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logo
NTDB id 391337 F9800 RS28995 WP 000941232.1  MLQPKTRWKEKEYNDERVSELASKLQLSPLVVSLFLGRGLDTEDKILDFLNTENQEFHDPFLLEGMDRTVERVNKAIQNG   80
NTDB id 354 BSU 27620 NP 390640.1            MLASKMRWEIQRPDQDKVKSLTEQLHITPLVASLLVKRGFDTAESARLFLHTKDADFYDPFEMKGMKEAADRIKQAISQQ   80
consensus                                    !! *! !!     *  **!  !   !***!!! !! * !! !! *    !!*!   *!*!!! * !!     *!*  !! *

logo
NTDB id 391337 F9800 RS28995 WP 000941232.1  EQILIFGDYDADGVSSTTVLYLALQELGAEVEFYIPNRFTEGYGPNEEAFRWAHSAGFSLIITVDTGIAAVHEANVAKEL  160
NTDB id 354 BSU 27620 NP 390640.1            EKIMIYGDYDADGVTSTSVMLHTLQKLSAQVDFYIPDRFKEGYGPNEQAFRSIKERGFSLIITVDTGIAAVHEAKVAKEL  160
consensus                                    ! !*!*!!!!!!!!*!!*!* * !! ! ! !*!!!! !! !!!!!!! !!!  *  !!!!!!!!!!!!!!!!!! !!!!!

logo
NTDB id 391337 F9800 RS28995 WP 000941232.1  GIDLIITDHHEPPPELPEALAIIHPKLEGGLYPFHYLAGVGVAFKVAHALLGRVPEHLLEIAVIGTVADLVSLHGENRLL  240
NTDB id 354 BSU 27620 NP 390640.1            GLDVIITDHHEPGPELPDVRAIVHPKQPGCTYPFKELAGVGVAFKLAHALLGELPDELLDLAAIGTIADLVPLHDENRLI  240
consensus                                    !*!*!!!!!!!!*!!!!*  !!*!!! *!* !!!* !!!!!!!!!*!!!!!! *!**!!**! !!!*!!!!*!! !!!!*

logo
NTDB id 391337 F9800 RS28995 WP 000941232.1  VKRGLKHMRMTKNIGLKALFKVANVSQSEITEESIGFSIAPRINAVGRLEDATPAVHLLLSDDPEEAKELAEEIDELNKL  320
NTDB id 354 BSU 27620 NP 390640.1            ATLGLERLRRTNRLGLKELIKLSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEAEELAAEIDQLNKE  320
consensus                                       !! **! !  *!!! ! !*      !  !!**!! *!!!*!!!!!*! ! !!!!!!*!*!* !! !!! !!! !!!

logo
NTDB id 391337 F9800 RS28995 WP 000941232.1  RKDIVKQITEEAIAEVENNFPPEENKVLVLAKEGWNPGVIGIVASKLVERFYRPTIVLCIDPIKETAKGSARSIAGFDLF  400
NTDB id 354 BSU 27620 NP 390640.1            RQKMVSKMTDEAIEMVEQQG..LDQTAIVVAKAGWNPGVVGIVASKLVDRFYRPAIVLGIDEEKGIAKGSARSIRGFNLF  398
consensus                                    !  *!  *!*!!!  !!** ** **  *!*!! !!!!!!*!!!!!!!!*!!!!! !!!*!!* !  !!!!!!!! !! !!

logo
NTDB id 391337 F9800 RS28995 WP 000941232.1  ANLSDCRELLPHFGGHPMAAGMTLHMNDVDELRRRLNEQADAILTEEDFIPITAVDVFCKVEDVTLAAIEDMQKLAPFGV  480
NTDB id 354 BSU 27620 NP 390640.1            ESLSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTLTEEDFIPVQEVDLVCGVEDITVESIAEMNMLSPFGM  478
consensus                                      !!*!!**!!!!!!!!!!!!!!!!*  !!**!! !!!! !!  !!!!!!!!*  !!* ! !!!*!*  ! *!* ! !!!*

logo
NTDB id 391337 F9800 RS28995 WP 000941232.1  GNPKPRIAVKDAELESIRAIGSDGSHLKMALRDGQATLDTIGFGFGAYAKEISPVAKVSVIGEASINEWNNFKKPQLMVQ  560
NTDB id 354 BSU 27620 NP 390640.1            LNPKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGELQEGIVPGSRISIVGEMSINEWNNRKKPQLMIK  558
consensus                                     !!!!** !  ! !! *! !!   *!*!! *!     !!**!!  !     ! !  **!**!! !!!!!!! !!!!!!*
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logo
NTDB id 391337 F9800 RS28995 WP 000941232.1  DIAVEAWQLFDWRSMRNAEANLAELPKEKITMVYFSEEVLNKFSLEDYKEKLMH.....ASEVTHLDDQYIVLLDLPNGT   635
NTDB id 354 BSU 27620 NP 390640.1            DAAVSEWQLFDLRGKRTWEDTVSALPSAKRAIVSFKEDSTTLLQTEDLRREVHVISSKDQAKAFDLDGAYIVLLDPPPSL   638
consensus                                    ! !!  !!!!! !  !  !  *  !!  !  *! ! !*       !! *  ********     *!!  !!!!!!*!*

logo
NTDB id 391337 F9800 RS28995 WP 000941232.1  DELRELFKVGFPARIYTLFYQENNHLFSTVPTREHFKWYYSFLSQKTPFSLRQYGEQLCRHKGWSKDTVNFMTQVFFELE   715
NTDB id 354 BSU 27620 NP 390640.1            DMLARLLEGKAPERIYFIFLNHEDHFLSTFPARDHFKWYYAFLLKRGAFDVKKHGSELAKHKGWSVETINFMTKVFFDLG   718
consensus                                    ! !  !     ! !!! *! **  !  !! ! !*!!!!!! !!  * *! ** *!  !**!!!!! *!*!!!! !!!*!

logo
NTDB id 391337 F9800 RS28995 WP 000941232.1  FVTIKDGVIFMADKKQKRDLIESNTYREKMNHLQLEKELVYSTYQQLYTWFETIRNHKEVEQLG....   779
NTDB id 354 BSU 27620 NP 390640.1            FVKIENGVLSVVSGAKKRDLTDSQTYQAKQQLMELDQKLNYSSAEELKEWLNKLMKQDSEAYESTRRT   786
consensus                                    !! !  !!* *     !!!! *!*!!  ! *** !*  ! !!*   !  !   *  *       ****
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X non conserved

X similar

X ≥ 50% conserved