```
logo
NTDB id 1013803 ABVR74 RS00780 WP 025017137.1  .........MKNEILMTRAFTLLESLLVLLIISFITTLFSSEIIQTVHLFKGELFVLQFENFYKRSQEEAALLQKSESLV   71
NTDB id 589 KW2 RS10840 WP 096816401.1         MMITMTRIKMNSAISMIRAFTLLESLLVLLIVSFITLFFSAELTQTVHLFKGELFVLQFENLYKISQENAALQSSPENLG   80
NTDB id 397 FSA28 RS09390 WP 002263438.1       .............MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLI   67
NTDB id 388 SMU RS09020 WP 002263438.1        .............MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLI   67
NTDB id 432 SGO RS09405 WP 012130939.1        ....MVRTIVRLRQLPIKAFTVLESLLVLMISSFILLALSSSVQATFEQIQAKIFFLEFEHFYQESQKLSVSSQRKLVLE   76
consensus                                               *   ** **!!!**!!!**!*!*!!*** *!*****!*  * ***!*! !!**!***!**** *   **!*

logo
NTDB id 1013803 ABVR74 RS00780 WP 025017137.1  AKNQELICEDRSITIPKEVAVKD.FTVKFDDKGENSSLQKLTISLPYEKKFITYQLEIGSGKFKKKIS   138
NTDB id 589 KW2 RS10840 WP 096816401.1         SKNGKLIYENKEIDIPKEVEMVE.FLIKFDEKGENSSLQKIKVYLPYEKKTILYQMEMGSGKYKKKIN   147
NTDB id 397 FSA28 RS09390 WP 002263438.1       LKSEYISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQT.SDEKRVTYQLYIGSGQYKKTES   134
NTDB id 388 SMU RS09020 WP 002263438.1        LKSEYISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQT.SDEKRVTYQLYIGSGQYKKTES   134
NTDB id 432 SGO RS09405 WP 012130939.1        ISSQEISNGYARLPIPKGIQAPESTQIYFDKAGGNSSLSKVQFQT.K.EGLVTYQLYIGNGKFKKTTN   142
consensus                                     ***  ****   * !!!** * *  ** !! *!*!!!! !* ***  *** **!!***!*!**!!* *
```

X non conserved

X similar

X $\geq$ 50% conserved