```
logo
NTDB id 256733 CV702 RS10880 WP 026138965.1  ....MIRIKMKNEILMTRAFTLLESLLVLLIISFITTLFSLEIIQTVHLFKGELFVLQFENFYKRSQEAAALLQKSESLV   76
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 256733 CV702 RS10880 WP 026138965.1  AKNQELICEDRSITIPKEVAVKD.FTVKFDDKGGNSSLQKLTIFLPYEKKFITYQLEIGSGKFKKKIS   143
NTDB id 589 KW2 RS10840 WP 096816401.1       SKNGKLIYENKEIDIPKEVEMVE.FLIKFDEKGENSSLQKIKVYLPYEKKTILYQMEMGSGKYKKKIN   147
NTDB id 397 FSA28 RS09390 WP 002263438.1     LKSEYISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLV.FQTSDEKRVTYQLYIGSGQYKKTES   134
NTDB id 388 SMU RS09020 WP 002263438.1       LKSEYISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLV.FQTSDEKRVTYQLYIGSGQYKKTES   134
NTDB id 432 SGO RS09405 WP 012130939.1       ISSQEISNGYARLPIPKGIQAPESTQIYFDKAGGNSSLSKVQ.FQTK.EGLVTYQLYIGNGKFKKTTN   142
consensus                                    ***  ****   * !!!** * *  ** !! *!*!!!! !*  *** *** **!!***!*!**!!* *
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X non conserved

X similar

X ≥ 50% conserved