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logo
NTDB id 1177 A1552VC RS01165 WP 000422572.1  ....MELNQYLDGMLTHKASDLYITVGAPILYRVDGELRAQGEAL.SVADVTALLHAMMDDARQAEFKQTREANFAVVRD   75
NTDB id 141685 PSPO RS10460 WP 010561718.1   ...MHGLDHFLKQMVESEASDLFVSTFLPVSVKVNGELRPIGHSDLTEEEALELVESAMTGKQKTEFHETKECNFAIARD   77
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 1020 ACIAD RS04210 WP 011182164.1    ....MDITELLAFSVQNRASDLHLSAGIPPMIRVDGEVRRINLPAMDHSEVHNLIYDIMNDKQRRDYEEKLETDFSFEVP   76
consensus                                     ***  *!  **   *!!********** *** !**  *    **  *   **  ** **** ** *  ! *!**

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NTDB id 1177 A1552VC RS01165 WP 000422572.1  ..SGRFRVSAFFQRELPGAVIRRIETRIPTFEELKLPEVLQNLAIAKRGLVLVVGATGSGKSTTMAAMTGYRNQHRTGHI  153
NTDB id 141685 PSPO RS10460 WP 010561718.1  ..SGRFRVSAFWQRDKAGMVVRRIVTTIPEVHDLGLPPVLTDVIMSKRGLVLFVGGTGTGKSTSLAALLGYRNRNQRGHI  155
NTDB id 1203 PSJM300 01605 AFN76401.1       G.IGRFRVSAFYQRNLAGMVLRRIETNIPTIEDLKLPDVLKKLALTKRGLVLFVGATGTGKSTSLAAMIGYRNKNSSGHI  155
NTDB id 1019 ACIAD RS04205 WP 004922051.1   EKNERFRVSAFQQRDMPGMVIRRIETKIPSPDDLHLPPILKELAMSKRGIIIFSGATGTGKSTSLASMIQHRNQHSKGHI  156
NTDB id 1307 DR RS10055 WP 010888596.1      D.KARFRVNAFMQRGNVGGVMRLIPTTVRTIAEMGLPQTVIDIANAPRGLVLVTGPTGSGKSTTLASMIDHINTTKKLHI  159
NTDB id 1020 ACIAD RS04210 WP 011182164.1   N.LARFRVNVFNQNRGAGAVFRTIPSQVLTLEELGLGKIFRDICDYPRGLVLVTGPTGSGKSTTLAAMLDYINNHRYDHI  155
consensus                                     *!!!!**! !*    ! !*!*! * ******* !* ** **** *!!***  !*!!*!!!!**!*** **!    *!!

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NTDB id 1177 A1552VC RS01165 WP 000422572.1  LTVEDPIEFVHEHKRCIVTQREVGLDTESYEVALKNSLRQAPDMILIGEIRSRETMEYAMTFAETGHLCMATLHANNANQ  233
NTDB id 141685 PSPO RS10460 WP 010561718.1  LTIEDPIEFVHEHQKSIVTQREVGLDTESFESALKSSLRQAPDVILIGEIRSQETMEYALSFAETGHLCVATLHANNANQ  235
NTDB id 1203 PSJM300 01605 AFN76401.1       ISIEDPIEFIHQHQNCIVTQREVGIDTDSFEVALKNTLRQAPDVILIGEVRTRETMDHAVAFAETGHLCLATLHANNANQ  235
NTDB id 1019 ACIAD RS04205 WP 004922051.1   ITIEDPIEFIHEHAGCIITQREVGIDTDSFEIALKNTLRQAPDVILIGEIRSREVMDYALAFAETGHLVFATLHANNANQ  236
NTDB id 1307 DR RS10055 WP 010888596.1      LTIEDPIEFMHPNKASIINQREVGSDTLSFDDALRAALRQAPDVILVGEMRDYETIKAAVIAAETGHLVMGTLHTNSAPE  239
NTDB id 1020 ACIAD RS04210 WP 011182164.1   LTVEDPIEFVHEPKKCLINQREVHRDTLGFNEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAK  235
consensus                                   ***!!!!!!*!**  ****!!!!*!!**** !!* *!!**!!*!!*!!*!* !*** !***!!!!!! **!!!***!**

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NTDB id 1177 A1552VC RS01165 WP 000422572.1  ALERILHLVPKEQREQFLLDLSLNLKGVIAQQLLRDKNGKGRHGVFEVLLNSPRIADLIRRGELHELKATMARSQEVGMQ  313
NTDB id 141685 PSPO RS10460 WP 010561718.1  AIDRIMHLVPKEKHDKLKYDLALNLRAIVAQQLVPTSDGHGRAAAIEILLNSPMVAELIKRGEVGAIKETMSKSREMGMQ  315
NTDB id 1203 PSJM300 01605 AFN76401.1       ALDRIINFFPADRQNQVWMDLSLNLKAIVAQQLVPTPDGKGRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLGMQ  315
NTDB id 1019 ACIAD RS04205 WP 004922051.1   TLERIIHFFDADRHQQIFMDLSLNLKAVISQQLIPTHDGQSRRAAVEVLINTPLLSDYIRQGEIYKIKELMARSGELGMQ  316
NTDB id 1307 DR RS10055 WP 010888596.1      SIDRIVDVFPEEQQEQIRVQLANNLVAVMTQQLLPRADGQGRVLAYEILLANPAVRALIREGKTFQITSVMQTGAREGMV  319
NTDB id 1020 ACIAD RS04210 WP 011182164.1   TIDRIIDVFPAEEKDMVRTMLSESLQAVISQTLL.KKEGGGRVAAHEIMIGIPAIRNLIREGKVAQMYSSIQTGATHGMT  314
consensus                                    **!!* *** *  *** **!***!**** !*!**  *! *! ** !*****! * **!* !**  **  * ** **!!*

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NTDB id 1177 A1552VC RS01165 WP 000422572.1  TFDQALYQLVVDDKISEQDALHSADSANDLRLMLKTKRGDDYGSG...SLQNVK....IDME.................  368
NTDB id 141685 PSPO RS10460 WP 010561718.1  TFDQALFDLYQQHRINYADALHHADSPNDLRLMIKLRNNEQQGAG...FLQGVT....VDGLDGDKNQ............  376
NTDB id 1203 PSJM300 01605 AFN76401.1       TFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGSETDGEHLTSVS.QGLT....LEMSDDDPGRSFR.........  381
NTDB id 1019 ACIAD RS04205 WP 004922051.1   TFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSEEANKPLDNRERMSEME....RKMTFDGQRRNLK.........  383
NTDB id 1307 DR RS10055 WP 010888596.1      TMDAYLAGLFRRHLITYDKGLERAIDPKEFARLANDASAGLGGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAAPATP  399
NTDB id 1020 ACIAD RS04210 WP 011182164.1   TLDQSLKQLLAKGIINLPTARAVAKSPESF.......................................................  344
consensus                                   !*!**!* !*     !** ***  !***********                 *
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                                                AAPAAGRSEPGRAGGFGRR
NTDB id 1177 A1552VC RS01165 WP 000422572.1     ....................   368
NTDB id 141685 PSPO RS10460 WP 010561718.1      ....................   376
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          AAPAAGRSEPGRAGGFGRR   418
NTDB id 1020 ACIAD RS04210 WP 011182164.1       ....................   344
consensus
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X non conserved
X similar
X $\geq$50% conserved