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logo

NTDB id 38794 BATR1942 RS13145 WP 003327153.1  MYKNINVYKYLSLVIIILSFVYVIYLSYISVFNIFVGADVKINEQGKLEVTNVIDYTDEYYSGVKKGDIILEVNGKKGSD   80
NTDB id 91 BSU 31690 NP 391047.2               ...MKNLIKKFTIAVIVLSILYISYTTYISMNGIIIGTKIHKNDKSQFMIEEISESSYGQFVGLRQGDIILKINKEKPSD   77
consensus                                      ***  !* !  ** *!*!! *!* ! *!!!*  ! *!  ** !*    *  * * *  * !** !!!!! *!  !*!!

logo

NTDB id 38794 BATR1942 RS13145 WP 003327153.1  VQLERGRLVNVESLTVERDNKSFRLQNVSLI...NEENLFMYLIPVISYSICLFCIFFVYRINKIRKSTSAYILILFLLF   157
NTDB id 91 BSU 31690 NP 391047.2               KHLKWGYLSHINSLDILRSGKKIHLKDFDLVTLNRPYSFFLFVLPLFFYFLSIICIFYILKVNKKRRSFAAYILILLLLD   157
consensus                                       *!  ! ! ** !! * !  !  *!    !**** *   !****!*  ! *** !!!** **!! !*!  !!!!!! !!

logo

NTDB id 38794 BATR1942 RS13145 WP 003327153.1  VSVAYVSACGARRGEEVSTYLLVLTLVSCPILYIHFIYKYFTELGTKLFEKKILIIMYMLPLINLLLEDVLSNRLKSSAF   237
NTDB id 91 BSU 31690 NP 391047.2               ISIAYISAGGPFRGHIINRYINLFTFISSPILYLQFIQRYLGEIGKTFLNRI..SFLYIIPIFNLGIEFFQDYLQVDIDF   235
consensus                                      *!*!!*!!*!* !!* *  !* * ! *!*!!!!**!! *!  !*!     * **  *!**!* !! *!             !

logo

NTDB id 38794 BATR1942 RS13145 WP 003327153.1  LSNLNLIPFFVLVLTVSILIHQGIRKIKQNEQKSILKVLAYTNILSFSPFIIFFIIPYVLFDTYVSPFS.LAAFTLVIPF   316
NTDB id 91 BSU 31690 NP 391047.2               LATLNLVSFATLTLFSFSAIYLHLNKYKYAEHSFILKLLILTNTLSFAPFLIFFVLPIIFTGNYIFPALASASLLVLIPF   315
consensus                                      !  !!!**!  ! !     !* ** ! !  !*  !!!*!  !! !!! !!*!!!**! *    !* !  * !   **!!!

logo

NTDB id 38794 BATR1942 RS13145 WP 003327153.1  SLVYQFMTNRLYNIDFVIGRLKYYGFLAITPTVIVVTIFNYIQKPKDNIYSVKLTIVIYLLMFAVFYFKEIIDFRFKLKR   396
NTDB id 91 BSU 31690 NP 391047.2               GLVYQFVANKMFDIEFILGRMRYYALLAMIPTLLIVGALVLFDVMDIQMNPVRQTVFFFVVMFAVFYFKEVMDFKFRLKR   395
consensus                                       !!!!!* !*** !*!**!!**!!* !!* !!***!       *  ** *!* !*  ***!!!!!!!!!**!!*!*!!!

logo

NTDB id 38794 BATR1942 RS13145 WP 003327153.1  FTEKFNYQDGIFKFTQIIKQASSVDQVLNHFKNTILEVLNIDNACVYDNSKGKITLLSKNNFKDIAYKNYEQYVQKFSGD   476
NTDB id 91 BSU 31690 NP 391047.2               FSEKFNYQDSIFKYTQLMRGVTSLQQVFKELKNTILDVLLVSKAYTFEVIPDHKVIFLDKHEVGPDWNFYQEEFENVTSE   475
consensus                                      !*!!!!!!! !!!*!!***  *!* !!  * !!!!!*!! *  !* ** ** *  *    *   * *  !       * *

logo

NTDB id 38794 BATR1942 RS13145 WP 003327153.1  IGKIIEFKQGFIIKIGERGDRSFLVLCFSSYNTFKLTRDEIAWLQTLAFYTNVSLENVVKIEELMVHLEDLKQQESNPVW   556
NTDB id 91 BSU 31690 NP 391047.2               IGKIIEVNQGFLMKVGERGGSSYVLLCLSNINTPRLTRDEISWLKTLSFYTSVSMENVLHIEELMEHLKDLKQEGTNPIW   555
consensus                                      !!!!!!  !!!**!*!!!!  !***!! !  !!**!!!!!! !! !! !!! !!*!!!**!!!!! !! !!!!  *!!*!
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logo

NTDB id 38794 BATR1942 RS13145 WP 003327153.1  LKKLMFAIEEKQRSDLARDLHDSVLQDLISLKRQCELFLADFKKE.DPCKNEVQDKLVQMNQQMSDVILMTRETCHELRP   635
NTDB id 91 BSU 31690 NP 391047.2               LKKLMFAIEEKQRSGLARDLHDSVLQDLISLKRQCELFLGDFKKDDNPCREEVQDKLVQMNEQMSDVISMTRETCHELRP   635
consensus                                      !!!!!!!!!!!!!! !!!!!!!!!!!!!
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