```
logo
NTDB id 1136749 DQL47 RS07360 WP 021320992.1  MSEVGFKYSILASGSTGNCFYLETPKKRLLIDAGLTGKKITSLLAEIDRRPEDLDAILITHEHSDHIKGVGIMARKYNLD   80
NTDB id 383 SMU RS06875 WP 002262928.1        MTETGFRYSILASGSSGNSFYLETPQKRILIDAGLSGKKISQLMAEIDRKVDDVDAILVTHEHKDHIHGVGVLARKYGLD   80
consensus                                     !*! !!*!!!!!!!!*!!*!!!!!! !!*!!!!!!*!!!!* !*!!!!!***!*!!!!*!!!! !!!*!!!**!!!! !!

logo
NTDB id 1136749 DQL47 RS07360 WP 021320992.1  VYANEKTWQMIDERHMIGNLDVTQKHYFERGKTMTFGDIDIESFGVSHDAVDPQFYRIMKDNKSFVMLTDTGYVSDRLAG  160
NTDB id 383 SMU RS06875 WP 002262928.1        VYANAATWQAMD..GMIGKIDVSQKHIFDLGKTITFGDIDIESFGVSHDAAQPQFYRLMKDGKSFVILTDTGYVSDRMAD  158
consensus                                     !!!!  !!! *!***!!! *!!*!!!*!* !!!*!!!!!!!!!!!!!!!!  !!!!!*!!! !!!!*!!!!!!!!!!*!

logo
NTDB id 1136749 DQL47 RS07360 WP 021320992.1  IIENADGYLIESNHDVEILRAGSYPWSLKQRILSDLGHLSNEDGAGAMIRSLGNKTKKIYLGHLSQENNIKELAHMTMVN  240
NTDB id 383 SMU RS06875 WP 002262928.1        LIENADGYLIESNHDIEILRSGIYPWSLKQRILSDKGHLSNDDGAETMVRTLGNKTKKIYLGHLSKENNVKELAHMTMKN  238
consensus                                     *!!!!!!!!!!!!!!!*!!!! ! !!!!!!!!!!!! !!!!!*!!!  !*!*!!!!!!!!!!!!!! !!!*!!!!!!!! !

logo
NTDB id 1136749 DQL47 RS07360 WP 021320992.1  QLAMADLGVGTDFTVHDTSPDTACPLTDI  269
NTDB id 383 SMU RS06875 WP 002262928.1        ALMQADLAVDHDFKIYDTSPDTALPLAKI  267
consensus                                      !  !!!*! *!! **!!!!!!!*!!  !
```

X non conserved

X similar

X $\geq$ 50% conserved