```
logo
NTDB id 153 SP RS04730 WP 000387344.1          .....MEAIIEKIKEYKIIVICTGLGLLVG..GFFLLKPAPQTPVKETNLQAEVAAVSKDSSTEKEVKKEEKEEPLEQDL   73
NTDB id 1078156 ACKFA7 RS12070 WP 038459268.1 MTKAMMTEPL..RRNLKILGACLVAGAVFIC..IWLFRSEKEEPVKQQSVLGQTA...........SASVADKKDTADDK   65
NTDB id 105 BSU 25590 NP 390437.1              .....MNWLN..QHKKAIIL..AASAAVFTAIMIFLATGKNKEPVKQAVPTETEN..........TVVKQEANNDESNET   61
consensus                                           !  **   *  *!** * * ****    **! *    *!!!*  * *  *          * * *   *  **

logo
NTDB id 153 SP RS04730 WP 000387344.1          ITVDVKGAVKSPGIYDLPVGSRVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEAVSQQTGSGTASST.SKE   152
NTDB id 1078156 ACKFA7 RS12070 WP 038459268.1 IMIDVKGAVRNPGVYEMKAGDRVTQAIEKAGGLKKKADELKVNLAEQLQDGTIVYIPSEGEEENRPKTAAG.....EKEN   140
NTDB id 105 BSU 25590 NP 390437.1              IVIDIKGAVQHPGVYEMRTGDRVSQAIEKAGGTSEQADEAQVNLAEILQDGTVVYIPKKGEETAVQQGGGGSVQSDGGKG   141
consensus                                      !**!*!!!!* !!*!*** !*!!**!**!!!!*****!!*  *!!!* **!**!!*!**!!!    *****!*  *   *

logo
NTDB id 153 SP RS04730 WP 000387344.1          KKVNLNKASLEELKQVKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGGKTIEKLKDYVTVD   216
NTDB id 1078156 ACKFA7 RS12070 WP 038459268.1 AAVNINTASLDELQAISGVGQKKAEAIIAYREEKGRFQTAEDLMNVSGFGEKSFERIKTSITVK   204
NTDB id 105 BSU 25590 NP 390437.1              ALVNINTATLEELQGISGVGPSKAEAIIAYREENGRFQTIEDITKVSGIGEKSFEKIKSSITVK   205
consensus                                      * !!*!*!*!*!!*****!*! **!**!!**!!**!*!******* *!!!*!*!**!**!****!!*
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X non conserved

X similar

X ≥ 50% conserved