```
logo
NTDB id 58846 QML87 RS04615 WP 001289493.1  MSEIGFKYSILASGSSGNSFYLETSKKKLLVDAGLSGKKITSLLAEINRKPEDLDAILITHEHSDHIHGVGVLARKYGMDLYANE   85
NTDB id 383 SMU RS06875 WP 002262928.1      MTETGFRYSILASGSSGNSFYLETPQKRILIDAGLSGKKISQLMAEIDRKVDDVDAILVTHEHKDHIHGVGVLARKYGLDVYANA   85
consensus                                   !*! !!*!!!!!!!!!!!!!!!!!* !**!*!!!!!!!!!* !*!!! !!**!*!!!!*!!!! !!!!!!!!!!!!!!*!*!!!

logo
NTDB id 58846 QML87 RS04615 WP 001289493.1  KTWQAMENSKYLGKVDSSQKHIFEMGKTKTFGDIDIESFGVSHDAVAPQFYRFMKDDKSFVLLTDTGYVSDRMAGIVENADGYLI   170
NTDB id 383 SMU RS06875 WP 002262928.1      ATWQAMDG..MIGKIDVSQKHIFDLGKTITFGDIDIESFGVSHDAAQPQFYRLMKDGKSFVILTDTGYVSDRMADLIENADGYLI   168
consensus                                    !!!!!* ** *!!*! !!!!!!**!!! !!!!!!!!!!!!!!!!  !!!!! !!! !!!!*!!!!!!!!!!!! **!!!!!!!!

logo
NTDB id 58846 QML87 RS04615 WP 001289493.1  EANHDVEILRSGSYAWRLKQRILSDLGHLSNEDGAEAMIRTLGNRTKKIYLGHLSKENNIKELAHMTMVNQLAQADLGVGVDFKV   255
NTDB id 383 SMU RS06875 WP 002262928.1      ESNHDIEILRSGIYPWSLKQRILSDKGHLSNDDGAETMVRTLGNKTKKIYLGHLSKENNVKELAHMTMKNALMQADLAVDHDFKI   253
consensus                                   ! !!!*!!!!!! !*! !!!!!!!! !!!!!*!!!! !*!!!!!*!!!!!!!!!!!!!!!*!!!!!!!! ! ! !!!!*! *!!!*

logo
NTDB id 58846 QML87 RS04615 WP 001289493.1  YDTSPDTATPLTEI   269
NTDB id 383 SMU RS06875 WP 002262928.1      YDTSPDTALPLAKI   267
consensus                                   !!!!!!!! !!  !
```

X non conserved

X similar

X $\geq$ 50% conserved