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logo
NTDB id 1064362 ACHD8M RS15755 WP 159354531.1  MKSKYNKISVVLILLTLVYVIYLTYISTNNLLVGATISENKQGEIKIKDVEDFTMAYYAGVQKGDIVKNINGQNVKSINI   80
NTDB id 91 BSU 31690 NP 391047.2               MKNLIKKFTIAVIVLSILYISYTTYISMNGIIIGTKIHKNDKSQFMIEEISFSSYGQFVGLRQGDIILKINKEKPSDKHL   80
consensus                                      !!     ! ** *!*!***!* ! !!!! ! ***!  !* !      ! ** * * * * !*  !!!*  !!   *   **

logo
NTDB id 1064362 ACHD8M RS15755 WP 159354531.1  QMNKLKNVNSMVVDRNGHDIEIKL....NLFNDRNFSTYLIPIIFYVVCLFCCFFIIKINEEKNLKSALVLIVFLLSASI  156
NTDB id 91 BSU 31690 NP 391047.2               KWGYLSHINSLDILRSGKKIHLKDFDLVTLNRPYSFFLFVLPLFFYFLSIICIFYILKVNKKRRSFAAYILILLLLDISI  160
consensus                                          ! **!!* * ! !* !**! **** !  *  !  ***!* !! *** !*!*!*!*!  *    ! *!!* !!  !!

logo
NTDB id 1064362 ACHD8M RS15755 WP 159354531.1  AYISAGGSAKGDLLSRCIMLLTLITVPLNYLLFLYHYFKELGTVLFSKKVFVLYLFPTINILFELFRSRLSF.GDFIPKF  235
NTDB id 91 BSU 31690 NP 391047.2               AYISAGGPFRGHIINRYINLFTFISSPILYLQFIQRYLGEIGKTFLNR.ISFLYIIPIFNLGIEFFQDYLQVDIDFLATL  239
consensus                                      !!!!!!!* *!*** !*! ! ! !* !* !! !* *!  !*!     ***  !!* !  !*  ! !   !  * !!**

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NTDB id 1064362 ACHD8M RS15755 WP 159354531.1  NLISFFLLFLVVAYYFIVILRRNKDTDQAHIIKVLVVINILAFSPFLCLYLIPYVFFNDYYVSSFLTASFMLLIPFSLVY  315
NTDB id 91 BSU 31690 NP 391047.2               NLVSFATLTLFSFSAIYLHLNKYKYAEHSFILKLLILTNTLSFAPFLIFFVLPIIFTGNYIFPALASASLLVLIPFGLVY  319
consensus                                      !!*!!  ! !       **! * !  ** *!*!*!** ! ! ! !!!* ***! *!   !  *   *!! **!!!! !!!

logo
NTDB id 1064362 ACHD8M RS15755 WP 159354531.1  QFMSNKIYNVDFILSKLKYYGFLALTPTIILVTTIELLREPEENFYNLKLTFLIYILMLAVFYYKEILDFRFKLKRVSER  395
NTDB id 91 BSU 31690 NP 391047.2               QFVANKMFDIEFILGRMRYYALLAMIPTLLIVGALVLFDVMDIQMNPVRQTVFFFVVMFAVFYFKEVMDFKFRLKRFSEK  399
consensus                                      !!* !!** **!!! ***!!* !!* !!***!  * !   ** *  *** !   ***! !!!!*!!**!!*!*!!! !!*

logo
NTDB id 1064362 ACHD8M RS15755 WP 159354531.1  FNYQDSIYKFTQLIRNSSSLNQVFGELKTTILEVTLTSKVYIFEVTYDGNVKAFDERDSS.LIDDYQVLLNSVTVEIGKI  474
NTDB id 91 BSU 31690 NP 391047.2               FNYQDSIFKYTQLMRGVTSLQQVFKELKNTILDVLLVSKAYTFEVTPDHKVIFLDKHEVGPDWNFYQEEFENVTSEIGKI  479
consensus                                      !!!!!!!*!*!!!*!  *!!*!!! !!! !!!*! ! !! ! !!!!*!* !   ! **  *    !!      !! !!!!!

logo
NTDB id 1064362 ACHD8M RS15755 WP 159354531.1  TEFNQGFAMKVGERGDRSFVVLCLSKMNTPRLSREEISWLKTLSFYVNVSIENVMKIEELMNHLEDLKKQQGSNPVWLKK  554
NTDB id 91 BSU 31690 NP 391047.2               IEVNQGFLMKVGERGGSSYVLLCLSNINTPRLTRDEISWLKTLSFYTSVSMENVLHIEELMEHLKDLK.QEGTNPIWLKK  558
consensus                                       ! !!!! !!!!!!!  !*!*!!!! *!!!!!*!*!!!!!!!!!!!  !!*!!!**!!!!! !! !!!*! !*!!*!!!!
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logo

NTDB id 1064362 ACHD8M RS15755 WP 159354531.1  LMFTIEEKQRSDLARDLHDSVLQDLISLKRQCELFLADFKK.EEPCQLEVQDKLHQMNEQMSDVILMTRETCHELRPQLL   633
NTDB id 91 BSU 31690 NP 391047.2               LMFAIEEKQRSGLARDLHDSVLQDLISLKRQCELFLGDFKKDDNPCREEVQDKLVQMNEQMSDVISMTRETCHELRPQLL   638
consensus                                      !!! !!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!** !!   !!!!!!*!!!!!!!!!!! !!!!!!!!!!!!!!

logo

NTDB id 1064362 ACHD8M RS15755 WP 159354531.1  YDLGLVKAVSKLAAQQQERAPFHIRLNTGRFTAALDLDTQLNLYRIIQEFLSNAMKHSKANEVLIMLISIQNKVILHYED   713
NTDB id 91 BSU 31690 NP 391047.2               YDLGLVKALSKLVAQQQERVPFHIRLNTGRFTASLDLDSQLNLYRIIQEFLSNAVKHSQATDVLIMLISIQNKIVLHYED   718
consensus                                      !!!!!!!!*!!! !!!!!! !!!!!!!!!!!!! !!!!*!!!!!!!!!!!!!!!!*!!! ! *!!!!!!!!!!!!**!!!!!

logo

NTDB id 1064362 ACHD8M RS15755 WP 159354531.1  DGVGCNQEEGGGQSMSMGLSGIKERVRALDGRMKIDTSEGNGFKADIEMEL   764
NTDB id 91 BSU 31690 NP 391047.2               DGVGFDQEKNTEHSMSMGLSGIKERVRALDGRLRIETSEGKGFKADIEIEL   769
consensus                                      !!!!* !!     *!!!!!!!!!!!!!!!!!!!!**!*!!!! !!!!!!!*!!
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X non conserved
X similar
X ≥ 50% conserved