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logo
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NTDB id 1177 A1552VC RS01165 WP 000422572.1  ....MELNQYLDGMLTHKASDLYITVGAPILYRVDGELRAQGEAL.SVADVTALLHAMMDDARQAEFKQTREANFAVVRD   75
NTDB id 1203 PSJM300 01605 AFN76401.1        ....MEFEKLLRLMVEKGGSDLFITAGVPPSMKVNGKILPVNKTPMSPEMTRETVHGVMNEQQRREFTENHECNFAISAR   76
NTDB id 1019 ACIAD RS04205 WP 004922051.1    ....MDFNDLLNLMAKHKASDLFITAGVEPSLKINGEITPVSKTKLDGDMIAQLIDSITTQKQRQEFIETHECNFAILNR   76
NTDB id 1307 DR RS10055 WP 010888596.1       MTSGIDITDILRLAADKGASDIILAAGLPPQFKLSGDYDAQGFEPLSASDTRKLMYSMMNERQQRIFEEKRELDFSFALG   80
NTDB id 1185 GTF74 RS11940 WP 000350195.1    ....MDIAELLEFSVKHNASDLHLSAGVPPMVRIDGEVRKLGVPAFTHSDVHRLIFEIMNDAQRSEYEEKLEVDFSFELP   76
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consensus                                     *** **! ***** *!!*****!********!******** *** ** ****************** *! *******

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NTDB id 1307 DR RS10055 WP 010888596.1       D.KARFRVNAFMQRGNVGGVMRLIPTTVRTIAEMGLPQTVIDIANAPRGLVLVTGPTGSGKSTTLASMIDHINTTKKLHI   159
NTDB id 1185 GTF74 RS11940 WP 000350195.1    N.VGRFRVNAFHQARGCSAVFRTIPTVIPTLEQLDAPEIFSKIANYEKGLVLVTGPTGSGKSTTLAAMVNYVNAHHNKHI   155
NTDB id 1176 A1552VC RS01160 WP 000350195.1  N.VGRFRVNAFHQARGCSAVFRTIPTVIPTLEQLDAPEIFSKIANYEKGLVLVTGPTGSGKSTTLAAMVNYVNAHHNKHI   155
NTDB id 1283 ACE17W RS02630 WP 010947729.1   N.LARFRVNAFNQSRGAAAVFRTIPSEILSMEDLGLPPIFKEMASFSRGLVLVTGPTGSGKSTTLAAVIDYINSSRYEHI   155
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consensus                                    * **!!!!**!*! *****!*!*!** * ******** ******   *!*****!*!!*!!!!**!******!  *  !!

logo
NTDB id 269136 C3F36 RS02055 WP 029314740.1  ITIEDPVEFVHRHQKSIINQREVGVDTRSFHAALKNTLRQAPDVILIGEIRDRETMEHALAFSETGHLAISTLHANNANQ   235
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NTDB id 1203 PSJM300 01605 AFN76401.1        ISIEDPIEFIHQHQNCIVTQREVGIDTDSFEVALKNTLRQAPDVILIGEVRTRETMDHAVAFAETGHLCLATLHANNANQ   235
NTDB id 1019 ACIAD RS04205 WP 004922051.1    ITIEDPIEFIHEHAGCIITQREVGIDTDSFEIALKNTLRQAPDVILIGEIRSREVMDYALAFAETGHLVFATLHANNANQ   236
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NTDB id 1185 GTF74 RS11940 WP 000350195.1    LTIEDPIEFVHSNNKCLINQREVHRDTHSFKNALRSALREDPDVILVGELRDQETISLALTAAETGHLVFGTLHTSSAAK   235
NTDB id 1176 A1552VC RS01160 WP 000350195.1  LTIEDPIEFVHSNNKCLINQREVHRDTHSFKNALRSALREDPDVILVGELRDQETISLALTAAETGHLVFGTLHTSSAAK   235
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NTDB id 1060 ABD1 RS04380 WP 000355489.1     LTVEDPIEFVHQSKKCLINQREVHRDTHGFNEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAK   235
NTDB id 1040 HON27 RS13360 WP 000355489.1    LTVEDPIEFVHQSKKCLINQREVHRDTHGFNEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAK   235
NTDB id 1081 FDQ49 RS05335 WP 006581387.1    LTVEDPIEFVHQSKKCLINQREVHRDTHGFNEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAK   235
NTDB id 1202 PSJM300 01600 AFN76400.1        LTIEDPIEFVHESKKCLVNQREVHRDTLGFSEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAK   235
NTDB id 1194 PAKAF RS01995 WP 003084552.1    LTIEDPIEFVHESKKCLVNQREVHRDTLGFSEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAK   235
consensus                                    ***!!!*!!*!   ******!!!!!**!!  **   !!***!!**!!*!!*!!*!* !****!****!!!!!***!!!* *! *
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logo
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NTDB id 1177 A1552VC RS01165 WP 000422572.1      ALERILHLVPKEQREQFLLDLSLNLKGVIAQQLLRDKNGKGRHGVFEVLLNSPRIADLIRRGELHELKATMARSQEVGMQ   313
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NTDB id 1019 ACIAD RS04205 WP 004922051.1        TLERIIHFFDADRHQQIFMDLSLNLKAVISQQLIPTHDGQSRRAAVEVLINTPLLSDYIRQGEIYKIKELMARSGELGMQ   316
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NTDB id 1283 ACE17W RS02630 WP 010947729.1       TINRIIDVFPAEEKSMVRSMLSESLQAVVAQSLLKK.NKGGRVAALEIMMCTGAIRNLIREDKIAQMYSSIQTGQAKGMQ   314
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NTDB id 1202 PSJM300 01600 AFN76400.1            TIDRVVDVFPAEEKSMVRSMLSESLQAVISQTLLKK.VGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSSIQTGGSLGMQ   314
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consensus                                        ***!********** *** *!***! ****! !***  ***!*** !***********!** ******* *****  !!*
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logo
                                                  TLDQLKGLVARGVLSYQTAREAAKS PELF
                                                  FLSNIEIYKNNTDRDVRRNRIFLKKDRDNKSMEEAIRKIDRNKSMMEXTARLFARRDVTRRRRIFLKKDRLMIRTRRDRNALHSTRRAIRKKE.VPAYNSASRRSTRRRDLFRRGAAAP
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NTDB id 1307 DR RS10055 WP 010888596.1           TMDAYLAGLFRRHLITYDKGLERAID.PKEF..ARLANDASAGLGGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAAP   396
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NTDB id 1259 GC085 RS10305 WP 010947729.1        TLDQHLTELVNKNIISKHTAHEAALN.KSLF...................................................   344
NTDB id 1060 ABD1 RS04380 WP 000355489.1         TLDQSLKGLVARGVISPQTARTAAKQ.PESF..L................................................   345
NTDB id 1040 HON27 RS13360 WP 000355489.1        TLDQSLKGLVARGVISPQTARTAAKQ.PESF..L................................................   345
NTDB id 1081 FDQ49 RS05335 WP 006581387.1        TLDQSLKGLVSRGVISPQTARTAAKQ.PESF..L................................................   345
NTDB id 1202 PSJM300 01600 AFN76400.1            TLDSCLKGLLAKGLISRDSAKEKAKQ.PENF...................................................   344
NTDB id 1194 PAKAF RS01995 WP 003084552.1        TLDMCLKGLVAKGLISRENAREKAKI.PENF...................................................   344
consensus                                        !*** *  !* *****   *   *      *
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logo
                                                  ATPAAPAAGRSEPGRAGGFGRR
NTDB id 269136 C3F36 RS02055 WP 029314740.1      .......................   378
NTDB id 1177 A1552VC RS01165 WP 000422572.1      .......................   368
NTDB id 1203 PSJM300 01605 AFN76401.1            .......................   381
NTDB id 1019 ACIAD RS04205 WP 004922051.1        .......................   383
NTDB id 1307 DR RS10055 WP 010888596.1           ATPAAPAAGRSEPGRAGGFGRR   418
NTDB id 1185 GTF74 RS11940 WP 000350195.1        .......................   345
NTDB id 1176 A1552VC RS01160 WP 000350195.1      .......................   345
NTDB id 1283 ACE17W RS02630 WP 010947729.1       .......................   344
NTDB id 1259 GC085 RS10305 WP 010947729.1        .......................   344
NTDB id 1060 ABD1 RS04380 WP 000355489.1         .......................   345
NTDB id 1040 HON27 RS13360 WP 000355489.1        .......................   345
NTDB id 1081 FDQ49 RS05335 WP 006581387.1        .......................   345
NTDB id 1202 PSJM300 01600 AFN76400.1            .......................   344
NTDB id 1194 PAKAF RS01995 WP 003084552.1        .......................   344
consensus
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X non conserved
X similar
X $\geq$ 50% conserved