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logo
NTDB id 412 SMU RS09535 WP 002262393.1      ....MKSEIIAVGTEILTGQIVNTNSQFLSEKFAELGIDVYFQTAVGDNEERLLSVLKIAKERSDLIVLCGGLGPTEDDLTKQTL   81
NTDB id 170 SP RS09755 WP 000642718.1       ....MKAEIIAVGTEILTGQIVNTNAQFLSEKLAEIGVDVYFQTAVGDNEVRLLSLLEIASQRSSLVILTGGLGPTEDDLTKQTL   81
NTDB id 280 KZH43 RS08670 WP 000642700.1    ....MKAEIIAVGTEILTGQIVNTNAQFLSEKLAEIGVDVYFQTAVGDNEVRLLSLLEIASQRSSLVILTGGLGATEDDLTKQTL   81
NTDB id 205 SPR RS08830 WP 000642700.1      ....MKAEIIAVGTEILTGQIVNTNAQFLSEKLAEIGVDVYFQTAVGDNEVRLLSLLEIASQRSSLVILTGGLGATEDDLTKQTL   81
NTDB id 239 SPD RS09270 WP 000642701.1      ....MKAEIIAVGTEILTGQIVNTNAQFLSEKLAEIGVDVYFQTAVGDNEVRLLSLLEIASQRSSLVILTGGLGATEDDLTKQTL   81
NTDB id 538 SMSK321 RS11060 WP 000642686.1  ....MKAEIIAVGTEILTGQIVNTNAQFLSEKLAEIGVDVYFQTAVGDNEARLLSLLEIASQRSSLVILTGGLGPTEDDLTKQTL   81
NTDB id 510 SM12261 RS08320 WP 000642690.1  ....MKAEIIAVGTEILTGQIVNTNAQFLSEKLAEIGVDVYFQTAVGDNEARLLSLLEIASQRSSLVILTGGLGPTEDDLTKQTL   81
NTDB id 304196 PB01 RS06970 WP 151699536.1 ....MKAEIIAVGSELLLGQISNTNARFISSQLAELGIDVYYHTVVGDNSKRLFETIKLAEQRADLIIFTGGLGPTKDDLTKETI   81
NTDB id 124 BSU 16930 NP 389575.2           MEFPKKAEIIAVGSELLLGQIANTNAQFISKQLAEIGVNVFYHTAVGDNPERLKQVIRIAEERSDFIIFSGGLGPTKDDLTKETI   85
consensus                                       *!*!!!!!!*!*!*!!!*!!!**!*!***!!*!**!***!*!!!!* !!******!**!*******!!!!*!*!!!!!*!*

logo
NTDB id 412 SMU RS09535 WP 002262393.1      AKFLKRELVFDKTAQERLDEFFASRPTSMRTPNNECQAQIIAGSQPLSNKTGLAVGGLLEADGVTYVVLPGPPSELKPMVNKELL  166
NTDB id 170 SP RS09755 WP 000642718.1       AKFLGKALVFDPQAQEKLDIFFTLRPDYARTPNNERQAQIVEGAIPLPNETGLAVGGKLEVDGVTYVVLPGPPSELKPMVLNQLL  166
NTDB id 280 KZH43 RS08670 WP 000642700.1    AKFLGKALVFDPQAQEKLDIFFALRPDYARTPNNERQAQIVEGAIPLPNETGLAVGGKLEVDGVTYVVLPGPPSELKPMVLNQLL  166
NTDB id 205 SPR RS08830 WP 000642700.1      AKFLGKALVFDPQAQEKLDIFFALRPDYARTPNNERQAQIVEGAIPLPNETGLAVGGKLEVDGVTYVVLPGPPSELKPMVLNQLL  166
NTDB id 239 SPD RS09270 WP 000642701.1      AKFLGKALVFDPQAQEKLDIFFALRPDYARTPNNERQVQIVEGAIPLPNETGLAVGGKLEVDGVTYVVLPGPPSELKPMVLNQLL  166
NTDB id 538 SMSK321 RS11060 WP 000642686.1  AKFLGKELVFDPQAQEKLDVFFAQRPDYARTPNNERQAQIVEGATPLPNETGLAVGGILEVDGVTYVVLPGPPSELKPMVLNQLL  166
NTDB id 510 SM12261 RS08320 WP 000642690.1  AKFLGKELVFDPQAQEKLDVFFAQRPDYARTPNNERQAQLVEGAIPLPNETGLAVGGILEVDGVTYVVLPGPPSELKPMVLNQLL  166
NTDB id 304196 PB01 RS06970 WP 151699536.1 ATYLNKKLIMNEQALKYIKEFFKR.MNREMTENNKKQALVLDGCTVLSNKHGMAPGMLLKGDKHAYILLPGPPKEMEPMFQFEAK  165
NTDB id 124 BSU 16930 NP 389575.2           ANTLGRPLVLNDEAFQSIEDYFKR.TKRTMSPNNRKQALVIEGSDVLANHFGMAPGMLTEHESRYYMLLPGPPSELRPMFENEAK  169
consensus                                   !**!** !*****!******!* ********!!**!*****!***!*!**!*!*!*********!**!!!!!*!**!!******

logo
NTDB id 412 SMU RS09535 WP 002262393.1      PYL...SKTSEKLYSRVLRFFGIGESHLVTLLHDLIAEQTDPTIAPYAKTGEVTIRLSTKAHRQKEADSKLDKLEKKIITID...  245
NTDB id 170 SP RS09755 WP 000642718.1       PKL...M.TGSKLYSRVLRFFGIGESQLVTILADLIDNQIDPTLAPYAKTGEVTLRLSTKASSQEEANQALDILENQILDCQTFE  247
NTDB id 280 KZH43 RS08670 WP 000642700.1    PKL...M.TGSKLYSRVLRFFGIGESQLVTILADLIDNQIDPTLAPYAKTGEVTLRLSTKASSQEEANQALDILENQILDCQTFE  247
NTDB id 205 SPR RS08830 WP 000642700.1      PKL...M.TGSKLYSRVLRFFGIGESQLVTILADLIDNQIDPTLAPYAKTGEVTLRLSTKASSQEEANQALDILENQILDCQTFE  247
NTDB id 239 SPD RS09270 WP 000642701.1      PKL...M.TGSKLYSRVLRFFGIGESQLVTILADLIDNQIDPTLAPYAKTGEVTLRLSTKASSQEEANQALDILENQILDCQTFE  247
NTDB id 538 SMSK321 RS11060 WP 000642686.1  PKL...M.TGSKLYSRVLRFFGIGESQLVTILSDLIDNQTDPTLAPYAKTGEVTLRLSTKASSQEEANQVLDILENQILGRQTFE  247
NTDB id 510 SM12261 RS08320 WP 000642690.1  PKL...M.TGSKLHSRVLRFFGIGESQLVTILSDLIDNQTDPTLAPYAKTGEVTLRLSTKASSQEEANQVLDILENQILGRQTFE  247
NTDB id 304196 PB01 RS06970 WP 151699536.1 PLLQKMLNGETIILSHVMRFYGIGEAELETKVQDLLDNQTNPTLAPLASDGEVTLRLTAKSETEEEAWEKIEMLQTEVFN.....  245
NTDB id 124 BSU 16930 NP 389575.2           PLLLKKMGSNEKIVSTVLRFFGIGESQLEADLEDIIDAQTNPTIAPLAADGEVTLRLTAKHADEKETERLLKETEAVILE.....  249
consensus                                   !*!    * ******!*!*!!*!!!!**!**** !****!**!!*!!*!**!!!!*!!**!*****!*** ********** ****

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NTDB id 412 SMU RS09535 WP 002262393.1      ..NLADYFYGYGEENSLPQVVFDLLKEKGKTITAAESLTAGLFQARLADFAGASDIFKGGFITYSIEEKARMLGIPFEDLQLHGV  328
NTDB id 170 SP RS09755 WP 000642718.1       GISLRDFCYGYGEETSLASIVVEELKRQGKTIAAAESLTAGLFQATVANFSGVSSIFKGGFVTYSLEEKSRMLDIPAKNLEEHGV  332
NTDB id 280 KZH43 RS08670 WP 000642700.1    GISLRDFCYGYGEETSLASIVVEELKRQGKTIAAAESLTAGLFQATVANFSGVSSIFEGGFVTYSLEEKSRMLDIPAKNLEEHGV  332
NTDB id 205 SPR RS08830 WP 000642700.1      GISLRDFCYGYGEETSLASIVVEELKRQGKTIAAAESLTAGLFQATVANFSGVSSIFEGGFVTYSLEEKSRMLDIPAKNLEEHGV  332
NTDB id 239 SPD RS09270 WP 000642701.1      GISLRDFCYGYGEETSLASIVVEELKRQGKTIAAAESLTAGLFQATVANFSGVSSIFEGGFVTYSLEEKSRMLDIPAKNLEEHGV  332
NTDB id 538 SMSK321 RS11060 WP 000642686.1  GLSLRDLCYGYGEEASLASIVVEELKKQGKTITAAESLTAGLFQATVADFSGASSIFKGGFVTYSLEEKSKMLDIPVKDLEEQGV  332
NTDB id 510 SM12261 RS08320 WP 000642690.1  GLSLRDLCYGYGEETSLASIVVEELKRQGKTITSAESLTAGLFQARVADFSGASSIFKGGFVTYSLEEKSKMLDIPVKDLEEQGV  332
NTDB id 304196 PB01 RS06970 WP 151699536.1 ..RVGEYVYGYNN.DSLASKLQEILSENHLTISAAESMTAGLFQSELASIPGMGAVLVGGMITYTEDIKVKQLGVKKQTINEYGV  327
NTDB id 124 BSU 16930 NP 389575.2           ..RVGEFFYGYDD.TSLVKELSIACKEKGITISAAESFTGGLFSEWLTDHSGASKLFAGGVVCYTNDVKQNVLGVKKETLDRFGA  331
consensus                                   ********!!!****!!*************!!**!!!*!*!!!***** **!***** !!***!****!***!*** * ****!*
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logo
NTDB id 412 SMU RS09535 WP 002262393.1      VSAFTAEKMAERSRQLTQADLAISLTGVAGPDSLEGQPAGTVFIGLSSSKRTMAIKVLIGGRSRSDVRYIAVLHAFNLVRQTLLS   413
NTDB id 170 SP RS09755 WP 000642718.1       VSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHPVGTVFIGLAQDQGTEVIKVNIGGRSRADVRHIAVMHAFNLVRKALLS   417
NTDB id 280 KZH43 RS08670 WP 000642700.1    VSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHPVGTVFIGLAQDQGTEVIKVNIGGRSRADVRHIAVMHAFNLVRKALLS   417
NTDB id 205 SPR RS08830 WP 000642700.1      VSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHPVGTVFIGLAQDQGTEVIKVNIGGRSRADVRHIAVMHAFNLVRKALLS   417
NTDB id 239 SPD RS09270 WP 000642701.1      VSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHPVGTVFIGLAQDQGTEVIKVNIGGRSRADVRHIAVMHAFNLVRKALLS   417
NTDB id 538 SMSK321 RS11060 WP 000642686.1  VSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHPAGTVFIGLAQEHGTKVIKVNIGGRSRADVRHIAVMHAFNLVRKALLS   417
NTDB id 510 SM12261 RS08320 WP 000642690.1  VSEFTAQKMAEQARIKTQSDFGLSLTGVAGPDSLEGHPAGTVFIGLAQEQGTEVIKVNIGGRSRADVRHIAVMHAFNLVRKALLS   417
NTDB id 304196 PB01 RS06970 WP 151699536.1 VSSECAEAMAMRIKEKFGTNIGIGITGAAGPDAHGGQPAGTVWIGIVLNDLAPVSYRLNLSGTRNTNRLRAVKFTIHYLIQL.LR   411
NTDB id 124 BSU 16930 NP 389575.2           VSKECASELAKGVQKLTGSDIGISFTGVAGPDAQEGHEPGHVFIGISANGKEEV.HEFHFAGSRTGIRKRGAKYGCHLILKL.LE   414
consensus                                   !!***!***!**************!!*!!!!***!** !*!*!!*******************!***!***************!*
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logo
NTDB id 412 SMU RS09535 WP 002262393.1      HKNLV..   418
NTDB id 170 SP RS09755 WP 000642718.1       D......   418
NTDB id 280 KZH43 RS08670 WP 000642700.1    D......   418
NTDB id 205 SPR RS08830 WP 000642700.1      D......   418
NTDB id 239 SPD RS09270 WP 000642701.1      D......   418
NTDB id 538 SMSK321 RS11060 WP 000642686.1  D......   418
NTDB id 510 SM12261 RS08320 WP 000642690.1  D......   418
NTDB id 304196 PB01 RS06970 WP 151699536.1  EQGFKTI   418
NTDB id 124 BSU 16930 NP 389575.2           QK.....   416
consensus                                   *
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X non conserved

X similar

X $\geq$ 50% conserved