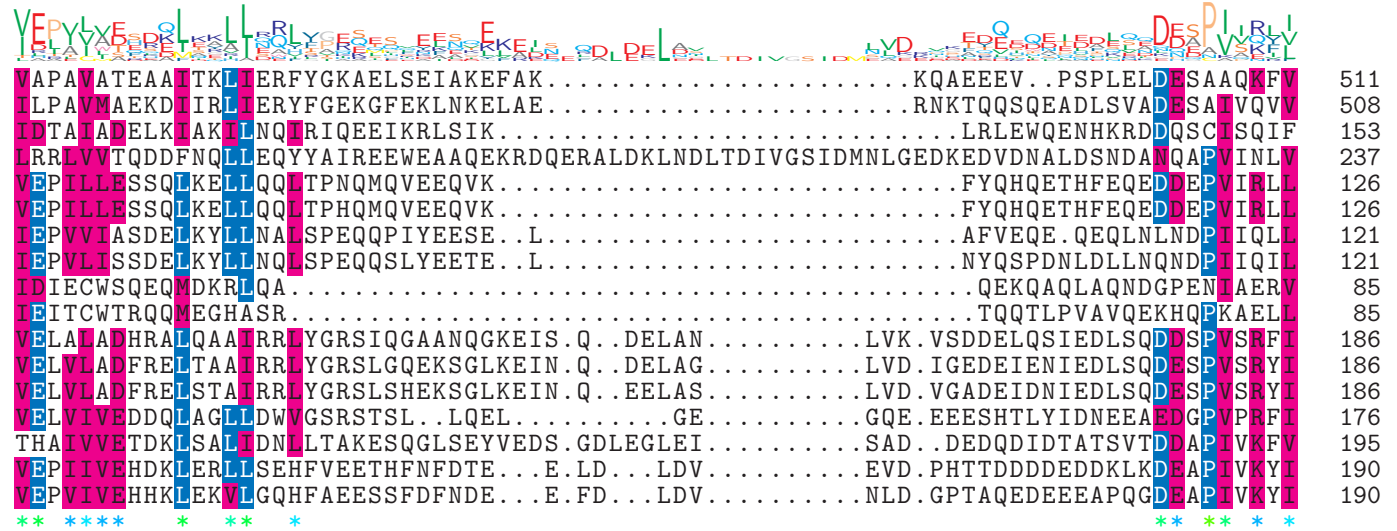
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NTDB id 1306 DR RS10060 WP 027479822.1	AAQLGYEFLDPIQNPP....DPKVALMIPEATARRVVVVRQLQG...NSLVVAMKDPRNVFALDDKLITG....KE	451
NTDB id 1227 CJJ81176 RS07050 WP 011812800.1	ANFCNLHFVDLDE.....DNDFIYEGIPFSLILKFKFLILKNND...GFMIRSKPCSLELLEQVKTF..MIC..EK	102
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NTDB id 1366 NTHI RS01950 WP 011271972.1	GEIF.....T...ISP....DLWERNQQQSSLLRLRYFALPLKEEN...NRLWLGVDLSLSNISACETIAF...ITG..KP	75
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NTDB id 1436 BW25113 RS00530 WP 001025145.1	MNIPQLTALCLRYHGVLLDASE...EVVHVAVVDAPSHELLDALHF...ATT..KR	48
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	SAEFAEYPFDLDVYDT....SQIPKDLVDQKLIKLRHILPLIQRG...QILYVATSNPSNIBATDAIRF...NSK..LL	128
consensus	*.....*	

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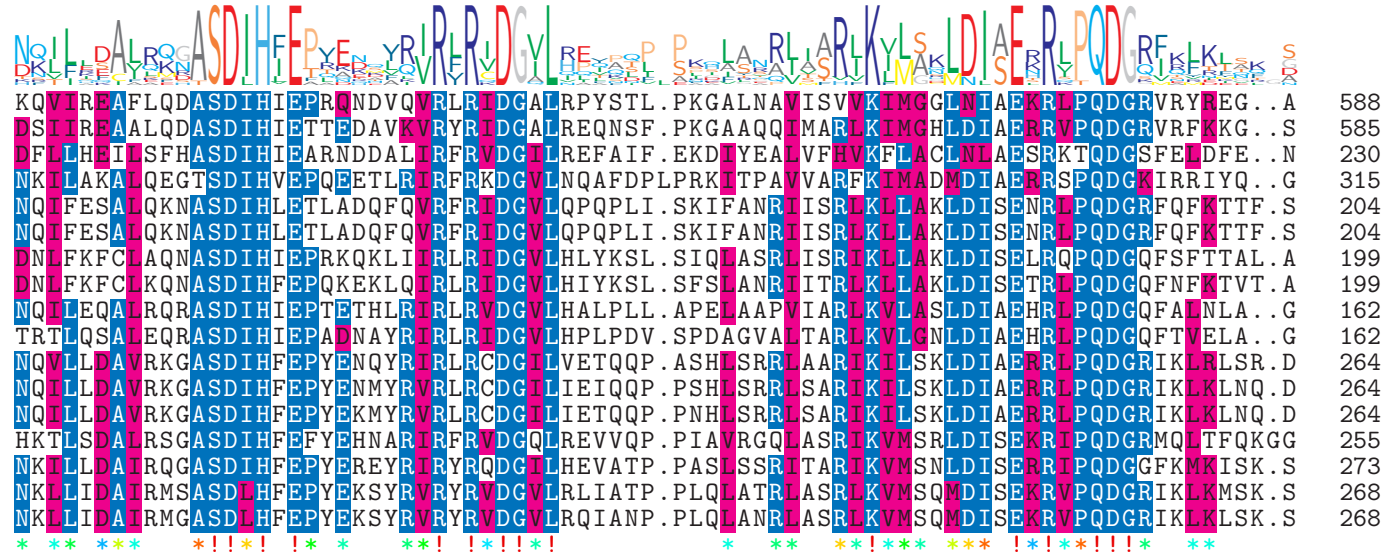
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
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RKVDFRVNTLPSRYGEKVCLRLIDNSATQGLDFLITNPETLQTVRELAARPYGLMLVTGPTGSGKSTTLYSVLAERNISP	395
DILDFRLSTLPTHGKEKIVLRAQQNKPVLSFSELGMTENQQQAFQRVLSQPQGLILVTGPTGSGKSISLYTALQWLNTP	284
DILDFRLSTLPTHGKEKIVLRAQQNKPVLSFAELGMTENQQQAFQRVLSQPQGLILVTGPTGSGKSISLYTALQWLNTP	284
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Sequence	Position
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DINICSVEDPVEIPIAGLNQQTIHPRAGLTFQGVLRALLRQDPDVIMIGEIRDGETAEIAIKAAQTGHLVLSTLHTNSTC	322
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NTDB id 1306 DR RS10060 WP 027479822.1
NTDB id 1227 CJJ81176 RS07050 WP 011812800.1
NTDB id 1410 SGL RS13950 WP 010873519.1
NTDB id 1355 HI 0298 AAC21962.1
NTDB id 1366 NTHI RS01950 WP 011271972.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1468 ASU2 RS05090 WP 014991734.1
NTDB id 77965 PMK1 RS12740 WP 004145935.1
NTDB id 1436 BW25113 RS00530 WP 001025145.1
NTDB id 1168 A1552VC RS11070 WP 000957200.1
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NTDB id 1293 VP RS12240 WP 005479695.1
NTDB id 1110 NGFG RS09235 WP 003689817.1
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[illegible]

logo

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NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved