```
logo
NTDB id 1307 DR RS10055 WP 010888596.1          MTSGIDITDILRLAADKGASDIILAAGLPPQFKLSGDYDAQGFEPLSASDTRKLMYSMMNERQQRIFEEKRELDFSFALG   80
NTDB id 1095974 ACMYQ0 RS06820 WP 415755301.1  ....MDLNAMLKILSSQDGSDLYLSTGAPPCAKFNGVLKALSAEPLKAGEVADIADAVMDVAQREEFERELEMNLAISIA   76
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
consensus                                          **** *!***    *!!*****!***  ***!** **   **** **  ** *** * ** *! *  ! ****

logo
NTDB id 1307 DR RS10055 WP 010888596.1          D.KARFRVNAFMQRGNVGGVMRLIPTTVRTIAEMGLPQTVIDIANAPRGLVLVTGPTGSGKSTTLASMIDHINTTKKLHI  159
NTDB id 1095974 ACMYQ0 RS06820 WP 415755301.1  G.VGRFRINIFKQRNEVSIVARNIKLDIPKFEDLKLPHVLLKTVMEKRGLVLFVGGTGSGKSTSLAALIDYRNRNSGGHI  155
NTDB id 1177 A1552VC RS01165 WP 000422572.1    ..SGRFRVSAFFQRELPGAVIRRIETRIPTFEELKLPEVLQNLAIAKRGLVLVVGATGSGKSTTMAAMTGYRNQHRTGHI  153
NTDB id 1203 PSJM300 01605 AFN76401.1          G.IGRFRVSAFYQRNLAGMVLRRIETNIPTIEDLKLPDVLKKLALTKRGLVLFVGATGTGKSTSLAAMIGYRNKNSSGHI  155
NTDB id 1019 ACIAD RS04205 WP 004922051.1      EKNERFRVSAFQQRDMPGMVIRRIETKIPSPDDLHLPPILKELAMSKRGIIIFSGATGTGKSTSLASMIQHRNQHSKGHI  156
consensus                                         *!!!***! !! * **!*!*!** *** ****!! **  *** *!!*****!*!!*!!!!**!*** **!  * *!!

logo
NTDB id 1307 DR RS10055 WP 010888596.1          LTIEDPIEFMHPNKASIINQREVGSDTLSFDDALRAALRQAPDVILVGEMRDYETIKAAVTAAETGHLVMGTLHTNSAPE  239
NTDB id 1095974 ACMYQ0 RS06820 WP 415755301.1  ITIEDPVEYVHKHKKSIINQREVGVDTRSFHAALKNTLRQAPDVILIGEIRDRETMEHALAFADTGHLAISTLHANNANQ  235
NTDB id 1177 A1552VC RS01165 WP 000422572.1    LTVEDPIEFVHEHKRCIVTQREVGLDTESYEVALKNSLRQAPDMILIGEIRSRETMEYAMTFAETGHLCMATLHANNANQ  233
NTDB id 1203 PSJM300 01605 AFN76401.1          ISIEDPIEFIHQHQNCIVTQREVGIDTDSFEVALKNTLRQAPDVILIGEVRTRETMDHAVAFAETGHLCLATLHANNANQ  235
NTDB id 1019 ACIAD RS04205 WP 004922051.1      ITIEDPIEFIHEHAGCIITQREVGIDTDSFEIALKNTLRQAPDVILIGEIRSREVMDYALAFAETGHLVFATLHANNANQ  236
consensus                                      ***!!!*!**! ** *!**!!!!!*!!*!***!!***!!!!!!*!!*!!*!**!*** !***!*!!!! **!!!*!*!**

logo
NTDB id 1307 DR RS10055 WP 010888596.1          SIDRIVDVFPEEQQEQIRVQLANNLVAVMTQQLLPRADGQGRVLAYEILLANPAVRALIREGKTFQITSVMQTGAREGMV  319
NTDB id 1095974 ACMYQ0 RS06820 WP 415755301.1  ALDRIINFFPEDRRPQLLNDLGNNLKAFVSQRLVKTIDGKR.RAAVEVLLGTPTIRDLIKRNEFSEIKEIMEKSKNLGMQ  314
NTDB id 1177 A1552VC RS01165 WP 000422572.1    ALERILHLVPKEQREQFLLDLSLNLKGVIAQQLLRDKNGKGRHGVFEVLLNSPRIADLIRRGELHELKATMARSQEVGMQ  313
NTDB id 1203 PSJM300 01605 AFN76401.1          ALDRIINFFPADRQNQVWMDLSLNLKAIVAQQLVPTPDGKGRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLGMQ  315
NTDB id 1019 ACIAD RS04205 WP 004922051.1      TLERIIHFFDADRHQQIFMDLSLNLKAVISQQLIPTHDGQSRRAAVEVLINTPLLSDYIRQGEIYKIKELMARSGELGMQ  316
consensus                                      ***!!* *** **  !* **!**!!*****!*!*** *!******!*!*!***! * **!**** *** *! ** **!!*

logo
NTDB id 1307 DR RS10055 WP 010888596.1          TMDAYLAGLFRRHLITYDKGLERAIDPKEFARLA...NDASA.GLGGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA  395
NTDB id 1095974 ACMYQ0 RS06820 WP 415755301.1  TFDQALIDLVNDGSIDEDEAVKNADSANNVRLKLKLYRDNPAAPVQAAAASAPAPAAA..EARSTTPAASAGEWGLELIL  392
NTDB id 1177 A1552VC RS01165 WP 000422572.1    TFDQALYQLVVDDKISEQDALHSADSANDLRLMLKTKRGDDY...GSGSLQN.....................VKIDME.  368
NTDB id 1203 PSJM300 01605 AFN76401.1          TFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGSETDGEHLTSVS.QG.....................LTLEMSD  373
NTDB id 1019 ACIAD RS04205 WP 004922051.1      TFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSEEANKPLDNRERMSE.....................MERKMTF  375
consensus                                      !*!**!* !*    !******  !** *********  *  *  *  ****                        * ****
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logo
                                                  PDTPEEP LPRSSGRNLRK
                                                  EAIEAAAAGPEEQRNAFGFGRR
NTDB id 1307 DR RS10055 WP 010888596.1            PATPAAPAAGRSEPGRAGGFGRR   418
NTDB id 1095974 ACMYQ0 RS06820 WP 415755301.1     EDIEEEP.LPPEDPGRKGI....   410
NTDB id 1177 A1552VC RS01165 WP 000422572.1       ........................   368
NTDB id 1203 PSJM300 01605 AFN76401.1             ...........DDPGRSFR....   381
NTDB id 1019 ACIAD RS04205 WP 004922051.1         ...........DGQRRNLK....   383
consensus                                                    *****
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X non conserved

X similar

X $\geq$ 50% conserved