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NTDB id 1381 A4U84 RS02010 WP 021113424.1     FRVEKSHFDEILNSLIKTHQLNLKKENGIYYIHQAQEHK......QHTTAQLVNALPKLITKTIKLHYSKASEVIESLTK   129
NTDB id 1347 HI 0435 AAC22094.1               LKLNNIDMPRLLQIIAKSKHLTLNKDDGIYYLNGSQSGK......GQVAGNLTTNEPHLVSHTVKLHFAKASELMKSLTT   141
NTDB id 1373 NTHI RS02645 WP 011272046.1      LKLNNIDMPRLLKIIAKSKHLTLNKDDGVYYLNGSQSGK......GQVAGNLTTNEPHLVSHTVKLHFAKASELMKSLTT   141
NTDB id 96540 FORC29 RS01940 WP 000815976.1   LHLTDVPWKQALQTVVKSAGLITRQEGNILSVHSIAWQNDNIARQEAEQAR.AQANLPLENRSITLQYADAGELAK....   134
NTDB id 1433 BW25113 RS17570 WP 000815987.1   LHLTDVPWKQALQTVVKSAGLITRQEGNILSVHSIAWQNNNIARQEAEQAR.AQANLPLENRSITLQYADAGELAK....   134
NTDB id 1193 PA5040 NP 253727.1               LRLQNVPWDQALDLVLKTKGLDKRKLGNVLLVAPADEIA.ARERQELEAQKQIAELAPLRRELIQVNYAKAADIAKLFQS   398
NTDB id 1186 GTF74 RS01320 WP 000788426.1     LRLDGVPWQQVLDIILQVKGLDKRVDGNVILIAPKEELD.LREKQALEKARLAEELGDLKSEIIKINFAKASDIAAMIGG   273
NTDB id 1175 A1552VC RS12105 WP 000788426.1   LRLDGVPWQQVLDIILQVKGLDKRVDGNVILIAPKEELD.LREKQALEKARLAEELGDLKSEIIKINFAKASDIAAMIGG   273
NTDB id 1408 DSB67 RS13980 WP 010643500.1     LRLDGVPWQQVLDIILQVKGLDKRVDGNVILVAPKDELD.LREKQQLEKQKLEKEIGELSSEIIKVNFAKASDIAEMING   274
consensus                                     ***  *** **! *******!  * ***** *    *      **  * ***   *  ! * ********!***** *

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NTDB id 1381 A4U84 RS02010 WP 021113424.1     .....GQGNLLSESGYLHFDDRSNSIIVKDSAASVKNFTQLIESLDKPTEQIAIEARIVTISSEHLQQLGVRWGLFSPNE   204
NTDB id 1347 HI 0435 AAC22094.1               .....GSGSLLSPAGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMDKPIEQIAIEARIVTITDESLKELGVRWGIFNPTE   216
NTDB id 1373 NTHI RS02645 WP 011272046.1      .....GSGSLLSSAGSITFDDRSNLLVIQDEPRFVQNIKKLIAEMDKPIEQIAIEARIVTITDESLKELGVRWGIFNPTE   216
NTDB id 96540 FORC29 RS01940 WP 000815976.1   .....AGEKLLSAKGSMTVDKRTNRLLLRDNKTALSALEQWVAQMDLPVGQVELSAHIVTINEKSLRELGVKWTLADAQH   209
NTDB id 1433 BW25113 RS17570 WP 000815987.1   .....AGEKLLSAKGSMTVDKRTNRLLLRDNKTALSALEQWVAQMDLPVGQVELSAHIVTINEKSLRELGVKWTLADAQH   209
NTDB id 1193 PA5040 NP 253727.1               VTSDGGQEGKEGGRGSITVDDRTNSIIAYQPQERLDELRRIVSQLDIPVRQVMIEARIVEANVGYDKSLGVRWGGAYHKG   478
NTDB id 1186 GTF74 RS01320 WP 000788426.1     ....EGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREIIESLDIPVKQVQIEARIVTVKEGNLEELGVRWGVMSTNG   349
NTDB id 1175 A1552VC RS12105 WP 000788426.1   ....EGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREIIESLDIPVKQVQIEARIVTVKEGNLEELGVRWGVMSTNG   349
NTDB id 1408 DSB67 RS13980 WP 010643500.1     ....EGNISMLSDRGSMTIDERTNSLLIRELPENIEVIREIIESLDIPVKQVQIEARIVTINEGNMDELGVRWGFTSING   350
consensus                                       **  *** *!****!*!*!****** *  *  ** **  *!*!* !* **!*!!** *  ***!!!*!**   *

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NTDB id 1381 A4U84 RS02010 WP 021113424.1     NHYKLAGNL.........................EGNGLTT..NNLNVNF..PVNPSASVALQIAAI.NSRVLDLELT   252
NTDB id 1347 HI 0435 AAC22094.1               NARRVAGSL.........................TGNSFENIADNLNVNFATTTTPAGSIALQVAKI.NGRLLDLELS   268
NTDB id 1373 NTHI RS02645 WP 011272046.1      NARRVAGSL.........................TGNSFENIADNLNVNFATTTTPAGSIALQVAKI.NGRLLDLELS   268
NTDB id 96540 FORC29 RS01940 WP 000815976.1   AGGV.GQVT.........................TLGSDLS...........VATATTHVGFNIGRI.NGRLLDLELS   249
NTDB id 1433 BW25113 RS17570 WP 000815987.1   AGGV.GQVT.........................TLGSDLS...........VATATTHVGFNIGRI.NGRLLDLELS   249
NTDB id 1193 PA5040 NP 253727.1               NWSG.YGKDG....NIGIKDEDGMNCGPIAGSCTFPTTGTSKSPSPFVDLG.....AKDATSGIGIGFITDNIILDLQLS   548
NTDB id 1186 GTF74 RS01320 WP 000788426.1     SHSV.GGSIESNLWQKGLLAD...................DEFPVDEFLNVNLASTSANASSIAFQVAKLGSGTLLDLELS   410
NTDB id 1175 A1552VC RS12105 WP 000788426.1   SHSV.GGSIESNLWQKGLLAD...................DEFPVDEFLNVNLASTSANASSIAFQVAKLGSGTLLDLELS   410
NTDB id 1408 DSB67 RS13980 WP 010643500.1     SHTV.GGSIENNLATIGLYDGGGE..........GDDGGSSVGIDDFLNVNLAATSPNATSIAFQVAKLGSDTLLDLELS   419
consensus                                       * ****                          *  *   * * **** * **  *********** ****!!!*!*

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NTDB id 1381 A4U84 RS02010 WP 021113424.1     ALESENNIEIIASPRLLTTDKKTASIKQGTEIPYAMYSKKEEITDIEFREAVLGLEVTPHISKQNQILLDLAISQNSPNN   332
NTDB id 1347 HI 0435 AAC22094.1               ALERENNVEIIASPRLLTTNKKSASIKQGTEIPYIVSNTRNDTQSVEFREAVLGLEVTPHISKDNNILLDLLVSQNSPGS   348
NTDB id 1373 NTHI RS02645 WP 011272046.1      ALERENNVEIIASPRLLTTNKKSASIKQGTEIPYIVSNTRNDTQSVEFREAVLGLEVTPHISKDNNILLDLLVSQNSPGS   348
NTDB id 96540 FORC29 RS01940 WP 000815976.1   ALEQKQQLDIIASPRLLASHLQPASIKQGSEIPYQVSSGESGATSVEFKEAVLGMEVTPTVLQKGRIRLKLHISQNVPGQ   329
NTDB id 1433 BW25113 RS17570 WP 000815987.1   ALEQKQQLDIIASPRLLASHLQPASIKQGSEIPYQVSSGESGATSVEFKEAVLGMEVTPTVLQKGRIRLKLHISQNVPGQ   329
NTDB id 1193 PA5040 NP 253727.1               AMEKTGNGEIVSQPKVVTSDKETAKILKGSEVPYQEAS.SSGATSTSFKEAALSLEVTPQITPDNRIIVEVKVTKDAPDY   627
NTDB id 1186 GTF74 RS01320 WP 000788426.1     ALQNESKAEIISSPRLITTNKQPAYIEQGTEIPYLESS.SSGASTVAFKKAVLSLKVTPQITPDNRLVLDLSVTQDRRGE   489
NTDB id 1175 A1552VC RS12105 WP 000788426.1   ALQNESKAEIISSPRLITTNKQPAYIEQGTEIPYLESS.SSGASTVAFKKAVLSLKVTPQITPDNRLVLDLSVTQDRRGE   489
NTDB id 1408 DSB67 RS13980 WP 010643500.1     ALQQESKAEIISSPRLITTNKKPAYIEQGTEIPYLESS.SSGATSVTFKKAVLSLKVTPQITPDNRLVLDLSVTQDRPGQ   498
consensus                                     !********!***!******* *!*!**!*!*!!****  *******!**!*!***!!! ** ******** **** **
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NTDB id 1381 A4U84 RS02010 WP 021113424.1     QIN...NTMVTIDKQEIKTQVLAKHGETIVLGGIFQHLIAKGEDKVPLLGSIPVIKRLFSQSQDKISKRELVIFVTPYIV   409
NTDB id 1347 HI 0435 AAC22094.1               RVAYGQNEVVSIDKQEINTQVFAKDGETIVLGGVFHDTITKSEDKVPLLGDIPVIKRLFSKESERHQKRELVIFVTPHIL   428
NTDB id 1373 NTHI RS02645 WP 011272046.1      RVAYGQNEVVSIDKQEINTQVFAKDGETIVLGGVFHDTITKSEDKVPLLGDIPVIKRLFSKESERHQKRELVIFVTPHIL   428
NTDB id 96540 FORC29 RS01940 WP 000815976.1   VLQQADGEVLAIDKQEIETQVEVKSGETLALGGIFTRKNKSGQDSVPLLGDIPWFGQLFRHDGKEDERRELVVFITPRLV   409
NTDB id 1433 BW25113 RS17570 WP 000815987.1   VLQQADGEVLAIDKQEIETQVEVKSGETLALGGIFTRKNKSGQDSVPLLGDIPWFGQLFRHDGKEDERRELVVFITPRLV   409
NTDB id 1193 PA5040 NP 253727.1               QNM..LNGVPPINKNEVNAKILVNDGETIVIGGVFSNEQSKSVEKVPFLGELPYLGRLFRRDTVTDRKNELLVFLTPRIM   705
NTDB id 1186 GTF74 RS01320 WP 000788426.1     TVKTGTGEAVSIDTQRIGTQVLVNNGETVVLGGIFQHSINNSVDKVPLLGDLPVLGALFRRTYEQMGKSELLIFVTPKVV   569
NTDB id 1175 A1552VC RS12105 WP 000788426.1   TVKTGTGEAVSIDTQRIGTQVLVNNGETVVLGGIFQHSINNSVDKVPLLGDLPVLGALFRRTYEQMGKSELLIFVTPKVV   569
NTDB id 1408 DSB67 RS13980 WP 010643500.1     VVKTGTGEAVAIDTQRIGTQVLVNNGETVVLGGIFQHSITNSVDKVPLLGDLPLLGALFRRSYENVGKSELLIFVTPKVV   578
consensus                                      *   * *****!***** ****** !!!***!!*!   **   * **!!*!!**!*** !!*** *    **!!**!*!!***
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NTDB id 1381 A4U84 RS02010 WP 021113424.1     KSEKIG.AEKQK.....   420
NTDB id 1347 HI 0435 AAC22094.1               KAGETLEALKQKSEGKK   445
NTDB id 1373 NTHI RS02645 WP 011272046.1      KAGETLEALKQKSAGKK   445
NTDB id 96540 FORC29 RS01940 WP 000815976.1   SSE..............   412
NTDB id 1433 BW25113 RS17570 WP 000815987.1   SSE..............   412
NTDB id 1193 PA5040 NP 253727.1               NNQAIAIGR........   714
NTDB id 1186 GTF74 RS01320 WP 000788426.1     IQ...............   571
NTDB id 1175 A1552VC RS12105 WP 000788426.1   IQ...............   571
NTDB id 1408 DSB67 RS13980 WP 010643500.1     IQ...............   580
consensus
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X non conserved
X similar
X ≥ 50% conserved