logo

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NTDB id 286 KZH43 RS05975 WP 000782664.1      MKMKQISDTTLKITISLEDLMDRGMEIADFLVPQEKTEEFFYAILDELEMPDSFLDTGMLSFRVTPKPDKVDVFVTKSKI   80
NTDB id 285 SPD RS06395 WP 000782664.1        MKMKQISDTTLKITISLEDLMDRGMEIADFLVPQEKTEEFFYAILDELEMPDSFLDTGMLSFRVTPKPDKVDVFVTKSKI   80
NTDB id 284 SPR RS06100 WP 000782664.1        MKMKQISDTTLKITISLEDLMDRGMEIADFLVPQEKTEEFFYAILDELEMPDSFLDTGMLSFRVTPKPDKVDVFVTKSKI   80
NTDB id 283 SP RS06670 WP 000782668.1         MKMKQISDTTLKITISLEDLMDRGMEIADFLVPQEKTEEFFYAILDELEMPDSFLDTGMLSFRVTPKPDKVDVFVTKSKI   80
NTDB id 299 STER RS01050 WP 011680667.1       MEMKQISETTLKIMITMEDLEEHGMELKDFLIPQEKTEEFFYTVMDELDLPDNFKNSGMLSFRVTPRKDRVDVFVTKSDL   80
NTDB id 331 STU RS10485 WP 011225365.1        MEMKQINETTLKIMITMEDLEEHGMELKDFLIPQEKTEEFFYTVMDELDLPDNFKNSGMLSFRVTPRKDRVDVFVTKSDL   80
NTDB id 1140213 DQM80 RS01250 WP 011106593.1  MEMKQISETTLKITISMDDLEERGMELKDFLIPQEKTEEFFYSVMDELDLPDNFKDSGMLSFRVTPRKDRLDVFVTKSEI   80
NTDB id 380 SMU RS01250 WP 002262749.1        MEMKQISETTLKITISMEDLEERGMELKDFLIPQEKTEEFFYTVMDELDLPENFKDSGMLSFRVTPRNDRIDVFVTKSEI   80
consensus                                     ! !!!!**!!!!!*!***!! **!!!* !!!*!!!!!!!!!! **!!!**!* ! **!!!!!!!!!* !**!!!!!!!! *
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logo

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NTDB id 286 KZH43 RS05975 WP 000782664.1      DQNLDFEDLSDLPDMEELAQMSPDEFIKTLEKSIADKTKDDIEAIQSLEQVEAKEEEQ..........EQAEQEAESKK   149
NTDB id 285 SPD RS06395 WP 000782664.1        DQNLDFEDLSDLPDMEELAQMSPDEFIKTLEKSIADKTKDDIEAIQSLEQVEAKEEEQ..........EQAEQEAESKK   149
NTDB id 284 SPR RS06100 WP 000782664.1        DQNLDFEDLSDLPDMEELAQMSPDEFIKTLEKSIADKTKDDIEAIQSLEQVEAKEEEQ..........EQAEQEAESKK   149
NTDB id 283 SP RS06670 WP 000782668.1         DQNLDFEDLSDLPDMEELAQMSPDEFIKTLEKSIADKTKDDIEAIQSLEQVEVKEEEQ..........EQAEQEAESKK   149
NTDB id 299 STER RS01050 WP 011680667.1       KEDLDFNDLSDM...EDYSGLSPEEFLKALEDNFMDK..GDIEAHHKLEEHDKTLKEVDETMTE.....PAKEVAEETIR   150
NTDB id 331 STU RS10485 WP 011225365.1        KEALDFNDLSDM...EDYSGLSPEEFLKALEDNFMDK..GDIEAHHKLEEHDKTLKEVDETMTE.....PAKEVAEETIR   150
NTDB id 1140213 DQM80 RS01250 WP 011106593.1  NKDINLEDLAEF...GDMSQMTPEDFFKSLEQSMREK..GDVKAHEKLEQIEEIMEDVVEATLANQSEAADPSTNHESEP   155
NTDB id 380 SMU RS01250 WP 002262749.1        NKNLNLEDLSDF...DDISKMSPEDFFNTLEETMREK..GDAAALDKLAEIEKREEEK..........TQQEKGETKEK   144
consensus                                       *****!!***    *** ***!**!***!! ** *!   !**!*  !**** ****              * *   ** *
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logo

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NTDB id 286 KZH43 RS05975 WP 000782664.1      EPYIYYILSFAKLADLVAFAKTV.TFEMETSELYKMNERYYLTILVDIENHPSPYPAWLLARMREFADDSDISRSVLQEY   228
NTDB id 285 SPD RS06395 WP 000782664.1        EPYIYYILSFAKLADLVAFAKTV.TFEMETSELYKMNERYYLTILVDIENHPSPYPAWLLARMREFADDSDISRSVLQEY   228
NTDB id 284 SPR RS06100 WP 000782664.1        EPYIYYILSFAKLADLVAFAKTV.TFEMETSELYKMNERYYLTILVDIENHPSPYPAWLLARMREFADDSDISRSVLQEY   228
NTDB id 283 SP RS06670 WP 000782668.1         EPYIYYILSFAKLADLVAFAKTV.TFEMETSELYKMNERYYLTILVDIENHPSPYPAWLLARMREFADDSDISRSVLQEY   228
NTDB id 299 STER RS01050 WP 011680667.1       EDYTHYVLAFSDFDQVVTFAQGLKNVSVEGSEFYKLGDVYYMTILLYLADEPDYYANNMYARFLEYANVADRTRPYLQEH   230
NTDB id 331 STU RS10485 WP 011225365.1        EDYTHYVLAFSDFDQVVTFAQGLKNVSVEGSEFYKLGDVYYMTILLYLADEPDYYANNMYARFLEYANVADRTRPYLQEH   230
NTDB id 1140213 DQM80 RS01250 WP 011106593.1  LDYVHYVLDFSTITEAVAFAKTI.DFSIEASELYKGSNCYHMTILLDVQQQPSYFANVMYARLIEHANPGSKTRAYLQEH   234
NTDB id 380 SMU RS01250 WP 002262749.1        RDYVHFVLDFPNIQQVINFAKTV.DYDVEASELFKESDAYHMTVLLNLEDKPDYYADLMFARMLEHAGRGTKTRAYLLEH   223
consensus                                     * !* **! !  * ****!!***  ***! !!**!* * !**!*!***** !* *   * !!* !*!   * *!  !*!
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logo

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NTDB id 286 KZH43 RS05975 WP 000782664.1      GQVLMSHDAVLNLQKIG..   245
NTDB id 285 SPD RS06395 WP 000782664.1        GQVLMSHDAVLNLQKIG..   245
NTDB id 284 SPR RS06100 WP 000782664.1        GQVLMSHDAVLNLQKIG..   245
NTDB id 283 SP RS06670 WP 000782668.1         GQVLMSHDAVLNLQKIG..   245
NTDB id 299 STER RS01050 WP 011680667.1       ATILMEEDALPVLQATKWS   249
NTDB id 331 STU RS10485 WP 011225365.1        ATILMEEDALPVLQATKWS   249
NTDB id 1140213 DQM80 RS01250 WP 011106593.1  GLQLMLDGAVEQLQKIELG   253
NTDB id 380 SMU RS01250 WP 002262749.1        GVQLIKADALQELQMIG..   240
consensus                                     * *!*   *!* *!!***
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X non conserved
X similar
X ≥ 50% conserved