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logo

NTDB id 478 HSISS4 RS06985 WP 021143811.1      MTENKNVILSARDIVVEFDVRDRVLTAIRGVSLDLVEGEVLALVGESGSGKSVLTKTFTGMLEENGRVASGSIDYRGQDL   80
NTDB id 327 STU RS16115 WP 011226304.1         MTENKNVILSARDIVVEFDVRDRVLTAIRGVSLDLVEGEVLALVGESGSGKSVLTKTFTGMLEENGRVASGSIDYRGKDL   80
NTDB id 295 STER RS06915 WP 011226304.1        MTENKNVILSARDIVVEFDVRDRVLTAIRGVSLDLVEGEVLALVGESGSGKSVLTKTFTGMLEENGRVASGSIDYRGKDL   80
NTDB id 1131117 H0514 RS09155 WP 232086215.1  ...MKETVLSVKNLHVNFHTYAGEVKAIRDVNFDLKKGETLAIVGESGSGKSVMTRTLMGLSEKNAKV.TGDITFKGRDM   76
NTDB id 361 SMU RS01310 WP 002262737.1         ..MSKEKILQVNNLHVNFHTYAGEVKAIRDVSFYLEKGETLAIVGESGSGKSVTTRTLMGLSAKNAEI.FGDIEFKGHNL   77
consensus                                      ****!***!******!*!********!!!*!***!**!!*!!*!!!!!!!!!!*!*!**!****!*****!*!***!***

logo

NTDB id 478 HSISS4 RS06985 WP 021143811.1      TKLKSHQDWAHIRGAKIATIFQDPMTSLDPIKTIGSQIVEVIVKHQGKSAKEAKKMAIDYMDKVGIPDAEKRFNEYPFQY  160
NTDB id 327 STU RS16115 WP 011226304.1         TKFKSHQDWAAIRGAKIATIFQDPMTSLNPIKTIGSQIIEVIVKHQGKTAKEAKKMAIDYMDKVGIPDAEKRFNEYPFQY  160
NTDB id 295 STER RS06915 WP 011226304.1        TKFKSHQDWAAIRGAKIATIFQDPMTSLNPIKTIGSQIIEVIVKHQGKTAKEAKKMAIDYMDKVGIPDAEKRFNEYPFQY  160
NTDB id 1131117 H0514 RS09155 WP 232086215.1  QALR.EEDWVKVRGNDIAMIFQDPMTSLDPTMKIGLQIAEPIMIHTKVSKKEALDMALQLMKDVGIPNAEEHINDYPHQW  155
NTDB id 361 SMU RS01310 WP 002262737.1         NDLK.EEDWVHIRGNDISMIFQDPMTSLDPTMRIGLQIAEPIIKHEKVTKKEALKRALDMMEKVGIPNAQEHINDYPHQW  156
consensus                                      *******!!* *!!**!**!!!!!!!!!*!***!!*!!*!*!**!*****!!!***!***!**!!!!*!****!*!!*!*

logo

NTDB id 478 HSISS4 RS06985 WP 021143811.1      SGGMRQRIVIAIALACRPDVLICDEPTTALDVTIQAQIIDLLKSLKEEYGFSVIFITHDLGVVASIADKVAVMYAGEIIE  240
NTDB id 327 STU RS16115 WP 011226304.1         SGGMRQRIVIAIALACRPDVLICDEPTTALDVTIQAQIIDLLKSLKEEYGFSVIFITHDLGVVASIADKVAVMYAGEIIE  240
NTDB id 295 STER RS06915 WP 011226304.1        SGGMRQRIVIAIALACRPDVLICDEPTTALDVTIQAQIIDLLKSLKEEYGFSVIFITHDLGVVASIADKVAVMYAGEIIE  240
NTDB id 1131117 H0514 RS09155 WP 232086215.1  SGGMRQRAVIAIALAADPEILIADEPTTALDVTIQAQILRLMKKIQEERDSSIIFITHDLGVVAGMADRVAVMYAGKILE  235
NTDB id 361 SMU RS01310 WP 002262737.1         SGGMRQRAVIAIALATNPEILVADEPTTALDVTIQAQILHLMKEIQKNTDSSIIFITHDLGVVAGMADRVAVMYAGKIVE  236
consensus                                      !!!!!!!*!!!!!!!**!**!**!!!!!!!!!!!!!!!!**!*!********!*!!!!!!!!!!!**!!*!!!!!!!*!*!

logo

NTDB id 478 HSISS4 RS06985 WP 021143811.1      YATVEEIFYEPCHPYTWSLLSSLPQLADDKGELFSIPGTPPSLYTPVVGDAFALRSDYAMQIDFEEKAPQFQVSDTHWAK  320
NTDB id 327 STU RS16115 WP 011226304.1         YATVEEIFYEPCHPYTWSLLSSLPQLADDNGKLFSIPGTPPSLYTPVVGDAFALRSDYALQIDFEEKAPQFQVSDTHWAK  320
NTDB id 295 STER RS06915 WP 011226304.1        YATVEEIFYEPCHPYTWSLLSSLPQLADDNGKLFSIPGTPPSLYTPVVGDAFALRSDYALQIDFEEKAPQFQVSDTHWAK  320
NTDB id 1131117 H0514 RS09155 WP 232086215.1  YGTVDEVFYNPQHPYTWGLLNSMPTIETESGSLEAIPGTPPDLLYPPKGDAFAPRNVFALAIDYQEEPPMFKVSDTHFAA  315
NTDB id 361 SMU RS01310 WP 002262737.1         YGTVDEVFYNPQHPYTWGLLNSMPTTNTASGSLHSIPGTPPDLLQPPQGDAFAPRNEFALDIDLKEEPPFFKVSDSHYAA  316
consensus                                      !*!!*!*!!!*!*!!!!!*!!!*!*!***** ! !**!!!!!!!*!**!**!!!!!*!***!**!!**!**!*!*!!!!*!*!*

logo

NTDB id 478 HSISS4 RS06985 WP 021143811.1      TWLLHEDAPKVEKPAVIQNLHEKIRANMGFAHLGDEEEGNA  361
NTDB id 327 STU RS16115 WP 011226304.1         TWLLHEDAPKVDKPAVIQNLHEKILANMGFAHLGDEEEGNA  361
NTDB id 295 STER RS06915 WP 011226304.1        TWLLHEDAPKVDKPAVIQNLHEKILANMGFAHLGDEEEGNA  361
NTDB id 1131117 H0514 RS09155 WP 232086215.1  TWLLDERAPKVTPPSPILKRWEKWSKMKEK...........  345
NTDB id 361 SMU RS01310 WP 002262737.1         TWLLDDRAPKITPPERILQRWEKWKSLQGEKHND.......  350
consensus                                      !!!!***!!!***!**!****!!* ****************
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X non conserved
X similar
X $\geq$ 50% conserved