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logo
                                                  MSXLIIGDRRLGAILLDRGYLIDNDLQRALERHREVGGRLADVLYDSGMYSERRIABIIEERFGIPLVNLHRVEIPPKXK
NTDB id 1306 DR RS10060 WP 027479822.1            .MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL   79
NTDB id 1028 TT RS08215 WP 011173991.1            MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK   80
NTDB id 1168 A1552VC RS11070 WP 000957200.1       ..................................................................................    0
NTDB id 1401 DSB67 RS12665 WP 010643259.1         ..................................................................................    0
NTDB id 1293 VP RS12240 WP 005479695.1            ..................................................................................    0
NTDB id 1251 GCO85 RS07725 WP 011946523.1         ..................................................................................    0
NTDB id 1110 NGFG RS09235 WP 003689817.1          ..................................................................................    0
NTDB id 83449 JLC71 RS00195 WP 200916685.1        ..................................................................................    0
NTDB id 1058 ABD1 RS01615 WP 001274986.1          ..................................................................................    0
NTDB id 1018 ACIAD RS01685 WP 004920473.1         ..................................................................................    0
consensus
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logo
                                                  RSIRPQTALNLQAFPFALEAGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA
NTDB id 1306 DR RS10060 WP 027479822.1            RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA  157
NTDB id 1028 TT RS08215 WP 011173991.1            ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P  158
NTDB id 1168 A1552VC RS11070 WP 000957200.1       ..................................................................................    0
NTDB id 1401 DSB67 RS12665 WP 010643259.1         ..................................................................................    0
NTDB id 1293 VP RS12240 WP 005479695.1            ..................................................................................    0
NTDB id 1251 GCO85 RS07725 WP 011946523.1         ..................................................................................    0
NTDB id 1110 NGFG RS09235 WP 003689817.1          ..................................................................................    0
NTDB id 83449 JLC71 RS00195 WP 200916685.1        ..................................................................................    0
NTDB id 1058 ABD1 RS01615 WP 001274986.1          ..................................................................................    0
NTDB id 1018 ACIAD RS01685 WP 004920473.1         ..................................................................................    0
consensus
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logo
                                                  SDAGRTGGKLGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL
NTDB id 1306 DR RS10060 WP 027479822.1            SDAGRTGGKLGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL  237
NTDB id 1028 TT RS08215 WP 011173991.1            SGEGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA  237
NTDB id 1168 A1552VC RS11070 WP 000957200.1       ..................................................................................    0
NTDB id 1401 DSB67 RS12665 WP 010643259.1         ..................................................................................    0
NTDB id 1293 VP RS12240 WP 005479695.1            ..................................................................................    0
NTDB id 1251 GCO85 RS07725 WP 011946523.1         ..................................................................................    0
NTDB id 1110 NGFG RS09235 WP 003689817.1          ..................................................................................    0
NTDB id 83449 JLC71 RS00195 WP 200916685.1        ..................................................................................    0
NTDB id 1058 ABD1 RS01615 WP 001274986.1          ..................................................................................    0
NTDB id 1018 ACIAD RS01685 WP 004920473.1         ..................................................................................    0
consensus
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logo
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NTDB id 1306 DR RS10060 WP 027479822.1      GSMLRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQRGRLGEQLVQEGTLSR   317
NTDB id 1028 TT RS08215 WP 011173991.1      LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYLALPQAWEELFRRAYPQKNRLGEVLVQEGKLSR   317
NTDB id 1168 A1552VC RS11070 WP 000957200.1 ................................................................................     0
NTDB id 1401 DSB67 RS12665 WP 010643259.1   ................................................................................     0
NTDB id 1293 VP RS12240 WP 005479695.1      ................................................................................     0
NTDB id 1251 GC085 RS07725 WP 011946523.1   ................................................................................     0
NTDB id 1110 NGFG RS09235 WP 003689817.1    ................................................................................     0
NTDB id 83449 JLC71 RS00195 WP 200916685.1  ................................................................................     0
NTDB id 1058 ABD1 RS01615 WP 001274986.1    ................................................................................     0
NTDB id 1018 ACIAD RS01685 WP 004920473.1   ................................................................................     0
consensus
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logo
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NTDB id 1306 DR RS10060 WP 027479822.1      DQLREALQVQAREGKVKPLGEVITELGFASPDEVDSALQKQNVGGGRLEDTLVQSGKLSPEMLARSLAAQLGYEFLDPIQ   397
NTDB id 1028 TT RS08215 WP 011173991.1      EALKEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEE   397
NTDB id 1168 A1552VC RS11070 WP 000957200.1 ..............MLTNLVAILRQAELISATQEQAVVTQVSASGTSVPEALLELSIFHAQELTEQLSHIFGLPETDLSR    66
NTDB id 1401 DSB67 RS12665 WP 010643259.1   ..............MLTNLSTVLRQAGILTFSQEESLLEQVKASGISMPEALLSSGFFQSHELTEHLSSVFGLSCTSLNQ    66
NTDB id 1293 VP RS12240 WP 005479695.1      ..............MHSNLSTILRQKGLLTFSQEESLIEQVKASGISMPEALLSSGFFTSSELAEHLSSIFGLSQPELSQ    66
NTDB id 1251 GC085 RS07725 WP 011946523.1   ......MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVKNKILSAEQIALTAAQNFGVPMLDINC    74
NTDB id 1110 NGFG RS09235 WP 003689817.1    ..............MSVGLLRILVQNQVVTVEQAEHYYNE.SQAGKEVLPMLFSDGVISPKSLAALIARVFSYSILDLRH    65
NTDB id 83449 JLC71 RS00195 WP 200916685.1  .........MSTTPSISGLARLLVQHGRLSDTDAETLQSTAATNKAPFIAQLLQSGKMNAKEVAEFSAQAFGYPLLDLEQ    71
NTDB id 1058 ABD1 RS01615 WP 001274986.1    ......MSALHTSPKFTGFFRRLVEEKHVSAATMQTALDAAKRAKQDTVAYLIEEVHLSPSLLAETISAEFAEPYFDLDV    74
NTDB id 1018 ACIAD RS01685 WP 004920473.1   ......MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDIVPYLIDNFSISPLTIAEIISLEFGEPLLDLGV    74
consensus                                         *       *****  **** **** ** ****   *  *  *** ** ** ***  *** *   ** * ***
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logo
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NTDB id 1306 DR RS10060 WP 027479822.1      NPPDPKVALMIPEATARRYVVVPVRLQGNSLVVAMKDPRNVFALDDLKLITG.....KEILPAVMAEKDIIRLIERYFGE   472
NTDB id 1028 TT RS08215 WP 011173991.1      DPPDPGAPLLLPEDLCRRYGVFPHRLEGNRLVLLMKDPRNILALDDVRLALKRKGLNYEVAPAVATEAAITKLIERFYGK   477
NTDB id 1168 A1552VC RS11070 WP 000957200.1 YDYANLCQQLGLRELITRYDALPIAKQGNLLLLAVSDPTLLQAEEEFRFATG.....LQVELALADHRALQAAIRRLYGR   141
NTDB id 1401 DSB67 RS12665 WP 010643259.1   YEYASLCQTLGLRDLITRHNALPLNRTSSTLILAVADPTNLQAEDDFRFATG.....LQVELVLADFRELTAAIRRLYGR   141
NTDB id 1293 VP RS12240 WP 005479695.1      YEYASLCQQLGLRELITRHNALPLHRTPSTLLLAVADPTNQQAEDDFRFATG.....LQVELVLADFRELSTAIRRLYGR   141
NTDB id 1251 GC085 RS07725 WP 011946523.1   IDVGTIPVNLVNEKLIKRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTG.....LNTHAIVVETDKLSALIDNLLTA   149
NTDB id 1110 NGFG RS09235 WP 003689817.1    YPRHRVLMGVLTEEQMVEFHCVPVFRRGDKVFFAVSDPTQMPQIQKTVSAAG.....IAVELVIVEDDQLAGLLDWVGSR   140
NTDB id 83449 JLC71 RS00195 WP 200916685.1  IDTSYIPQGVLDTKVMQAQRIVPLFKRGTKLFIGISDITNLQAAEEVRFQTG.....AQIEPIVVEDNKLVKLLDHLLEA   146
NTDB id 1058 ABD1 RS01615 WP 001274986.1    YDTSQIPKDLVDQKLILKHRILPLIQRGQILYVATSNPSNIEAIDAIRFNSK.....LLVEPVIVEHHKLEKVLGQHFAE   149
NTDB id 1018 ACIAD RS01685 WP 004920473.1   FDPALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNIEAMDAIRFNSK.....LKVEPIIVEHDKLERLLSEHFVE   149
consensus                                   ** * *   ** ****  ** **!* ***  * ********* **********     *********   * *********
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logo

NTDB id 1306 DR RS10060 WP 027479822.1        KGFEKL......NK........ELAERNKTQQSQEADLSVADESAIVQVVDSIIREAALQDASDIHIETTEDAVKVRYRI   538
NTDB id 1028 TT RS08215 WP 011173991.1        AELSEI......AK........EFAKKQAEEEV..PSPLELDESAAQKFVKQVIREAFLQDASDIHIEPRQNDVQVRLRI   541
NTDB id 1168 A1552VC RS11070 WP 000957200.1   SIQGAANQGKEISQDE..LANL...VKVSDDELQSIEDLSQDDSPVSRFINQVLLDAVRKGASDIHFEPYENQYRIRLRC   216
NTDB id 1401 DSB67 RS12665 WP 010643259.1     SLGQEKSGLKEINQDE..LAGL...VDIGEDEIENIEDLSQDESPVSRYINQILLDAVRKGASDIHFEPYENMYRVRLRC   216
NTDB id 1293 VP RS12240 WP 005479695.1        SLSHEKSGLKEINQEE..LASL...VDVGADEIDNIEDLSQDESPVSRYINQILLDAVRKGASDIHFEPYEKMYRVRLRC   216
NTDB id 1251 GC085 RS07725 WP 011946523.1     KESQGLS...EYVEDSGDLEGLEISADD.EDQDIDTATSVTDDAPIVKFVNKILLDAIRQGASDIHFEPYEREYRIRYRQ   225
NTDB id 1110 NGFG RS09235 WP 003689817.1      STSLLQ....ELGE..........GQEEEESHTLYIDNEEAEDGPVPRFIHKTLSDALRSGASDIHFEFYEHNARIRFRV   206
NTDB id 83449 JLC71 RS00195 WP 200916685.1    .SGATLK...NLEMDESEL...EISGGDPEP.QNDTANIDVDDAPVVKYLQKILLDAINANASDIHFEPYEKYYRIRYRQ   218
NTDB id 1058 ABD1 RS01615 WP 001274986.1      ESSFDF......NDEEFDL...DVNLDGPTAQEDEEEAPQGDEAPIVKYINKLLIDAIRMGASDLHFEPYEKSYRVRYRV   220
NTDB id 1018 ACIAD RS01685 WP 004920473.1     ETHFNF......DTEELDL...DVEVDPHTTDDDDEDDKLKDEAPIVKYINKLLIDAIRMSASDLHFEPYEKSYRVRYRV   220
consensus                                               **  **  *   ** **  ***  * * *  ***************!** *!!!*!*!**** ***!*!*

logo

NTDB id 1306 DR RS10060 WP 027479822.1        DGALREQNSFPKGAAQQIMARLKIMGHLDIAERRVPQDGRVRF..KKGSIDLDLRLSTLPTVYGEKAVMRLLQKASNIPE   616
NTDB id 1028 TT RS08215 WP 011173991.1        DGALRPYSTLPKGALNAVISVVKIMGGLNIAEKRLPQDGRVRY..REGAIDVDLRLSTLPTVYGEKAVMRLLKKASDIPE   619
NTDB id 1168 A1552VC RS11070 WP 000957200.1   DGILVETQQPASHLSRRLAARIKILSKLDIAERRLPQDGRIKLRLSR.DTAIDMRVSTLPTLWGEKIVLRLLDSSAANLD   295
NTDB id 1401 DSB67 RS12665 WP 010643259.1     DGILIEIQQPPSHLSRRLSARIKILSKLDIAERRLPQDGRIKLKLNQ.DTAIDMRVSTLPTLFGEKIVLRLLDSSSAALD   295
NTDB id 1293 VP RS12240 WP 005479695.1        DGILIETQQPPNHLSRRLSARIKILSKLDIAERRLPQDGRIKLKLNQ.DTAIDMRVSTLPTLFGEKIVLRLLDSSSASLD   295
NTDB id 1251 GC085 RS07725 WP 011946523.1     DGILHEVATPPASLSSRITARIKVMSNLDISERRIPQDGGFKMKISK.SRAIDFRVSTCPTSAGEKVVMRVLDSGAAKLG   304
NTDB id 1110 NGFG RS09235 WP 003689817.1      DGQLREVVQPPIAVRGQLASRIKVMSRLDISEKRIPQDGRMQLTFQKGGKPVDFRVSTLPTLFGEKVVMRILNSDAASLN   286
NTDB id 83449 JLC71 RS00195 WP 200916685.1    DGILREIAQPPLAIKEKIASRIKVISKLDIAEKRVPQDGRMKLVLSK.SRAIDFRVSTLPTLHGEKIVMRILDPTSATLG   297
NTDB id 1058 ABD1 RS01615 WP 001274986.1      DGVLRQIANPPLQLANRLASRLKVMSQMDISEKRVPQDGRIKLKLSK.SKAIDFRVNSLPTLFGEKLVLRILDPSSAMLG   299
NTDB id 1018 ACIAD RS01685 WP 004920473.1     DGVLRLIATPPLQLATRLASRLKVMSQMDISEKRVPQDGRIKLKMSK.SKTIDFRVNSLPTLFGEKIVLRILDPASAMLG   299
consensus                                     !!*!*** ***   *  *** **!******!*!*!*!!!!********  ****!*!****!!**!!!*!*!*!***** **

logo

NTDB id 1306 DR RS10060 WP 027479822.1        LEQLGFSEYNYARYTEIIERPNGIFLVTGPTGSGKSFTCFSTLKRIAKPEKNTTTIEDPIEYEVPGIVQSQVNNSTGMTF   696
NTDB id 1028 TT RS08215 WP 011173991.1        IEDLGFAPGVFERFKEVISKPYGIFLITGPTGSGKSFTTFSILKRIATPDKNTQTIEDPVEYEIPGINQTQVNPQAGLTF   699
NTDB id 1168 A1552VC RS11070 WP 000957200.1   IDKLGYNPQQKQLYLNALKRPQGMILMTGPTGSGKTVSLYTGLRILNTSQINISTAEDPVEINLSGINQVQVQPKIGFGF   375
NTDB id 1401 DSB67 RS12665 WP 010643259.1     INKLGYSDSQKQLYLDALRRPQGMILMTGPTGSGKTVSLYTGLSILNKPEINISTAEDPVEIYLPGINQVQVQPKIGFGF   375
NTDB id 1293 VP RS12240 WP 005479695.1        IDKLGYSEQQKQLYLEALRRPQGMILMTGPTGSGKTVSLYTGLNILNKPEINISTAEDPVEINLSGINQVQVQPKIGFGF   375
NTDB id 1251 GC085 RS07725 WP 011946523.1     IEALGFNPVQRTNFLKAIQRPQGMILVTGPTGSGKTVTLYTALNILNTIEVNISTAEDPVEIKVPGINQVNINPKAGLTF   384
NTDB id 1110 NGFG RS09235 WP 003689817.1      IDQLGFEPFQKKLLLEAIHRPYGMVLVTGPTGSGKTVSLYTCLNILNTESVNIATAEDPAEINLPGINQVNVNDKQGLTF   366
NTDB id 83449 JLC71 RS00195 WP 200916685.1    VDALGYEPDQKAALLNAVTRPYGMVLVTGPTGSGKTVSLYTCLNILNEPGTNISTAEDPAEINLPGVNQVNVNDKAGLTF   377
NTDB id 1058 ABD1 RS01615 WP 001274986.1      IDALGYEEDQKALFMEALDKPQGMLLITGPTGSGKTVSLYTGLNILNTESSNISTAEDPVEINLEGINQVNVNPKVGLTF   379
NTDB id 1018 ACIAD RS01685 WP 004920473.1     IDALGYEPEQKALFMEALNKPQGMLLITGPTGSGKTVSLYTGLNILNTEHANISTAEDPVEINLEGVNQVNVNPKVGLTF   379
consensus                                     ** !!* * ** ****** *!*!**!*!!!!!!!!*******!*******!**!*!!!*!****!**!*******!**!
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logo

NTDB id 1306 DR RS10060 WP 027479822.1        ARALRAFLRQDPDIIFVGEIRDQETAKIAVEAALTGHMVLATLHTNDAPGAVTRLEEMGIENFNISAAVMGVLAQRLVRR   776
NTDB id 1028 TT RS08215 WP 011173991.1        ARALRAFLRQDPDIIMVGEIRDSETAKIATEAALTGHLVIATLHTNDAAQAITRLDEMGVEPFNISAALIGVLSQRLVRR   779
NTDB id 1168 A1552VC RS11070 WP 000957200.1   AEALRSFLRQDPDVVMVGEIRDLETAEIAVKAAQTGHLVLSTLHTNSAAETVIRLANMGVEPFNLASSLSLIIAQRLARR   455
NTDB id 1401 DSB67 RS12665 WP 010643259.1     AEALRSFLRQDPDVVMVGEIRDLDTAEIAIKASQTGHLVLSTLHTNSAAETVIRLSNMGVESFNLASSLSLIIAQRLARK   455
NTDB id 1293 VP RS12240 WP 005479695.1        AEALRSFLRQDPDVVMVGEIRDLDTAEIAIKASQTGHLVLSTLHTNSAAETIIRLSNMGVESFNLASSLSLIIAQRLARK   455
NTDB id 1251 GC085 RS07725 WP 011946523.1     SGALRSFLRQDPDIIMVGEIRDLETAEIAIKAAQTGHLVLSTLHTNSAAETLNRLVNMGIPTFNIASSVTLIIAQRLARK   464
NTDB id 1110 NGFG RS09235 WP 003689817.1      AAALKSFLRQDPDIIMVGEIRDLETADIAIKAAQTGHMVFSTLHTNNAPATLSRMLNMGVAPFNIASSVSLIMAQRLLRR   446
NTDB id 83449 JLC71 RS00195 WP 200916685.1    SAALKAFLRQDPDIIMVGEIRDLETADIAIKAAQTGHMVFSTLHTNDAPGTLTRLLNMGVAPFNVASSVILITAQRLARR   457
NTDB id 1058 ABD1 RS01615 WP 001274986.1      AAALKSFLRQDPDIIMVGEIRDLETAEIAIKAAQTGHMVMSTLHTNSAPETLTRLRNMGVPSFNIATSVNLVIAQRLARR   459
NTDB id 1018 ACIAD RS01685 WP 004920473.1     AAALRSFLRQDPDIIMVGEIRDLETAEIAIKAAQTGHLVMSTLHTNNAAETLTRLRNMGVASFNIATSVNLVIAQRLARR   459
consensus                                     **!!**!!!!!!!***!!!!!!**!!*!!**!**!!!*!**!!!!!*!*****!* *!!***!!**********!!!*!*

logo

NTDB id 1306 DR RS10060 WP 027479822.1        VCSECKQPTNADPEV.LRRLGISERDIRG..ANLMRGTGCPRCGGTGYKGRMGIHELMVMDDSLRRTIGAGRPAAEIRDV   853
NTDB id 1028 TT RS08215 WP 011173991.1        VCEHCKVEVKPDPET.LRRLGLSEAEIQG..ARLYKGMGCERCGGTGYKGRYAIHELLVVDDEIRHAIVAGKSATEIKEI   856
NTDB id 1168 A1552VC RS11070 WP 000957200.1   LCKHCKIAVRPSA.LLQSQFAFQPNEI....LYEANAAGCNECT.GGYSGRVGIYEVMAFNTELAEAIMQRASIHQIERL   529
NTDB id 1401 DSB67 RS12665 WP 010643259.1     LCPYCRQPQEPNS.QLQH.IGIANTEQ....IFQANPDGCNECT.HGYSGRTGIYEVMKFDESLSEALIKGASVHELEKL   528
NTDB id 1293 VP RS12240 WP 005479695.1        LCPYCKQPQEHTV.QLQH.LGIQTTDN....IFRANPDGCNECT.HGYSGRTGIYEVMRFDESLSEALIKGASVHELEKL   528
NTDB id 1251 GC085 RS07725 WP 011946523.1     LCNQCKAVRDDFTNQGLIELGFKEADLVN..LKLYKAVGCEQCT.SGYRGRVGLFEVLPMTKELGQLIMSGGNSLDILKL   541
NTDB id 1110 NGFG RS09235 WP 003689817.1      LCSSCKQEVERPSASALKEVGFTDEDLAK.DWKLYRAVGCDRCRGQGYKGRAGVYEVMPISEEMQRVIMNNGTEVGILDV   525
NTDB id 83449 JLC71 RS00195 WP 200916685.1    LCSHCKAPADIPNEA.LLRAGFTEGDLDG.SWTPYKAIGCDLCKDTGYKGRVGIYQVMPISDEINRIIMRNGTAIDVADQ   535
NTDB id 1058 ABD1 RS01615 WP 001274986.1      LCSQCKIPADIPKQS.LLEMGFTEQDLAHPDFRVFQPVGCLECR.EGYKGRVGIYEVMKVTPEISKIIMEDGNALEIAAA   537
NTDB id 1018 ACIAD RS01685 WP 004920473.1     LCSQCKRPIQVPERS.LLEMGFTPEDLAQPEFQIFEPVGCHDCR.EGYKGRVGIYEVMKITPEISKIIMEDGNALEIAAT   537
consensus                                     *!* !* * *       *  ***** **      *****!! *!   !!*!!******* ***** **** *******  *

logo

NTDB id 1306 DR RS10060 WP 027479822.1        ALGESGLRSLRQDGIEKALQGLTTLEEVLAVTAS   887
NTDB id 1028 TT RS08215 WP 011173991.1        AR.RKGMKTLREDGLYKALQGITTLEEVLARTIE   889
NTDB id 1168 A1552VC RS11070 WP 000957200.1   AK.ANGMQTLQESGLEKLREGITSFAELQRVLYF   562
NTDB id 1401 DSB67 RS12665 WP 010643259.1     AI.ANGMQTLQMSGIEKLKQGITSFRELQRVLYF   561
NTDB id 1293 VP RS12240 WP 005479695.1        AI.ANGMSTLQMSGIEKLKQGITSFSELQRVLYF   561
NTDB id 1251 GC085 RS07725 WP 011946523.1     AQ.SEGMLTIFQSGIEKVKEGITTIEEVNRVTVD   574
NTDB id 1110 NGFG RS09235 WP 003689817.1      AY.KEGMVDLRRAGILKIMQGITSLEEVTANTND   558
NTDB id 83449 JLC71 RS00195 WP 200916685.1    AR.LEGVRDLRQSGLLKVKQGMTSLEEVMAVTNE   568
NTDB id 1058 ABD1 RS01615 WP 001274986.1      SE.KLGFNNLRRSGLKKVMQGVTSLQEVNRVTSE   570
NTDB id 1018 ACIAD RS01685 WP 004920473.1     AE.TLGFNNLRRSGLKKVMQGVTSLQEINRVTSE   570
consensus                                     *     !* *** *!**!* *!*!***!***** *
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X non conserved

X similar

X $\geq$ 50% conserved