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logo
NTDB id 1090791 ACL6EQ RS15475 WP 099085219.1  MHRNNNFYKYLSLAFIILSFVYVIYLSYISVFNIFVGADVKINEQGKLEVTKVVDYTDEYYSGVKKGDIILEVNGQKGSE   80
NTDB id 91 BSU 31690 NP 391047.2               ...MKNLIKKFTIAVIVLSILYISYTTYISMNGIIIGTKIHKNDKSQFMIEEISESSYGQFVGLRQGDIILKINKEKPSD   77
consensus                                      ***  !  !  **! !*!! *!* ! *!!!*  ! *!  ** !*    *  * * *  * !** !!!!! *!  !*!*

logo
NTDB id 1090791 ACL6EQ RS15475 WP 099085219.1  IQLESGRLVNVESLTVERDNKSFHLQNVSLIS...EENLFMYLIPLISYSICLFCIFFVYRINKVRKSPSAFVLILFLLF   157
NTDB id 91 BSU 31690 NP 391047.2               KHLKWGYLSHINSLDILRSGKKIHLKDFDLVTLNRPYSFFLFVLPLFFYFLSIICIFYILKVNKKRRSFAAYILILLLD   157
consensus                                       *!  ! ! ** !! * !  !  !!    !******   !****!!  ! *** !!!** **!! !*!* !**!!! !!

logo
NTDB id 1090791 ACL6EQ RS15475 WP 099085219.1  VSVAYVSACGARRGEEVSTYLLVLTLVSCPILYIHFIYKYFAELGTKLFKKKILIIMYILPLVNLLLEAVLSNRLNSSAF   237
NTDB id 91 BSU 31690 NP 391047.2               ISIAYISAGGPFRGHIINRYINLFTFISSPILYLQFIQRYLGEIGKTFLNRI..SFLYIIPIFNLGIEFFQDYLQVDIDF   235
consensus                                      *!*!!*!!*!* !!* *  !* * ! *!*!!!!**!! *! *!*!     * **  *!!*!* !! *!            !

logo
NTDB id 1090791 ACL6EQ RS15475 WP 099085219.1  LSNLNLISFFVLVLTVSILIHQGIRKIKQNEQKSILKILAYINILSFSPFIIFFVIPYVLFDKYLSPFS.LAAFTLVIPF   316
NTDB id 91 BSU 31690 NP 391047.2               LATLNLVSFATLTLFSFSAIYLHLNKYKYAEHSFILKLLILTNTLSFAPFLIFFVLPIIFTGNYIFPALASASLLVLIPF   315
consensus                                      !  !!!*!!  ! !     !* ** ! !  !*  !!!*!   ! !!! !!*!!!!*! *    !* !  * !   **!!!

logo
NTDB id 1090791 ACL6EQ RS15475 WP 099085219.1  SLVYQFMTNRLYNIDFVVGRLKYYGFLAITPTVIVVTIFNYIQKPEDNFYSVKLTVVIYLLMFAVFYFKEIIDFRFKLKR   396
NTDB id 91 BSU 31690 NP 391047.2               GLVYQFVANKMFDIEFILGRMRYYALLAMIPTLLIVGALVLFDVMDIQMNPVRQTVFFFVVMFAVFYFKEVMDFKFRLKR   395
consensus                                       !!!!!* !*** !*!**!!**!!* !!* !!***!        ** *  *!* !!  ***!!!!!!!!!**!!*!*!!!

logo
NTDB id 1090791 ACL6EQ RS15475 WP 099085219.1  FTEKFNYQDGIFKFTQLIKQASSVDQVLNHFKNTILEVLNIDNACVYDNSKGEITLLSKNNFKDLTYKNHEQYVKNFSGD   476
NTDB id 91 BSU 31690 NP 391047.2               FSEKFNYQDSIFKYTQLMRGVTSLQQVFKELKNTILDVLLVSKAYTFEVTPDHKVIFLDKHEVGPDWNFYQEEFENVTSE   475
consensus                                      !*!!!!!!! !!!*!!!**  *!* !!  * !!!!!*!! *  !* ** ** *  *    *   * *  *     ! * *

logo
NTDB id 1090791 ACL6EQ RS15475 WP 099085219.1  IGKIKEFKQGFIIKIGERGDRSFLVLCFSSYNTFKLTRDEIAWLQTLAFYTNVSLENVMKIEELMVHLEDLKQQESNPVW   556
NTDB id 91 BSU 31690 NP 391047.2               IGKIIEVNQGFLMKVGERGGSSYVLLCLSNINTPRLTRDEISWLKTLSFYTSVSMENVLHIEELMEHLKDLKQEGTNPIW   555
consensus                                      !!!! !  !!!**!*!!!!  !***!! !  !!**!!!!!! !! !! !!! !!*!!!**!!!!! !! !!!!  *!!*!
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logo

NTDB id 1090791 ACL6EQ RS15475 WP 099085219.1  LKKLMYTIEEKQRSDLARDLHDSVLQDLISLKRQCELFLGDFKKGDNPCREEVQDKLVQMNEQMSDVISMTRETCHELRP  636
NTDB id 91 BSU 31690 NP 391047.2               LKKLMFAIEEKQRSGLARDLHDSVLQDLISLKRQCELFLGDFKKDDNPCREEVQDKLVQMNEQMSDVISMTRETCHELRP  635
consensus                                      !!!!!* !!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1090791 ACL6EQ RS15475 WP 099085219.1  QLLYDLGLVKALSKLVAQQQERVPFHIRLNTGRFTASLDLDSQLNLYRIIQEFLSNAVKHSQATDVLIMLISIQNKIVLH  716
NTDB id 91 BSU 31690 NP 391047.2               QLLYDLGLVKALSKLVAQQQERVPFHIRLNTGRFTASLDLDSQLNLYRIIQEFLSNAVKHSQATDVLIMLISIQNKIVLH  715
consensus                                      !!!!!!!!!!!!!!!!!!!!
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