```
logo
NTDB id 1013070 ABQ274 RS10545 WP 023164614.1  ................MTKAFTLLESLLVLLITSFITTLFSLEIIQTIHLFKGELFVLQFENFYKRSQEDAALLQKSESLV   65
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 1013070 ABQ274 RS10545 WP 023164614.1  AKNQELICEDRSITIPKEVAVKD.FTVKFDDKGENSSLQKLTISLPYEKKFITYQLEIGSGKFKKKIS   132
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                                      ***  ****   * !!!** * *  ** !! *!*!!!! !* ***  *** **!!***!*!**!!* *
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X non conserved

X similar

X $\geq$ 50% conserved