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logo
NTDB id 1169879 ACKWRP RS23720 WP 000941264.1  MLQPKTRWKEKEYNGERVSELASKLQLSPLVVSLFLGRGLDTEDKILDFLNTENQEFHDPFLLEGMDRTVERVNKAIQNG   80
NTDB id 354 BSU 27620 NP 390640.1              MLASKMRWEIQRPDQDKVKSLTEQLHITPLVASLLVKRGFDTAESARLFLHTKDADFYDPFEMKGMKEAADRIKQAISQQ   80
consensus                                      !! *! !!    *  **!  !   !***!!! !! * !! !! *    !!*!   *!*!!! * !!    *!*  !! *

logo
NTDB id 1169879 ACKWRP RS23720 WP 000941264.1  EQILIFGDYDADGVSSTTVLYLALQELGADVEFYIPNRFTEGYGPNEEAFRWAHSAGFSLIITVDTGIAAVHEAKVAKEL  160
NTDB id 354 BSU 27620 NP 390640.1              EKIMIYGDYDADGVTSTSVMLHTLQKLSAQVDFYIPDRFKEGYGPNEQAFRSIKERGFSLIITVDTGIAAVHEAKVAKEL  160
consensus                                      ! !*!*!!!!!!!!*!!*!* * !! ! ! !*!!!! !! !!!!!!! !!!  *  !!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 1169879 ACKWRP RS23720 WP 000941264.1  GIDLIITDHHEPPPELPEALAIIHPKLDGGVYPFHYLAGVGVAFKVAHALLGRVPEHLLEIAVIGTVADLVSLHGENRLL  240
NTDB id 354 BSU 27620 NP 390640.1              GLDVIITDHHEPGPELPDVRAIVHPKQPGCTYPFKELAGVGVAFKLAHALLGELPDELLDLAAIGTIADLVPLHDENRLI  240
consensus                                      !*!*!!!!!!!!*!!!!*  !!*!!! *!* !!!* !!!!!!!!!*!!!!!! *!**!!**! !!!*!!!!*!! !!!!*

logo
NTDB id 1169879 ACKWRP RS23720 WP 000941264.1  VKRGLKHMRMTKNIGLKALFKVANVSQSEITEESIGFSIAPRINAVGRLEDATPAVHLLLSEDPEEAKELAEEIDELNKL  320
NTDB id 354 BSU 27620 NP 390640.1              ATLGLERLRRTNRLGLKELIKLSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEAEELAAEIDQLNKE  320
consensus                                         !! **! !  *!!! ! !*      !  !!**!! *!!!*!!!!!*! ! !!!!!!*!!!* !! !!! !!! !!!

logo
NTDB id 1169879 ACKWRP RS23720 WP 000941264.1  RKDIVKQITEEAIAEVENNFPPEENKVLVLAKEGWNPGVIGIVASKLVERFYRPTIVLCIDPVKETAKGSARSIAGFDLF  400
NTDB id 354 BSU 27620 NP 390640.1              RQKMVSKMTDEAIEMVEQQG..LDQTAIVVAKAGWNPGVVGIVASKLVDRFYRPAIVLGIDEEKGIAKGSARSIRGFNLF  398
consensus                                      !  *!  *!*!!!  !!** ** **  *!*!! !!!!!!*!!!!!!!!!*!!!!! !!!*!!* !  !!!!!!!! !! !!

logo
NTDB id 1169879 ACKWRP RS23720 WP 000941264.1  ANLSDCRELLPHFGGHPMAAGMTLHMNDVDELRRRLNEQAEVILTEEDFIPITAVDAFCKVEDVTLAAIEDMQKLAPFGV  480
NTDB id 354 BSU 27620 NP 390640.1              ESLSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTLTEEDFIPVQEVDLVCGVEDITVESIAEMNMLSPFGM  478
consensus                                        !!*!!**!!!!!!!!!!!!!!!*  !!**!! !!!! !*  !!!!!!!!*  !!  ! !!!*!*  ! *!* ! !!!*

logo
NTDB id 1169879 ACKWRP RS23720 WP 000941264.1  GNPKPRIAVKDAELESIRAIGSDGSHLKMALRDGQATLDTIGFGFGAYAKEISPVAKVSVIGEASINEWNNFKKPQLMVQ  560
NTDB id 354 BSU 27620 NP 390640.1              LNPKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGELQEGIVPGSRISIVGEMSINEWNNRKKPQLMIK  558
consensus                                       !!!!** !  ! !! *! !!   *!*!! *!     !!**!!  !     ! !  **!**!! !!!!!!! !!!!!!*
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logo
NTDB id 1169879 ACKWRP RS23720 WP 000941264.1  DIAVEAWQLFDWRSMRNVEANVAELPKEKITMVYFSKEVLHKFSLEDYKEHMM.....HASEVTELDEQYIVLLDLPKGT   635
NTDB id 354 BSU 27620 NP 390640.1              DAAVSEWQLFDLRGKRTWEDTVSALPSAKRAIVSFKEDSTTLLQTEDLRREVHVISSKDQAKAFDLDGAYIVLLDPPPSL   638
consensus                                      ! !!  !!!!! !  !  !  !  !!  !  *! !  *  *    !! * *********     *!!  !!!!!!*!*

logo
NTDB id 1169879 ACKWRP RS23720 WP 000941264.1  DELRDLFKVGFPSRIYTLFYQENNHLFSTVPTRDHFKWYYSFLSQKSPFSLRQYGEQLCQHKGWSKDTVNFMTQVFFELE   715
NTDB id 354 BSU 27620 NP 390640.1              DMLARLLEGKAPERIYFIFLNHEDHFLSTFPARDHFKWYYAFLLKRGAFDVKKHGSELAKHKGWSVETINFMTKVFFDLG   718
consensus                                      ! !  !     ! !!! *! **  !  !! ! !!!!!!!! !!  * *! ** *!  !* !!!!! *!*!!!! !!!*!

logo
NTDB id 1169879 ACKWRP RS23720 WP 000941264.1  FVTIKDGVIFMADKKQKRDLIESNTYREKMNHLQLEKELVYSTYQQLYTWFETIRNHKEVEQLG....   779
NTDB id 354 BSU 27620 NP 390640.1              FVKIENGVLSVVSGAKKRDLTDSQTYQAKQQLMELDQKLNYSSAEELKEWLNKLMKQDSEAYESTRRT   786
consensus                                      !! !  !!* *     !!!! *!*!!  ! *** !*  ! !!*   !  !   *  *       ****
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X non conserved

X similar

X $\geq$ 50% conserved