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logo
NTDB id 85 BSU 00860 NP 387967.1              ..MMFGRFTERAQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGIAAKALQALGLGSEKIQKEVESLIG........R   70
NTDB id 610 V4T04 RS10165 WP 012897346.1      MKFENIKYTPTLDRILEKAEEYAHQYQYGTIESAHLLAAMATTSGSIAYSLLAGMNVDSSDLLIDLE........DLSSH   72
NTDB id 291172 DEH83 RS03485 WP 255777069.1  .....MKHSKALTESIESAQLLASHFETDYLESWQLLIAMANNPYSVAGSVLNDYPLEIDDFE..KA...AFHITGKVYQ   70
NTDB id 287 SP RS11210 WP 001109712.1        .....MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE..EV...ALELTETDYS   70
NTDB id 290 KZH43 RS10025 WP 001109677.1     .....MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE..EV...ALELTETDYS   70
NTDB id 289 SPD RS10700 WP 001109677.1       .....MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE..EV...ALELTETDYS   70
NTDB id 377 SMU RS09275 WP 002262344.1       ....MTDYSLKMQEVFRLAQFEAARFESSYLESWHVLLAMVEIDSSVAGLSFAEFEADVRFEDYQAAAILAIGKKPKSSS   76
NTDB id 329 STU RS10020 WP 011225298.1       ....MTIYSRKMQAIFHRAQLEAERFESPFLETWHVLLAMVEVPGSVAYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLS   76
NTDB id 297 STER RS00545 WP 011680614.1      ....MTIYSRKMQAIFHRAQLEAERFESPFLETWHVLLAMVEVTGSVAYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLS   76
consensus                                       * *** ****** !** !  *   *******!***   * **!* *****  *** **        ***    * *
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logo
NTDB id 85 BSU 00860 NP 387967.1              GQEMSQTIHYTPRAKKVIELSMDEARKLGHSYVGTEHILLGLIREGEGVAARVLNNLGVSLNKARQQVLQL.........   141
NTDB id 610 V4T04 RS10165 WP 012897346.1      VKVKRSTLRFSPRAEEVMTAASFLAIHNNSEAVGTEHLLYALLQVEDGFGLQLLKL.........QKINIVSLRKELEKR   143
NTDB id 291172 DEH83 RS03485 WP 255777069.1  QEGAFTIWPFSYRMKVLFLTAEQIAEAVHAKNLGTEHVLLAMLFDRGSLAARILEFIGFSYEDKEGALRMTDLRKNLEHK   150
NTDB id 287 SP RS11210 WP 001109712.1        QDETFTELPFSRRLQVLFDEAEYVASVVHAKVLGTEHVLYAILHDSNALATRILERAGFSYEDKKDQVKIAALRRNLEER   150
NTDB id 290 KZH43 RS10025 WP 001109677.1     QDETFTELPFSRRLQILFDEAEYVASVVHAKVLGTEHVLYAILHDGNALATRILERAGFSYEDKKDQVKIAALRRNLEER   150
NTDB id 289 SPD RS10700 WP 001109677.1       QDETFTELPFSRRLQILFDEAEYVASVVHAKVLGTEHVLYAILHDGNALATRILERAGFSYEDKKDQVKIAALRRNLEER   150
NTDB id 377 SMU RS09275 WP 002262344.1       N..DIMLLEQSHALKRTLAEAAAISQVTHAKEVGTEHVLFAMLLNPNLLATRILELVGFHAKDDGESIRLLDLRKVIERY   154
NTDB id 329 STU RS10020 WP 011225298.1       ESDIIDLRAQSPALEAMLQEAQGIASVTGAVEVGSEHVLMAFLLHKDLMVCRLLEVAGFQYKDDSDKPRIIDLRRSLERN   156
NTDB id 297 STER RS00545 WP 011680614.1      ESDIIDLRAQSPALEAMLQEAQEIASVTGAVEVGSEHVLMAFLLHKDLMVCRLLEVAGFQYKDDSDKPRIIDLRRSLERN   156
consensus                                    * *  * * ** **  ** **  ******** *!*!!*!**** *  *** **!* ***** *  * **** *** ** *
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logo
NTDB id 85 BSU 00860 NP 387967.1              LGSNETGS...................SAAGTNSNANTPTLDSLARDLTAIAKEDSLDPVIGRSKEIQRVIEVLSRRTKN   202
NTDB id 610 V4T04 RS10165 WP 012897346.1      TGLKVPESKKAVTPMSKRKM.......AKG.VAENSTTPTLDSVSSDLTEEARLGKLDPMIGREAEIDRLIHILSRRTKN   215
NTDB id 291172 DEH83 RS03485 WP 255777069.1  ASWNKEDLKA.IRHLNKNAA.ATRQTMANMMGMPPATSGGLEDYTRDLTEMAHMGLLEPVIGRDKEISRMLQILSRKTKN   228
NTDB id 287 SP RS11210 WP 001109712.1        AGWTREDLKA.LRQRHRTVA.DKQNSMANMMGMPQTPSGGLEDYTHDLTEQARSGKLEPVIGRDKEISRMIQILSRKTKN   228
NTDB id 290 KZH43 RS10025 WP 001109677.1     AGWTREDLKA.LRQRHRTVA.DKQNSMANMMGMPQTPSGGLEDYTHDLTEQARSGKLEPVIGRDKEISRMIQILSRKTKN   228
NTDB id 289 SPD RS10700 WP 001109677.1       AGWTREDLKA.LRQRHRTVA.DKQNSMANMMGMPQTPSGGLEDYTHDLTEQARSGKLEPVIGRDKEISRMIQILSRKTKN   228
NTDB id 377 SMU RS09275 WP 002262344.1       AGFSKEDIKA.IFEMRKPKKVKNSSSFSDLMKPP.SVTGELADFTRDLTELARQGRLEPVIGREQEISRMVQILSRKTKN   232
NTDB id 329 STU RS10020 WP 011225298.1       AGLSKQDLKA.IHDLRKPKKSKASANFANMMQPPQSSTGELADYTKDLTALAESGNLDPIIGRDEEISRMIQVLSRKTKN   235
NTDB id 297 STER RS00545 WP 011680614.1      AGLSKQDLKA.IHDLRKPKKSKASANFANMMQPPQSSTGELADYTKDLTALAESGNLDPVIGRDEEISRMIQVLSRKTKN   235
consensus                                    ********** *  * *        * ****  *** ** !*****!!!**!****!*!*!!!**!!*!****!!!*!!!
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logo
NTDB id 85 BSU 00860 NP 387967.1              NPVLIGEPGVGKTAIAEGLAQQIINNEVPEILRDKRVMTLDMGTVVAGTKYRGEFEDRLKKVMDEIRQAGNIILFIDELH   282
NTDB id 610 V4T04 RS10165 WP 012897346.1      NPVLVGEPGVGKSAIIEGLAQRIVNGQVPIGLMNSRIMALNMATVVAGTKFRGEFEDRLTAIVEEVSSDPDVIIFIDELH   295
NTDB id 291172 DEH83 RS03485 WP 255777069.1  NPVLVGDAGVGKTALALGLAQRVAAGQVPNELAKMRVLELDLMNVVAGTRFRGDFEERMNNIINDIEADGHVILFIDELH   308
NTDB id 287 SP RS11210 WP 001109712.1        NPVLVGDAGVGKTALALGLAQRIASGDVPAEMAKMRVLELDLMNVVAGTRFRGDFEERMNNIIKDIEEDGQVILFIDELH   308
NTDB id 290 KZH43 RS10025 WP 001109677.1     NPVLVGDAGVGKTALALGLAQRIASGDVPAEMAKMRVLELDLMNVVAGTRFRGDFEERMNNIIKDIEEDGQVILFIDELH   308
NTDB id 289 SPD RS10700 WP 001109677.1       NPVLVGDAGVGKTALALGLAQRIASGDVPAEMAKMRVLELDLMNVVAGTRFRGDFEERMNNIIKDIEEDGQVILFIDELH   308
NTDB id 377 SMU RS09275 WP 002262344.1       NPVLVGDAGVGKTALAYGLAQRIVDSAVPFELADMKVLELDMMSVVAGTRFRGDFEERMNQIIADIEADGHIVLFIDELH   312
NTDB id 329 STU RS10020 WP 011225298.1       NPVLVGEAGVGKTALALGLAQRIASGEVPFELADMRILELDMMSVVAGTRFRGDFEERMNQIIDEIEADGKIILFIDELH   315
NTDB id 297 STER RS00545 WP 011680614.1      NPVLVGEAGVGKTALALGLAQRIASGEVPFELADMRILELDMMSVVAGTRFRGDFEERMNQIIDEIEADGKIILFIDELH   315
consensus                                    !!!!*!**!!!!*!***!!!!******!! *** *****!****!!!!!**!!*!!*!***** *** ** ***!!!!!!
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logo
NTDB id 85 BSU 00860 NP 387967.1                TLIGAGG.AEGAIDASNILKPSLARGELQCIGATTLDEYRKYIEKDAALERRFQPIQVDQPSVDESIQILQGLRDRYEAH   361
NTDB id 610 V4T04 RS10165 WP 012897346.1        TIIGAGGGMDSVNDAANILKPALARGDFQMVGATTYHEYQKYIEKDEALERRLARINVDEPSPDEAIAILQGLREKFEDY   375
NTDB id 291172 DEH83 RS03485 WP 255777069.1     TIMGSGSGIDSTLDAANILKPALARGTLRTVGATTQEEYQKYIEKDAALSRRFAKVAIEEPTVADSIAILQGLKKTYEDH   388
NTDB id 287 SP RS11210 WP 001109712.1           TIMGSGSGIDSTLDAANILKPALARGTLRTVGATTQEEYQKHIEKDAALSRRFAKVTIEEPSVADSMTILQGLKATYEKH   388
NTDB id 290 KZH43 RS10025 WP 001109677.1        TIMGSGSGIDSTLDAANILKPALARGTLRTVGATTQEEYQKHIEKDAALSRRFAKVTIEEPSVADSMTILQGLKATYEKH   388
NTDB id 289 SPD RS10700 WP 001109677.1          TIMGSGSGIDSTLDAANILKPALARGTLRTVGATTQEEYQKHIEKDAALSRRFAKVTIEEPSVADSMTILQGLKATYEKH   388
NTDB id 377 SMU RS09275 WP 002262344.1          TIMGSGSGIDSTLDAANILKPALARGSLRTIGATTQEEYQKHIEKDAALSRRFAKVTIEEPSEEEAYQILLGLKKSYETY   392
NTDB id 329 STU RS10020 WP 011225298.1          TIIGSGSGIDSTLDAANILKPALARGTLHMVGATTQAEYQKHIEKDAALSRRFAKITIEEPSVSEAIDILNGLRSSYEDY   395
NTDB id 297 STER RS00545 WP 011680614.1         TIIGSGSGIDSTLDAANILKPALARGTLHMVGATTQAEYQKHIEKDAALSRRFAKITIEEPSVSEAIDILNGLRSSYEDY   395
consensus                                       !**!*!*******!!*!!!!!*!!!!*****!!!!**!!*!*!!!!*!!*!!********!** *** !!*!!* **! *

logo
NTDB id 85 BSU 00860 NP 387967.1                HRVSITDDAIEAAVKLSDRYISDRFLPDKAIDLIDEAGSKVRLRSFTTPPNLKELEQKLDEVRKEKDAAVQSQEFEKAAS   441
NTDB id 610 V4T04 RS10165 WP 012897346.1        HQVKFTDQAIKSAVMLSVRYMTSRKLPDKAIDLLDEAAAAVKISVKNQQTKRLDLEKELAEAQEELSEAVIKLDIKASRT   455
NTDB id 291172 DEH83 RS03485 WP 255777069.1     HKVVITDSAIETAVKYAHRYLTSKHLPDSAIDLLDEAAATVQNKGPQNH.VKVEL....SVAD....EALMAGNWNKVST   459
NTDB id 287 SP RS11210 WP 001109712.1           HRVQITDEAVETAVKMAHRYLTSRHLPDSAIDLLDEAAATVQNKAKHVKADDSDL....SPAD....KALMDGKWKQAAQ   460
NTDB id 290 KZH43 RS10025 WP 001109677.1        HRVQITDEAVETAVKMAHRYLTSRHLPDSAIDLLDEAAATVQNKAKHVKADDSDL....SPAD....KALMDGKWKQAAQ   460
NTDB id 289 SPD RS10700 WP 001109677.1          HRVQITDEAVETAVKMAHRYLTSRHLPDSAIDLLDEAAATVQNKAKHVKADDSDL....SPAD....KALMDGKWKQAAQ   460
NTDB id 377 SMU RS09275 WP 002262344.1          HHVIISDAAVMAAVKGAHRYLSGKNLPDSAIDLLDEASATVQGLVKK..KAPAYL....TDLD....HALITNDYQLAKR   462
NTDB id 329 STU RS10020 WP 011225298.1          HRVTITDAAVETAVKAAHRYLTSKNLPDSAIDLLDEASATVQVRIKK..EAKREI....TPLD....EALISGDIGAAVK   465
NTDB id 297 STER RS00545 WP 011680614.1         HRVTITDAAVETAVKAAHRYLTSKNLPDSAIDLLDEASATVQVRIKK..EDKREI....TPLD....EALISGDIGAAVK   465
consensus                                       !*! **! !***!!*****!**** !!!*!!!!*!!!***!* * *       **     ****      !** ***  *

logo
NTDB id 85 BSU 00860 NP 387967.1                LRDTEQRLREQVEDTKKSWKEKQGQENSEVTVDDIAMVVSSWTGVPVSKIAQTETDKLLNMENILHSRVIGQDEAVVAVA   521
NTDB id 610 V4T04 RS10165 WP 012897346.1        KEKAVEKIADKIYKF.....SVKEDKRQEVTDQAVVAVASTLTGVPITQMTKSESDRLINLEKELHKRVVGQEEAISAVS   530
NTDB id 291172 DEH83 RS03485 WP 255777069.1     LLEKE.........S.....QPIVY.KLRVKDEDVLATLSSLSGIPVQKLTQTDAKKYLNLEKELHKRVIGQNEAISAIS   524
NTDB id 287 SP RS11210 WP 001109712.1           LIAKE.........E.....EVPVY.KDLVTESDILTTLSRLSGIPVQKLTQTDAKKYLNLEAELHKRVIGQDQAVSSIS   525
NTDB id 290 KZH43 RS10025 WP 001109677.1        LIAKE.........E.....EVPVY.KDLVTESDILTTLSRLSGIPVQKLTQTDAKKYLNLEAELHKRVIGQDQAVSSIS   525
NTDB id 289 SPD RS10700 WP 001109677.1          LIAKE.........E.....EVPVY.KDLVTESDILTTLSRLSGIPVQKLTQTDAKKYLNLEAELHKRVIGQDQAVSSIS   525
NTDB id 377 SMU RS09275 WP 002262344.1          LLKKE.........K.....KPLLKKRAEVKEEDILATLSKLSGIPVKKLTQADSKKYLNLERELHKRVIGQDDAVSSIS   528
NTDB id 329 STU RS10020 WP 011225298.1          QYKAN.........Q.....KAKFPKPALVDADQIMQTLSRLSGIPVEKMTQTDSKRYLNLESELHKRVIGQDEAVSAIS   531
NTDB id 297 STER RS00545 WP 011680614.1         QYKAN.........Q.....KAKFPKPALVDADQIMQTLSRLSGIPVEKMTQTDSKRYLNLESELHKRVIGQDEAVSAIS   531
consensus                                       ** **                 *  * *!****** **!***!*!* ***********!*! *!!*!!*!!**!*****

logo
NTDB id 85 BSU 00860 NP 387967.1                KAVRRARAGLKDPKRPIGSFIFLGPTGVGKTELARALAESIFGDEESMIRIDMSEYMEKHSTSRLVGSPPGYVGYDEGGQ   601
NTDB id 610 V4T04 RS10165 WP 012897346.1        RAIRRARSGVADSRRPMGSFMFLGPTGVGKTELAKALADSVFGSEDNMIRVDMSEFMEKHSTSRLIGAPPGYVGYDEGGQ   610
NTDB id 291172 DEH83 RS03485 WP 255777069.1     RAIRRNQSGIRTSKRPIGSFMFLGPTGVGKTELAKALAEVLFDDETALIRFDMSEYMEKFAASRLNGAPPGYVGYEEGGE   604
NTDB id 287 SP RS11210 WP 001109712.1           RAIRRNQSGIRSHKRPIGSFMFLGPTGVGKTELAKALAEVLFDDESALIRFDMSEYMEKFAASRLNGAPPGYVGYEEGGE   605
NTDB id 290 KZH43 RS10025 WP 001109677.1        RAIRRNQSGIRSHKRPIGSFMFLGPTGVGKTELAKALAEVLFDDESALIRFDMSEYMEKFAASRLNGAPPGYVGYEEGGE   605
NTDB id 289 SPD RS10700 WP 001109677.1          RAIRRNQSGIRSHKRPIGSFMFLGPTGVGKTELAKALAEVLFDDESALIRFDMSEYMEKFAASRLNGAPPGYVGYEEGGE   605
NTDB id 377 SMU RS09275 WP 002262344.1          RAIRRNQSGIRVGKRPIGSFMFLGPTGVGKTELAKALAEVLFDDESALIRFDMSEYMEKFAASRLNGAPPGYVGYEEGGE   608
NTDB id 329 STU RS10020 WP 011225298.1          RAIRRNQSGIRTGKRPIGSFMFLGPTGVGKTELAKALAEVLFDDESALLRFDMSEYMEKFAASRLNGAPPGYVGYDEGGE   611
NTDB id 297 STER RS00545 WP 011680614.1         RAIRRNQSGIRTGKRPIGSFMFLGPTGVGKTELAKALAEVLFDDESALLRFDMSEYMEKFAASRLNGAPPGYVGYDEGGE   611
consensus                                       *!*!!***!*** *!!*!!!*!!!!!!!!!!!!!!*!!!***!**!****!*!!!!*!!!***!!!!*!*!!!!!!!*!!!*
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logo

NTDB id 85 BSU 00860 NP 387967.1          LTEKVRRKPYSVVLLDEIEKAHPDVFNILLQVLEDGRLTDSKGRTVDFRNTILIMTSNVGASELKRNKYVGFNVQDETQN   681
NTDB id 610 V4T04 RS10165 WP 012897346.1  LTERVRNKPYSVVLLDEVEKAHPDVFNIMLQILDDGFVTDTKGRKVDFRNTIIIMTSNLGATALRDDKTVGFGAKNITAD   690
NTDB id 291172 DEH83 RS03485 WP 255777069.1 LTEKVRNRPYSVLLFDEVEKAHPDIFNVLLQVLDDGVLTDSKGRKVDFSNTIIIMTSNLGATSLRDDKTVGFGARDVRFD   684
NTDB id 287 SP RS11210 WP 001109712.1     LTEKVRNKPYSVLLFDEVEKAHPDIFNVLLQVLDDGVLTDSKGRKVDFSNTIIIMTSNLGATALRDDKTVGFGAKDIRFD   685
NTDB id 290 KZH43 RS10025 WP 001109677.1  LTEKVRNKPYSVLLFDEVEKAHPDIFNVLLQVLDDGVLTDSKGRKVDFSNTIIIMTSNLGATALRDDKTVGFGAKDIRFD   685
NTDB id 289 SPD RS10700 WP 001109677.1    LTEKVRNKPYSVLLFDEVEKAHPDIFNVLLQVLDDGVLTDSKGRKVDFSNTIIIMTSNLGATALRDDKTVGFGAKDIRFD   685
NTDB id 377 SMU RS09275 WP 002262344.1    LTEKVRNKPYSVLLFDEVEKAHPDIFNILLQVLDDGVLTDSRGRKVDFSNTIIIMTSNLGATALRDDKTVGFGAKSFSQD   688
NTDB id 329 STU RS10020 WP 011225298.1    LTEKVRNKPYSVLLFDEIEKAHPDIFNVLLQVLDDGVLTDSRGRKVDFSNTIIIMTSNLGATALRDDKTVGFGAQTISHN   691
NTDB id 297 STER RS00545 WP 011680614.1   LTEKVRNKPYSVLLFDEIEKAHPDIFNILLQVLDDGVLTDSRGRKVDFSNTIIIMTSNLGATALRDDKTVGFGAQTISHN   691
consensus                                 !!!*!!**!!!!*!*!!*!!!!!!*!!**!!*!*!!**!!**!!*!!!*!!!*!!!!!*!!**!***!*!!!******  *

logo

NTDB id 85 BSU 00860 NP 387967.1          HKDMKDKVMGELKRARRPEFINRIDEIIVFHSLEKKHLTEIVSLMSDQLTKRLKEQDLSIELTDAAKAKVAEEGVDLEYG   761
NTDB id 610 V4T04 RS10165 WP 012897346.1  YSAMKSRILEELKRHYRPEFLNRIDENIVFHSLESQEIEQIVKIMSKSLIKRLAEQDIHVKLTPSAVKLIAEVGFDPEYG   770
NTDB id 291172 DEH83 RS03485 WP 255777069.1 HENMEKRMMEELKKAYRPEFINRIDEKVVFHSLTSEDMQEVVKIMVQPLISSLAEKGIKLKFQPSALKLLAQEGYDPEMG   764
NTDB id 287 SP RS11210 WP 001109712.1     QENMEKRMFEELKKAYRPEFINRIDEKVVFHSLSSDHMQEVVKIMVKPLVASLTEKGIDLKLQASALKLLANQGYDPEMG   765
NTDB id 290 KZH43 RS10025 WP 001109677.1  QENMEKRMFEELKKAYRPEFINRIDEKVVFHSLSSDHMQEVVKIMVKPLVASLAEKGIDLKLQASALKLLANQGYDPEMG   765
NTDB id 289 SPD RS10700 WP 001109677.1    QENMEKRMFEELKKAYRPEFINRIDEKVVFHSLSSDHMQEVVKIMVKPLVASLAEKGIDLKLQASALKLLANQGYDPEMG   765
NTDB id 377 SMU RS09275 WP 002262344.1    YKAMESRILEELKKVYRPEFINRIDEKVVFHNLGQEDIRHIVKIMVAPLIAHLADQGITLKFQPSALKHLALAGYDAEMG   768
NTDB id 329 STU RS10020 WP 011225298.1    HQAMQARIMEELKKSYRPEFINRIDEKVVFHSLEEEQLHDIVKIMVKPLISALADKGISLKFQPAALKHLAKDGYDIEMG   771
NTDB id 297 STER RS00545 WP 011680614.1   HQAMQARIMEELKKSYRPEFINRIDEKVVFHSLEEEQLHDIVKIMVKPLISALADKGISLKFQPAALKHLAKDGYDIEMG   771
consensus                                    !* ****!!!***!!!!*!!!!!**!!!*!  ** * **!**!***!*   !***** ******!****!   !*!*!*!

logo

NTDB id 85 BSU 00860 NP 387967.1          ARPLRRAIQKHVEDRLSEELLRGNIHKGQHIVLDVEDGEFVVKTTAKTN   810
NTDB id 610 V4T04 RS10165 WP 012897346.1  ARPLRKALQKEVEDLLSEQLLSGEIKAGNHVSIGASNKKIKIAQIV...   816
NTDB id 291172 DEH83 RS03485 WP 255777069.1 ARPLRRILQTRVEDHLSELLLAGELKTGQSLKVGVKAGKLKFEVL....   809
NTDB id 287 SP RS11210 WP 001109712.1     ARPLRRTLQTEVEDKLAELLLKGDLVAGSTLKIGVKAGQLKFDIA....   810
NTDB id 290 KZH43 RS10025 WP 001109677.1  ARPLRRTLQTEVEDKLAELLLKGDLVAGSILKIGVKAGQLKFDIA....   810
NTDB id 289 SPD RS10700 WP 001109677.1    ARPLRRTLQTEVEDKLAELLLKGDLVAGSILKIGVKAGQLKFDIA....   810
NTDB id 377 SMU RS09275 WP 002262344.1    ARPLRRTLQTEVEDKLAELILSGKLASGQTLKIGIQKEKLKFDIV....   813
NTDB id 329 STU RS10020 WP 011225298.1    ARPLRRTIQTQVEDKLSELLLGGQVVSGQTLKIGCSKDKLTFTVV....   816
NTDB id 297 STER RS00545 WP 011680614.1   ARPLRRTIQTQVEDKLSELLLGGQVVSGQTLKIGCSKDKLTFTVV....   816
consensus                                 !!!!!***!**!!!*!*!**!  !***  !*******  ********
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X non conserved

X similar

X ≥ 50% conserved