```
logo
NTDB id 1198 PSJM300 03950 AFN76868.1          MAQKAIKNSVFTWEGLDRQGAKIKGELSGVSPALVKAQLRKQGINPQKVRKKSISL.......FSAGKKIKPMDIALFTR   73
NTDB id 391143 DAAJ005 RS08405 WP 151846721.1 .......MAVFEYRARDRSGKILKSQMEAETEAQVRDTLRAKNLMIVEIKAPKSGLNADLKIPFLDDRPPSLKQVSIFSK   73
NTDB id 1029 TT RS02230 WP 011228203.1         .......MPVYQYKARDRQGRLVEATIEAEDLRTAARLLRDRGLFVAEIKEPGKGLQAEVRIPALE.RGPGLKDLAIFSR   72
consensus                                              * !* ****!!*!***** ***** * **  !! *** * *** *  *!*********  * * ******!**

logo
NTDB id 1198 PSJM300 03950 AFN76868.1          QMATMMKAGVPLLQSFDIIGEGFDNPNMRKLVDDLKQEVAAGNSFASSLRKKPQYFDDLYCNLVDSGEQSGSLETLLDRV  153
NTDB id 391143 DAAJ005 RS08405 WP 151846721.1 QLATLINAGVPLVQSLAVLQGQIEHKGFQKTVKSLRSEVEAGTPLSEAIAKYPKVFNRLYVNLVRAGETSGTLDSVLERI  153
NTDB id 1029 TT RS02230 WP 011228203.1         QLATMLGAGLTLLQALAILERQTENRKFREILKQVRTDVEGGMAFSEALSK.HKIFSRLYVNLVRAGETSGGLDLILDRL  151
consensus                                      !*!!** !!**!*!*****  * *** ****** ****!**!  ***** ! ***! *!!*!!!**!!*!!*!***!*!*

logo
NTDB id 1198 PSJM300 03950 AFN76868.1          ATYKEKTEALKAKIKKAMNYPIAVVLVAVIVTAILLIKVVPQFQDVFANFGAELPAFTLMVIGLSEALQA..WWYVVLAG  231
NTDB id 391143 DAAJ005 RS08405 WP 151846721.1 AAFQEKELILRGKIKSALTYPVVVLVFAILLITYFLLTTIVPQFAGILAQLNAPLPFITKMLMAVSEFLQHSAWLLVVFAV  233
NTDB id 1029 TT RS02230 WP 011228203.1         ASFLEKELELRGKIRSAMTYPVIVFVFAVGVAYFLLTGIVPQFAQILTDLGSELPLLTRFLIAVSDLLRAATLPLLLLAV  231
consensus                                      !** !!** !**!!**!**!!**!***!******!!* *!!!!* ********!! *!******!* !**  * ****!*

logo
NTDB id 1198 PSJM300 03950 AFN76868.1          FVGAAYAFKQAHTRSEKFRNWFDRLMLKVPVVGDILYKSSVARFARTLSTTFAAGVPLVDALDSVAGATGNVVFRSATNK  311
NTDB id 391143 DAAJ005 RS08405 WP 151846721.1 IITFAYRWVYK...MPKGRVIIDDIKLKLPVFGNLTQKSAIASFARTFGLLISSGVNIIEALEITKGTADNAIVEETIEN  310
NTDB id 1029 TT RS02230 WP 011228203.1         ALFFAYRWYYG...TPQGRRVIDRLKLRLPVFGNLNRKTAVARFSRTLALLLSSGVNIVEALDITKGTAGNSVVEEIVEA  308
consensus                                       * *!!** **   ****! **!***!**!!*!**  !***!*!*!!*******!!****!!****!***!  **** **

logo
NTDB id 1198 PSJM300 03950 AFN76868.1          IKSDVSSGMQLNFSMR.TTGTFPSMAVQMTAIGEESGALDEMLSKVATFYEDEVDNMVDGLTALMEPMIMAVLGVLVGGL  390
NTDB id 391143 DAAJ005 RS08405 WP 151846721.1 AKNVVMVGEQMSSSLA.TSKVFPPMVVSMISIGEETGALDSMLSKVGDFYDREVDEAVESMTAAIEPIMIVFLGGIVGTI  389
NTDB id 1029 TT RS02230 WP 011228203.1         AKLKIQQGDPLNLTLAQHPFVFPPMVSSMVAIGEETGALDTMLSKVADFYEREVDEAVASLTAAIEPLMIIFLGVIVGMI  388
consensus                                      *!  *  !*** *** ** *!!*!***!*!!!!*!!!!*!!!!!**!!**!!!**!***!!**!!*****!!**!!  *

logo
NTDB id 1198 PSJM300 03950 AFN76868.1          IIAMYLPIFQMGSVV...   405
NTDB id 391143 DAAJ005 RS08405 WP 151846721.1 VAGMFLPMFSIIGTLSQ.   406
NTDB id 1029 TT RS02230 WP 011228203.1         VAGMFLPLFKIIGTLSVQ   406
consensus                                      ***!*!!*! ******
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X non conserved

X similar

X ≥ 50% conserved