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logo

                                                MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR
NTDB id 1410 SGL RS13950 WP 010873519.1         ................................................................................    0
NTDB id 1306 DR RS10060 WP 027479822.1          MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR   80
NTDB id 1168 A1552VC RS11070 WP 000957200.1     ................................................................................    0
NTDB id 1293 VP RS12240 WP 005479695.1          ................................................................................    0
NTDB id 1401 DSB67 RS12665 WP 010643259.1       ................................................................................    0
NTDB id 1110 NGFG RS09235 WP 003689817.1        ................................................................................    0
NTDB id 1251 GC085 RS07725 WP 011946523.1       ................................................................................    0
NTDB id 153740 AKN87 RS10145 WP 053103341.1     ................................................................................    0
NTDB id 1058 ABD1 RS01615 WP 001274986.1        ................................................................................    0
NTDB id 1018 ACIAD RS01685 WP 004920473.1       ................................................................................    0
consensus


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                                                SIRPQTALNLQAFPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA
NTDB id 1410 SGL RS13950 WP 010873519.1         ................................................................................    0
NTDB id 1306 DR RS10060 WP 027479822.1          SIRPQTALNLQAFPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA  160
NTDB id 1168 A1552VC RS11070 WP 000957200.1     ................................................................................    0
NTDB id 1293 VP RS12240 WP 005479695.1          ................................................................................    0
NTDB id 1401 DSB67 RS12665 WP 010643259.1       ................................................................................    0
NTDB id 1110 NGFG RS09235 WP 003689817.1        ................................................................................    0
NTDB id 1251 GC085 RS07725 WP 011946523.1       ................................................................................    0
NTDB id 153740 AKN87 RS10145 WP 053103341.1     ................................................................................    0
NTDB id 1058 ABD1 RS01615 WP 001274986.1        ................................................................................    0
NTDB id 1018 ACIAD RS01685 WP 004920473.1       ................................................................................    0
consensus


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                                                GRTGGKLGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM
NTDB id 1410 SGL RS13950 WP 010873519.1         ................................................................................    0
NTDB id 1306 DR RS10060 WP 027479822.1          GRTGGKLGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM  240
NTDB id 1168 A1552VC RS11070 WP 000957200.1     ................................................................................    0
NTDB id 1293 VP RS12240 WP 005479695.1          ................................................................................    0
NTDB id 1401 DSB67 RS12665 WP 010643259.1       ................................................................................    0
NTDB id 1110 NGFG RS09235 WP 003689817.1        ................................................................................    0
NTDB id 1251 GC085 RS07725 WP 011946523.1       ................................................................................    0
NTDB id 153740 AKN87 RS10145 WP 053103341.1     ................................................................................    0
NTDB id 1058 ABD1 RS01615 WP 001274986.1        ................................................................................    0
NTDB id 1018 ACIAD RS01685 WP 004920473.1       ................................................................................    0
consensus
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NTDB id 1410 SGL RS13950 WP 010873519.1      .....................................................MTSSSS      6
NTDB id 1306 DR RS10060 WP 027479822.1       LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQRGRLGEQLVQEGTLSRDQL    320
NTDB id 1168 A1552VC RS11070 WP 000957200.1  ....................................................................      0
NTDB id 1293 VP RS12240 WP 005479695.1       ....................................................................      0
NTDB id 1401 DSB67 RS12665 WP 010643259.1    ....................................................................      0
NTDB id 1110 NGFG RS09235 WP 003689817.1     ....................................................................      0
NTDB id 1251 GCO85 RS07725 WP 011946523.1    ....................................................................      0
NTDB id 153740 AKN87 RS10145 WP 053103341.1  ....................................................................      0
NTDB id 1058 ABD1 RS01615 WP 001274986.1     ....................................................................      0
NTDB id 1018 ACIAD RS01685 WP 004920473.1    ....................................................................      0
consensus
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NTDB id 1410 SGL RS13950 WP 010873519.1      SKKSRAVALRNYFSPFGNKLVAGGHVDAEQLRQALVQVKKTGRSLPEELKAVTGRELSPELLRQYKKNQLFELKVLYGVD     86
NTDB id 1306 DR RS10060 WP 027479822.1       REALQVQAREGKVKPLGEVITELGFASPDEVDSALQKQNVGGGRLEDTLVQ..SGKLSPEMLAR.......SLAAQLGYE    391
NTDB id 1168 A1552VC RS11070 WP 000957200.1  ...........MLTNLVAILRQAELISATQEQAVVTQVSASGTSVPEALLE..LSIFHAQELTE.......QLSHIFGLP     60
NTDB id 1293 VP RS12240 WP 005479695.1       ...........MHSNLSTILRQKGLLTFSQEESLIEQVKASGISMPEALLS..SGFFTSSELAE.......HLSSIFGLS     60
NTDB id 1401 DSB67 RS12665 WP 010643259.1    ...........MLTNLSTVLRQAGILTFSQEESLLEQVKASGISMPEALLS..SGFFQSHELTE.......HLSSVFGLS     60
NTDB id 1110 NGFG RS09235 WP 003689817.1     ...........MSVGLLRILVQNQVVTVEQAE.HYYNESQAGKEVLPMLFS..DGVISPKSLAA.......LIARVFSYS     59
NTDB id 1251 GCO85 RS07725 WP 011946523.1    ...MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVK..NKILSAEQIAL.......TAAQNFGVP     68
NTDB id 153740 AKN87 RS10145 WP 053103341.1  ......MELSSGLIGMAGQIVKLGLADYKKVLNSVADARLAGQPIISWLVH..NKIVTSREVMT.......LATSQFGVA     65
NTDB id 1058 ABD1 RS01615 WP 001274986.1     ...MSALHTSPKFTGFFRRLVEEKHVSAATMQTALDAAKRAKQDTVAYLIE..EVHLSPSLLAE.......TISAEFAEP     68
NTDB id 1018 ACIAD RS01685 WP 004920473.1    ...MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDIVPYLID..NFSISPLTIAE.......IISLEFGEP     68
consensus                                              ****  ***  ****  **   * * *  * **** **      ***  ***         ** ****
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NTDB id 1410 SGL RS13950 WP 010873519.1      SVDPEVAPIATQQMAELIGRFFPLDTCRRFKFLPLAQQEGDPPSVLVAMVDPDNLAAQDELNRILRVKGFELRRLVVTQD    166
NTDB id 1306 DR RS10060 WP 027479822.1       FLDPIQNPPDP.....KVALMTPEATARRYVVVPVRLQ...GNSLVVAMKDPRNVFALDDLK...LITGKEILPAVMAEK    460
NTDB id 1168 A1552VC RS11070 WP 000957200.1  ETDLSRYDYAN.....LCQQLGLRELITRYDALPIAKQ...GNLLLLAVSDPTLLQAEEFR...FATGLQVELALADHR    129
NTDB id 1293 VP RS12240 WP 005479695.1       QPELSQYEYAS.....LCQQLGLRELITRHNALPLHRT...PSTLLLAVADPTNQQAEDDFR...FATGLQVELVLADFR    129
NTDB id 1401 DSB67 RS12665 WP 010643259.1    CTSLNQYEYAS.....LCQTLGLRDLITRHNALPLNRT...SSTLILAVADPTNLQAEDDFR...FATGLQVELVLADFR    129
NTDB id 1110 NGFG RS09235 WP 003689817.1     ILDLRHYPRHR.....VLMGVLTEEQMVEFHCVPVFRR...GDKVFFAVSDPTQMPQIQKTV...SAAGIAVELVIVEDD    128
NTDB id 1251 GCO85 RS07725 WP 011946523.1    MLDINCIDVGT.....IPVNLVNEKLIKRHAMVPLFSR...GTNLYLATDDPSKQASLKEIQ...FHTGLNTHAIVVETD    137
NTDB id 153740 AKN87 RS10145 WP 053103341.1  SFDLSCFKIKL.....LSKDLLSERIVRKHRVLPLAKR...GNLLYVALADHGDQLVLNEIK...FTTGCAVELVVVEQD    134
NTDB id 1058 ABD1 RS01615 WP 001274986.1     YFDLDVYDTSQ.....IPKDLVDQKLILKHRILPLIQR...GQILYVATSNPSNIEAIDAIR...FNSKLLVEPVIVEHH    137
NTDB id 1018 ACIAD RS01685 WP 004920473.1    LLDLGVFDPAL.....FLKDKIDEKLIQKYRIMPLVHR...GHVLYVATSNPTNIEAMDAIR...FNSKLKVEPIIVEHD    137
consensus                                     ***  ** *       *   ** **** ** **!* **    *  * *!* ***** ******   * *** ******* *
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NTDB id 1410 SGL RS13950 WP 010873519.1      DFNQLLEQYYAIREEWEAAQEKRDQERALDKLNDLTDIVGSI.DMNLGEDKEDVDNALDSNDANQAPVINLVNKILAKAL  245
NTDB id 1306 DR RS10060 WP 027479822.1       DIIRLIERYFGEKGFE..............KL...NKELAER.......NKTQQSQEADLSVADESAIVQVVDSIIREAA  516
NTDB id 1168 A1552VC RS11070 WP 000957200.1  ALQAAIRRLYGRSIQG..............AANQGKEISQDE.LANLVKVSDDELQSIEDLSQDDSPVSRFINQVLLDAV  194
NTDB id 1293 VP RS12240 WP 005479695.1       ELSTAIRRLYGRSLSH..............EKSGLKEINQEE.LASLVDVGADEIDNIEDLSQDESPVSRYINQILLDAV  194
NTDB id 1401 DSB67 RS12665 WP 010643259.1    ELTAAIRRLYGRSLGQ..............EKSGLKEINQDE.LAGLVDIGEDEIENIEDLSQDESPVSRYINQILLDAV  194
NTDB id 1110 NGFG RS09235 WP 003689817.1     QLAGLLDWVGSRST.S..............LLQELGEGQEEE.........ESHTLYIDNEEAEDGPVPRFIHKTLSDAL  184
NTDB id 1251 GCO85 RS07725 WP 011946523.1    KLSALIDNLLTAKESQ..............GLSEYVEDSGDLEGLETSADDEDQDIDTATSVTDDAPIVKFVNKILLDAI  203
NTDB id 153740 AKN87 RS10145 WP 053103341.1  KLDLLIEQLYDSLL.E..............SFESLADEVLEE.....SEY.IEIQNELIIDAIDEAPIVKFVNSILLRAI  193
NTDB id 1058 ABD1 RS01615 WP 001274986.1     KLEKVLGQHFAEES................SFD.FNDEEFDL.DVNLDGP.TAQEDEEEAPQGDEAPIVKYINKLLIDAI  198
NTDB id 1018 ACIAD RS01685 WP 004920473.1    KLERLLSEHFVEET................HFN.FDTEELDL.DVEVDPH.TTDDDDEDDKLKDEAPIVKYINKLLIDAI  198
consensus                                     *  *** ***                    * **  **    *    **   ***     **********************!*
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NTDB id 1410 SGL RS13950 WP 010873519.1      QEGTSDIHVEPQEETLRIRFRKDGVLNQAFDPLPRKIIPAVVARFKIMADMDIAERRSPQDGKIRRIYQ..GRKVDFRVN  323
NTDB id 1306 DR RS10060 WP 027479822.1       LQDASDIHIETTEDAVKVRYRIDGALREQNSF.PKGAAQQIMARLKIMGHLDIAERRVPQDGRVRFK..KGSIDLDLRLS  593
NTDB id 1168 A1552VC RS11070 WP 000957200.1  RKGASDIHFEPYENQYRIRLRCDGILVETQQP.ASHLSRRLAARIKILSKLDIAERRLPQDGRIKLRLSR.DTAIDMRVS  272
NTDB id 1293 VP RS12240 WP 005479695.1       RKGASDIHFEPYEKMYRVRLRCDGILIETQQP.PNHLSRRLSARIKILSKLDIAERRLPQDGRIKLKLNQ.DTAIDMRVS  272
NTDB id 1401 DSB67 RS12665 WP 010643259.1    RKGASDIHFEPYENMYRVRLRCDGILIETQQP.PSHLSRRLSARIKILSKLDIAERRLPQDGRIKLKLNQ.DTAIDMRVS  272
NTDB id 1110 NGFG RS09235 WP 003689817.1     RSGASDIHFEFYEHNARIRFRVDGQLREVVQP.PIAVRGQLASRIKVMSRLDISEKRIPQDGRMQLTFQKGGKPVDFRVS  263
NTDB id 1251 GCO85 RS07725 WP 011946523.1    RQGASDIHFEPYEREYRIRYRQDGILHEVATP.PASLSSRITARIKVMSNLDISERRIPQDGGFKMKISK.SRAIDFRVS  281
NTDB id 153740 AKN87 RS10145 WP 053103341.1  KIGASDLHFEPYENSYRVRFRTDGILHEVSKQ.PVGLRDRISTRLKVMASIDISERRKPQDGRIKLKVSS.VKSINFRVN  271
NTDB id 1058 ABD1 RS01615 WP 001274986.1     RMGASDLHFEPYEKSYRVRYRVDGVLRQIANP.PLQLANRLASRLKVMSQMDISEKRVPQDGRIKLKLSK.SKAIDFRVN  276
NTDB id 1018 ACIAD RS01685 WP 004920473.1    RMSASDLHFEPYEKSYRVRYRVDGVLRLIATP.PLQLATRLASRLKVMSQMDISEKRVPQDGRIKLKMSK.SKTIDFRVN  276
consensus                                    * **!!*!*!**!   ***!*! !!*! ** **  *  ** ** *!*!*** *!! !*!*!!!!********   *****!**
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NTDB id 1410 SGL RS13950 WP 010873519.1      TLPSRYGEKVCLRILDNSATQLGLDFLITNPETLQTVRELAARPYYGLMLVTGPTGSGKSTTLYSVLAERNSPEVNINTAE  403
NTDB id 1306 DR RS10060 WP 027479822.1       TLPTVYGEKAVMRLLQKASNIPELEQLGFSEYNYARYTEIIERPNGIFLVTGPTGSGKSFTCFSTLKRIAKPEKNTTTIE  673
NTDB id 1168 A1552VC RS11070 WP 000957200.1  TLPTLWGEKIVLRLLDSSAANLDIDKLGYNPQQKQLYLNALKRPQGMILMTGPTGSGKTVSLYTGLRILNTSQINISTAE  352
NTDB id 1293 VP RS12240 WP 005479695.1       TLPTLFGEKIVLRLLDSSSASLDIDKLGYSEQQKQLYLEALRRPQGMILMTGPTGSGKTVSLYTGLNILNKPEINISTAE  352
NTDB id 1401 DSB67 RS12665 WP 010643259.1    TLPTLFGEKIVLRLLDSSSAALDINKLGYSDSQKQLYLDALRRPQGMILMTGPTGSGKTVSLYTGLSILNKPEINISTAE  352
NTDB id 1110 NGFG RS09235 WP 003689817.1     TLPTLFGEKVVMRILNSDAASLNIDQLGFEPFQKKLLLEAIHRPYGMVLVTGPTGSGKTVSLYTCLNILNTESVNIATAE  343
NTDB id 1251 GCO85 RS07725 WP 011946523.1    TCPTSAGEKVVMRVLDSGAAKLGIEALGFNPVQRTNFLKAIQRPQGMILVTGPTGSGKTVTLYTALNILNTIEVNISTAE  361
NTDB id 153740 AKN87 RS10145 WP 053103341.1  TLPTLWGEKIVLRILDSDSTKLGIDSLGYESAQKKLYLEALDLPQGMILVTGPTGSGKSVSLYTGINIINTVERNISTAE  351
NTDB id 1058 ABD1 RS01615 WP 001274986.1     SLPTLFGEKLVLRILDPSSAMLGIDALGYEEDQKALFMEALDKPQGMLLITGPTGSGKTVSLYTGLNILNTESSNISTAE  356
NTDB id 1018 ACIAD RS01685 WP 004920473.1    SLPTLFGEKIVLRILDPASAMLGIDALGYEPEQKALFMEALNKPQGMLLITGPTGSGKTVSLYTGLNILNTEHANISTAE  356
consensus                                    **!***!!!***!*!***** **** !** * ** ****** *!*!**!*!!!!!!!!********************  **!**!*!
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NTDB id 1410 SGL RS13950 WP 010873519.1      DPIEYALPGITQVQVIREKGMDFSNILRAFMRQDPDVILVGETRDKETAKTAIEAALTGHLVLTTLHTNDAAGAIARLDE  483
NTDB id 1306 DR RS10060 WP 027479822.1       DPIEYEVPGIVQSQVNNSTGMTFARALRAFLRQDPDIIFVGEIRDQETAKIAVEAALTGHMVLATLHTNDAPGAVTRLEE  753
NTDB id 1168 A1552VC RS11070 WP 000957200.1  DPVEINLSGINQVQVQPKIGFGFAEALRSFLRQDPDVVMVGEIRDLETAEIAVKAAQTGHLVLSTLHTNSAAETVIRLAN  432
NTDB id 1293 VP RS12240 WP 005479695.1       DPVEINLSGINQVQVQPKIGFGFAEALRSFLRQDPDVVMVGEIRDLDTAEIAIKASQTGHLVLSTLHTNSAAETIIRLSN  432
NTDB id 1401 DSB67 RS12665 WP 010643259.1    DPVEIYLPGINQVQVQPKIGFGFAEALRSFLRQDPDVVMVGEIRDLDTAEIAIKASQTGHLVLSTLHTNSAAETVIRLSN  432
NTDB id 1110 NGFG RS09235 WP 003689817.1     DPAEINLPGINQVNVNDKQGLTFAAALKSFLRQDPDIIMVGEIRDLETADIAIKAAQTGHMVFSTLHTNNAPATLSRMLN  423
NTDB id 1251 GCO85 RS07725 WP 011946523.1    DPVEIKVPGINQVNINPKAGLTFSGALRSFLRQDPDIIMVGEIRDLETAEIAIKAAQTGHLVLSTLHTNSAAETLNRLVN  441
NTDB id 153740 AKN87 RS10145 WP 053103341.1  DPVEICLEGINQVNVNFKQGMDFAQALRAFLRQDPDVIMIGEIRDLETAEIAIKAAQTGHMVMSTLHTNSAADTLTRLRN  431
NTDB id 1058 ABD1 RS01615 WP 001274986.1     DPVEINLEGINQVNVNPKVGLTFAAALKSFLRQDPDIIMVGEIRDLETAEIAIKAAQTGHMVMSTLHTNSAPETLTRLRN  436
NTDB id 1018 ACIAD RS01685 WP 004920473.1    DPVEINLEGVNQVNVNPKVGLTFAAALRSFLRQDPDIIMVGEIRDLETAEIAIKAAQTGHLVMSTLHTNNAAETLTRLRN  436
consensus                                    !!*!****!**!*******!**!*  *!**!*!!!!!*****!!*!!**!!**!**!**!!!*!**!!!!!*!*****!*  *
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NTDB id 1410 SGL RS13950 WP 010873519.1      MGVEPFMVSGSLLGVLAQRLMRKVCTECRIAYH.PSKEELSRFGLSASGDEVVTFYKANSLSPDEIHLARQKGNLCQKCS  562
NTDB id 1306 DR RS10060 WP 027479822.1       MGIENFNISAAVMGVLAQRLVRRVCSECKQPTN.ADPEVLRRLGISERDI..RGA.............NLMRGTGCPRCG  817
NTDB id 1168 A1552VC RS11070 WP 000957200.1  MGVEPFNLASSLSLIIAQRLARRLCKHCKIAVR.PSALLQSQFAFQPNEI....L.............YEANAAGCNECT  494
NTDB id 1293 VP RS12240 WP 005479695.1       MGVESFNLASSLSLIIAQRLARKLCPYCKQPQE.HTVQL.QHLGIQTTDN....I.............FRANPDGCNECT  493
NTDB id 1401 DSB67 RS12665 WP 010643259.1    MGVESFNLASSLSLIIAQRLARKLCPYCRQPQE.PNSQL.QHIGIANTEQ....I.............FQANPDGCNECT  493
NTDB id 1110 NGFG RS09235 WP 003689817.1     MGVAPFNIASSVSLIMAQRLLRRLCSSCKQEVERPSASALKEVGFTDEDLLA.KDW.............KLYRAVGCDRCR  489
NTDB id 1251 GCO85 RS07725 WP 011946523.1    MGIPTFNIASSVTLIIAQRLARKLCNQCKAVRDDFTNQGLIELGFKEADL..VNL.............KLYKAVGCEQCT  506
NTDB id 153740 AKN87 RS10145 WP 053103341.1  MGVAAFNLAISINLIIAQRLARRLCNICKSVKK.IPVDVLLQEGFSLEQV..GTF.............KVYEAVGCDKCN  495
NTDB id 1058 ABD1 RS01615 WP 001274986.1     MGVPSFNIATSVNLVIAQRLARRLCSQCKIPAD.IPKQSLLEMGFTEQDLAHPDF.............RVFQPVGCLECR  502
NTDB id 1018 ACIAD RS01685 WP 004920473.1    MGVASFNIATSVNLVIAQRLARRLCSQCKRPIQ.VPERSLLEMGFTPEDLAQPEF.............QIFEPVGCHDCR  502
consensus                                    !!***!**********!!!!*!**!* !* * *  *  **  ****  **    *                ** ***!  *!*
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NTDB id 1410 SGL RS13950 WP 010873519.1      GSGYKGRVGVYEVMRNSERIAELINQGATTDRIKDCAV.EEGMITLLAYSLNLVQEGYTTLEEVERVTFTDTGLESEMRA  641
NTDB id 1306 DR RS10060 WP 027479822.1       GTGYKGRMGIHELMVMDDSLRRTIGAGRPAAEIRDVALGESGLRSLRQDGIEKALQGLTTLEEVLAVTAS.........  887
NTDB id 1168 A1552VC RS11070 WP 000957200.1  .GGYSGRVGIYEVMAFNTELAEAIMQRASIHQIERLAK.ANGMQTLQESGLEKLREGITSFAELQRVLYF.........  562
NTDB id 1293 VP RS12240 WP 005479695.1       .HGYSGRTGIYEVMRFDESLSEALIKGASVHELEKLAI.ANGMSTLQMSGIEKLKQGITSFSELQRVLYF.........  561
NTDB id 1401 DSB67 RS12665 WP 010643259.1    .HGYSGRTGIYEVMKFDESLSEALIKGASVHELEKLAI.ANGMQTLQMSGIEKLKQGITSFRELQRVLYF.........  561
NTDB id 1110 NGFG RS09235 WP 003689817.1     GQGYKGRAGVYEVMPISEEMQRVIMNNGTEVGILDVAY.KEGMVDLRRAGILKIMQGITSLEEVTANTND.........  558
NTDB id 1251 GCO85 RS07725 WP 011946523.1    .SGYRGRVGLFEVLPMTKELGQLIMSGGNSLDILKLAQ.SEGMLTIFQSGIEKVKEGITTIEEVNRVTVD.........  574
NTDB id 153740 AKN87 RS10145 WP 053103341.1  .AGYKGRIGIYEVVKITPTLQAIIMEGGNPIEINQQIR.KEGFNDLRTSALEKVIQGVVSLEEVNRVTKD.........  563
NTDB id 1058 ABD1 RS01615 WP 001274986.1     .EGYKGRVGIYEVMKVTPEISKIIMEDGNALEIAAASE.KLGFNNLRRSGLKKVMQGVTSLQEVNRVTSE.........  570
NTDB id 1018 ACIAD RS01685 WP 004920473.1    .EGYKGRVGIYEVMKITPEISKIIMEDGNALEIAATAE.TLGFNNLRRSGLKKVMQGVTSLQEINRVTSE.........  570
consensus                                      !!*!!*!**!*********  *** *** ***  **     !***** ********!*****!***** *
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logo


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NTDB id 1410 SGL RS13950 WP 010873519.1       KRKSALTCRTCAAELQQEWLDCPYCMTSRFS   672
NTDB id 1306 DR RS10060 WP 027479822.1        ................................  887
NTDB id 1168 A1552VC RS11070 WP 000957200.1   ................................  562
NTDB id 1293 VP RS12240 WP 005479695.1        ................................  561
NTDB id 1401 DSB67 RS12665 WP 010643259.1     ................................  561
NTDB id 1110 NGFG RS09235 WP 003689817.1      ................................  558
NTDB id 1251 GCO85 RS07725 WP 011946523.1     ................................  574
NTDB id 153740 AKN87 RS10145 WP 053103341.1   ................................  563
NTDB id 1058 ABD1 RS01615 WP 001274986.1      ................................  570
NTDB id 1018 ACIAD RS01685 WP 004920473.1     ................................  570
consensus
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X non conserved
X similar
X ≥ 50% conserved