```
logo
NTDB id 36 MW RS08230 WP 001793467.1          MVLLYQFLLRYKDFLTQWKLYIISA...VVLIMVLIGFIFWRQDDYTSRNFENKDNALKQSTSENSSLSKVEDVQVKDGD    77
NTDB id 38783 BATR1942 RS10540 WP 003325391.1 ...........MNWFYQHKKIILTAGAAVIFIVILL...NFPGA....KK....QEPVQQITAS..PETAGTRVQQQESK    56
NTDB id 105 BSU 25590 NP 390437.1             ...........MNWLNQHKKAIILAASAAVFTAIMI...FLATG....KN....KEPVKQAV....PTETENTVVKQEAN    54
consensus                                                 **** !*!* !**!* *****+***   **  *    **     ****! *    * *    !* ***

logo
NTDB id 36 MW RS08230 WP 001793467.1          NSKNKGPVYVDVKGAVKHPNVYKMTSKDRVVDLLDKAQ.LLEDADVSQINLSEKLTDQKMIFIPHKGQKNVEPQIGVNSV   156
NTDB id 38783 BATR1942 RS10540 WP 003325391.1 GDADGEKIIIDIKGAVKHPGVYELKTGDRVSQAIEKAGGINSEADEKQVNLAELLQDGTVVYIPSEGEEPTQSTTANASV   136
NTDB id 105 BSU 25590 NP 390437.1             NDESNETIVIDIKGAVQHPGVYEMRTGDRVSQAIEKAGGTSEQADEAQVNLAEILQDGTVVYIPKKGEETAVQQGGGGSV   134
consensus                                     **   * ***!*!!!!*!!*!!***!!!*****!!*** **!!* !*!!*!*!*!*****!! *!**    * * *!!

logo
NTDB id 36 MW RS08230 WP 001793467.1          HVKNGNTNNTKVNLNTASVSELMSVPGVGQAKANAIVEYRNQQGAFQEIDDLKKVKGFGSKTFDKLKSYFTI.   228
NTDB id 38783 BATR1942 RS10540 WP 003325391.1 ..QNGAGKEALVNINTASLEELQAISGVGPSKAEAIIAYREENGKFQAVEDITNVSGIGEKSFEKIKSAITVK   207
NTDB id 105 BSU 25590 NP 390437.1             ..QSDGGKGALVNINTATLEELQGISGVGPSKAEAIIAYREENGRFQTIEDITKVSGIGEKSFEKIKSSITVK   205
consensus                                       ****** **!!*!!!***!!*****!!!**!!*!!**!!***!*!! **!***!*!*!*!*!*!*!! *!**
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X non conserved

X similar

X ≥ 50% conserved