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logo
NTDB id 1307 DR RS10055 WP 010888596.1      MTSGIDITDILRLAADKGASDIILAAGLPPQFKLSGDYDAQGFEPLSASDTRKLMYSMMNERQQRIFEEKRELDFSFALG   80
NTDB id 110951 RVX RS05300 WP 116306376.1   ...MARIDQFFHILHGLGGSDLHLSSGCTPMVRVHGTLQRLDQPALDDADLAAMLDFIIPPARRAEFAETGDLDFGYAVP   77
NTDB id 1086 NMV RS00245 WP 002218487.1     ....MQITDLLAFGAKNKASDLHLSSGISPMIRVHGDMRRINLPEMSAEEVGNMVTSVMNDHQRKIYQQNLEVDFSFELP   76
NTDB id 1116 NGFG RS10615 WP 003690200.1    ....MQITDLLAFGAKNKASDLHLSSGISPMIRVHGDMRRINLPEMSAEEVGNMVTSVMNDHQRKIYQQNLEVDFSFELP   76
NTDB id 1185 GTF74 RS11940 WP 000350195.1   ....MDIAELLEFSVKHNASDLHLSAGVPPMVRIDGEVRKLGVPAFTHSDVHRLIFEIMNDAQRSEYEEKLEVDFSFELP   76
NTDB id 1176 A1552VC RS01160 WP 000350195.1 ....MDIAELLEFSVKHNASDLHLSAGVPPMVRIDGEVRKLGVPAFTHSDVHRLIFEIMNDAQRSEYEEKLEVDFSFELP   76
NTDB id 1283 ACE17W RS02630 WP 010947729.1  ....MDIAELLAFSVKNKSSDLHISAGMPPMIRVDGDLRKINLPPLEHKDVIKIIYDIMNDRQRKEYEEFFETDFSFEIA   76
NTDB id 1259 GC085 RS10305 WP 010947729.1   ....MDIAELLAFSVKNKSSDLHISAGMPPMIRVDGDLRKINLPPLEHKDVIKIIYDIMNDRQRKEYEEFFETDFSFEIA   76
NTDB id 1202 PSJM300 01600 AFN76400.1       ....MDITELLAFSAKQGASDLHLSSGLPPMIRVDGDVRRINLPAMDHKQVHALIYDIMNDKQRKDFEEFLETDFSFEVP   76
NTDB id 1194 PAKAF RS01995 WP 003084552.1   ....MDITELLAFSAKQGASDLHLSAGLPPMIRVDGDVRRINLPPLEHKQVHALIYDIMNDKQRKDFEEFLETDFSFEVP   76
NTDB id 1020 ACIAD RS04210 WP 011182164.1   ....MDITELLAFSVQNRASDLHLSAGIPPMIRVDGEVRRINLPAMDHSEVHNLIYDIMNDKQRRDYEEKLETDFSFEVP   76
NTDB id 1060 ABD1 RS04380 WP 000355489.1    ....MDITELLAFSVKNGASDLHLSAGMPPMIRVDGEVRRINLPALEHKDVHRLVYDIMNDKQRRDYEEKLETDFSFEVP   76
NTDB id 1040 HON27 RS13360 WP 000355489.1   ....MDITELLAFSVKNGASDLHLSAGMPPMIRVDGEVRRINLPALEHKDVHRLVYDIMNDKQRRDYEEKLETDFSFEVP   76
NTDB id 1081 FDQ49 RS05335 WP 006581387.1   ....MDITELLAFSVKNGASDLHLSAGMPPMIRVDGEVRRINLPALEHKDVHRLVYDIMNDKQRRDYEEKLETDFSFEVP   76
consensus                                    **!***********  *!!*****!**!*****!************ ********************  ***!!*****

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NTDB id 1307 DR RS10055 WP 010888596.1      DKARFRVNAFMQRGNVGGVMRLIPTTVRTIAEMGLPQTVIDIANAPRGLVLVTGPTGSGKSTTLASMIDHINTTKKLHIL  160
NTDB id 110951 RVX RS05300 WP 116306376.1  GMARYRANFYRHERGVGAAFREIPSNIPSVEALGLPPLLRDLAMLPKGLVLLTGPTGSGKSTTLAAMIDHANTHRRDHII  157
NTDB id 1086 NMV RS00245 WP 002218487.1    NVARFRVNAFNIGRGPAAVFRTIPSTVLSLEELKAPSIFQKIAESPRGMVLVTGPTGSGKSTTLAAMINYINETQPAHIL  156
NTDB id 1116 NGFG RS10615 WP 003690200.1   NVARFRVNAFNTGRGPAAVFRTIPSTVLSLEELKAPSIFQKIAESPRGMVLVTGPTGSGKSTTLAAMINYINETQPAHIL  156
NTDB id 1185 GTF74 RS11940 WP 000350195.1  NVGRFRVNAFHQARGCSAVFRTIPTVIPTLEQLDAPEIFSKIANYEKGLVLVTGPTGSGKSTTLAAMVNYVNAHHNKHIL  156
NTDB id 1176 A1552VC RS01160 WP 000350195.1 NVGRFRVNAFHQARGCSAVFRTIPTVIPTLEQLDAPEIFSKIANYEKGLVLVTGPTGSGKSTTLAAMVNYVNAHHNKHIL  156
NTDB id 1283 ACE17W RS02630 WP 010947729.1 NLARFRVNAFNQSRGAAAVFRTIPSEILSMEDLGLPPIFKEMASFSRGLVLVTGPTGSGKSTTLAAVIDYINSSRYEHIL  156
NTDB id 1259 GC085 RS10305 WP 010947729.1  NLARFRVNAFNQSRGAAAVFRTIPSEILSMEDLGLPPIFKEMASFSRGLVLVTGPTGSGKSTTLAAVIDYINSSRYEHIL  156
NTDB id 1202 PSJM300 01600 AFN76400.1      GVARFRVNAFNQNRGAGAVFRTIPSKVLTMEDLGMGEVFRKITDVPRGLVLVTGPTGSGKSTTLAAMLDYLNNTKYHHIL  156
NTDB id 1194 PAKAF RS01995 WP 003084552.1  GVARFRVNAFNQNRGAGAVFRTIPSKVLTMEELGMGEVFKRVSDVPRGLVLVTGPTGSGKSTTLAAMLDYLNNTKYHHIL  156
NTDB id 1020 ACIAD RS04210 WP 011182164.1  NLARFRVNVFNQNRGAGAVFRTIPSQVLTLEELGLGKIFRDICDYPRGLVLVTGPTGSGKSTTLAAMLDYINNHRYDHIL  156
NTDB id 1060 ABD1 RS04380 WP 000355489.1   NVARFRVNAFNQNRGAGAVFRTIPSKVLTMEDLGLGQIFKDICDYPRGIVLVTGPTGSGKSTTLAAMLDYINENRYDHIL  156
NTDB id 1040 HON27 RS13360 WP 000355489.1  NVARFRVNAFNQNRGAGAVFRTIPSKVLTMEDLGLGQIFKDICDYPRGIVLVTGPTGSGKSTTLAAMLDYINENRYDHIL  156
NTDB id 1081 FDQ49 RS05335 WP 006581387.1  NVARFRVNAFNQNRGAGAVFRTIPSKVLTMEDLGLGQIFKDICDYPRGIVLVTGPTGSGKSTTLAAMLDYINENRYDHIL  156
consensus                                  ***!*!*!**** *******!*!!* ********** **********!*!!*!!!!!!!!!!!!!!******!   **  !!*

logo
NTDB id 1307 DR RS10055 WP 010888596.1      TIEDPIEFMHPNKASIINQREVGSDTLSFDDALRAALRQAPDVILVGEMRDYETIKAAVTAAETGHLVMGTLHTNSAPES  240
NTDB id 110951 RVX RS05300 WP 116306376.1  TIEDPVEFVHEPRGCLINQREVGRHTKSFAAALRGALREDPDIILVGEMRDLETISLALEAAETGHLVLSTVHTISAPKT  237
NTDB id 1086 NMV RS00245 WP 002218487.1    TIEDPIEFVHQSKKSLINQRELHQHTLSFANALRSALREDPDVILVGEMRDPETIGLALTAAETGHLVFGTLHTTGAAKT  236
NTDB id 1116 NGFG RS10615 WP 003690200.1   TIEDPIEFVHQSKKSLINQRELHQHTLSFANALSSALREDPDVILVGEMRDPETIGLALTAAETGHLVFGTLHTTGAAKT  236
NTDB id 1185 GTF74 RS11940 WP 000350195.1  TIEDPIEFVHSNNKCLINQREVHRDTHSFKNALRSALREDPDVILVGELRDQETISLALTAAETGHLVFGTLHTSSAAKT  236
NTDB id 1176 A1552VC RS01160 WP 000350195.1 TIEDPIEFVHSNNKCLINQREVHRDTHSFKNALRSALREDPDVILVGELRDQETISLALTAAETGHLVFGTLHTSSAAKT  236
NTDB id 1283 ACE17W RS02630 WP 010947729.1 TVEDPIEFVHQSKKCLVNQREVHKDTLSFNAALRSALREDPDIILVGELRDLETIRLAMTAAETGHLVFGTLHTNSATKT  236
NTDB id 1259 GC085 RS10305 WP 010947729.1  TVEDPIEFVHQSKKCLVNQREVHKDTLSFNAALRSALREDPDIILVGELRDLETIRLAMTAAETGHLVFGTLHTNSATKT  236
NTDB id 1202 PSJM300 01600 AFN76400.1      TIEDPIEFVHESKKCLVNQREVHRDTLGFSEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAKT  236
NTDB id 1194 PAKAF RS01995 WP 003084552.1  TIEDPIEFVHESKKCLVNQREVHRDTLGFSEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAKT  236
NTDB id 1020 ACIAD RS04210 WP 011182164.1  TVEDPIEFVHEPKKCLINQREVHRDTLGFNEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAKT  236
NTDB id 1060 ABD1 RS04380 WP 000355489.1   TVEDPIEFVHQSKKCLINQREVHRDTHGFNEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAKT  236
NTDB id 1040 HON27 RS13360 WP 000355489.1  TVEDPIEFVHQSKKCLINQREVHRDTHGFNEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAKT  236
NTDB id 1081 FDQ49 RS05335 WP 006581387.1  TVEDPIEFVHQSKKCLINQREVHRDTHGFNEALRSALREDPDIILVGEMRDLETIRLALTAAETGHLVFGTLHTTSAAKT  236
consensus                                  !*!!!*!!*! ******!!!!****!**!   !!**!!!**!!*!!!!!*!!*!!!**!**!!!!!!!!!**!*!!**!***
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NTDB id 1307 DR RS10055 WP 010888596.1        IDRIVDVFPEEQQEQIRVQLANNLVAVMTQQLLPRADGQGRVLAYEILLANPAVRALIREGKTFQITSVMQTGAREGMVT   320
NTDB id 110951 RVX RS05300 WP 116306376.1     IDRIVEVFPSEMQAQIRTSLSETLQAVISQVLFPRADRPGRVAALEIMLGVPAVRNLIRESKTYQLTSVLQTGKNAGMQT   317
NTDB id 1086 NMV RS00245 WP 002218487.1       VDRIVDVFPAGEKEMVRSMLSESLTAVISQNLLKTHDGNGRVASHEILIANPAVRNLIRENKITQINSVLQTGQASGMQT   316
NTDB id 1116 NGFG RS10615 WP 003690200.1      VDRIVDVFPAGEKEMVRSMLSESLTAVISQNLLKTHDGNGRVASHEILIANPAVRNLIRENKITQINSVLQTGQASGMQT   316
NTDB id 1185 GTF74 RS11940 WP 000350195.1     IDRIIDVFPGSDKDMVRSMLSESLRAVIAQKLLKR.VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAAHGMQT   315
NTDB id 1176 A1552VC RS01160 WP 000350195.1   IDRIIDVFPGSDKDMVRSMLSESLRAVIAQKLLKR.VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAAHGMQT   315
NTDB id 1283 ACE17W RS02630 WP 010947729.1    INRIIDVFPAEEKSMVRSMLSESLQAVVAQSLLKK.NKGGRVAALEIMMCTGAIRNLIREDKIAQMYSSIQTGQAKGMQT   315
NTDB id 1259 GC085 RS10305 WP 010947729.1     INRIIDVFPAEEKSMVRSMLSESLQAVVAQSLLKK.NKGGRVAALEIMMCTGAIRNLIREDKIAQMYSSIQTGQAKGMQT   315
NTDB id 1202 PSJM300 01600 AFN76400.1         IDRVVDVFPAEEKSMVRSMLSESLQAVISQTLLKK.VGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSSIQTGGSLGMQT   315
NTDB id 1194 PAKAF RS01995 WP 003084552.1     IDRVVDVFPAEEKAMVRSMLSESLQSVISQTLIKK.IGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSAIQTGGSLGMQT   315
NTDB id 1020 ACIAD RS04210 WP 011182164.1     IDRIIDVFPAEEKDMVRIMLSESLQAVISQTLLKK.EGGGRVAAHEIMIGIPAIRNLIREGKVAQMYSSIQTGATHGMTT   315
NTDB id 1060 ABD1 RS04380 WP 000355489.1      IDRVIDVFPAEEKDMVRAMLSESLQAVISQTLLKK.NGGGRVAAHEIMIGIPAIRNLIRENKVAQMYSAIQTGANHGMTT   315
NTDB id 1040 HON27 RS13360 WP 000355489.1     IDRVIDVFPAEEKDMVRAMLSESLQAVISQTLLKK.NGGGRVAAHEIMIGIPAIRNLIRENKVAQMYSAIQTGANHGMTT   315
NTDB id 1081 FDQ49 RS05335 WP 006581387.1     IDRVIDVFPAEEKDMVRAMLSESLQAVISQTLLKK.NGGGRVAAHEIMIGIPAIRNLIRENKVAQMYSAIQTGANHGMTT   315
consensus                                     **!***!!!*******!**!***!**!**!*!***  **!!!***!!*** *!*!*!!!! !**!**! *!!!*  !!*!
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NTDB id 1307 DR RS10055 WP 010888596.1        MDAYLAGLFRRHLITYDKGLERAIDPKEFARLANDASAGLGGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAAPATPA   400
NTDB id 110951 RVX RS05300 WP 116306376.1     LDDDILRLLGAGLIHPRDAAVHVADKPRFKPYL....AGVGPTAMGRAA..........SASDATDDDF...........   372
NTDB id 1086 NMV RS00245 WP 002218487.1       MDQSLQSLVRQGLIAPEVARRRAQNS..ESMSF.................................................   347
NTDB id 1116 NGFG RS10615 WP 003690200.1      MDQSLQSLVRQGLIAPEAARRRAQNS..ESMSF.................................................   347
NTDB id 1185 GTF74 RS11940 WP 000350195.1     MEQNAKQLIARGVVDAQEVQSKIELD..LKAF...................................................   345
NTDB id 1176 A1552VC RS01160 WP 000350195.1   MEQNAKQLIARGVVDAQEVQSKIELD..LKAF...................................................   345
NTDB id 1283 ACE17W RS02630 WP 010947729.1    LDQHLTELVNKNIISKHTAHEAALNK...SLF...................................................   344
NTDB id 1259 GC085 RS10305 WP 010947729.1     LDQHLTELVNKNIISKHTAHEAALNK...SLF...................................................   344
NTDB id 1202 PSJM300 01600 AFN76400.1         LDSCLKGLLAKGLISRDSAKEKAKQP...ENF...................................................   344
NTDB id 1194 PAKAF RS01995 WP 003084552.1     LDMCLKGLVAKGLISRENAREKAKIP...ENF...................................................   344
NTDB id 1020 ACIAD RS04210 WP 011182164.1     LDQSLKQLLAKGIINLPTARAVAKSP...ESF...................................................   344
NTDB id 1060 ABD1 RS04380 WP 000355489.1      LDQSLKGLVARGVISPQTARTAAKQP...ESFL..................................................   345
NTDB id 1040 HON27 RS13360 WP 000355489.1     LDQSLKGLVARGVISPQTARTAAKQP...ESFL..................................................   345
NTDB id 1081 FDQ49 RS05335 WP 006581387.1     LDQSLKGLVSRGVISPQTARTAAKQP...ESFL..................................................   345
consensus                                     *** ** !*******    **  * *       *
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logo
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NTDB id 1307 DR RS10055 WP 010888596.1        APAAGRSEPGRAGGFGRR   418
NTDB id 110951 RVX RS05300 WP 116306376.1     ..................   372
NTDB id 1086 NMV RS00245 WP 002218487.1       ..................   347
NTDB id 1116 NGFG RS10615 WP 003690200.1      ..................   347
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 1202 PSJM300 01600 AFN76400.1         ..................   344
NTDB id 1194 PAKAF RS01995 WP 003084552.1     ..................   344
NTDB id 1020 ACIAD RS04210 WP 011182164.1     ..................   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
consensus
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X non conserved
X similar
X $\geq$ 50% conserved