```
logo
NTDB id 266948 C2E56 RS03845 WP 001289498.1  MSEIGFKYSILASGSTGNCFYLETPQKRLLIDAGLTGKKVTSLLAEINRKPEDLDAILVTHEHSDHIKGVGVLARKYHLD   80
NTDB id 383 SMU RS06875 WP 002262928.1       MTETGFRYSILASGSSGNSFYLETPQKRILIDAGLSGKKISQLMAEIDRKVDDVDAILVTHEHKDHIHGVGVLARKYGLD   80
consensus                                    !*! !!*!!!!!!!!*!!*!!*!!!!!!*!!!!!!*!!!** !*!!! !!**!*!!!!!!!!! !!!*!!!!!!!!!!*!!

logo
NTDB id 266948 C2E56 RS03845 WP 001289498.1  IYANEQTWKVMDERNMLGKVDVSQKHVFGRGKTLTFGDLDIESFGVSHDAVDPQFYRMMKDDKSFVMLTDTGYVSDRMAS  160
NTDB id 383 SMU RS06875 WP 002262928.1       VYANAATWQAMD..GMIGKIDVSQKHIFDLGKTITFGDIDIESFGVSHDAAQPQFYRLMKDGKSFVILTDTGYVSDRMAD  158
consensus                                    *!!!  !!  !!** !*!!*!!!!!!*!  !!!*!!!!*!!!!!!!!!!!!  !!!!!*!!! !!!!*!!!!!!!!!!!!!

logo
NTDB id 266948 C2E56 RS03845 WP 001289498.1  LIENADGYLIESNHDIEILRSGSYPWTLKQRILSDKGHLSNEDGSETMYRTLGNRTKKIYLGHLSKENNYKELAHMTMEN  240
NTDB id 383 SMU RS06875 WP 002262928.1       LIENADGYLIESNHDIEILRSGIYPWSLKQRILSDKGHLSNDDGAETMVRTLGNKTKKIYLGHLSKENNVKELAHMTMKN  238
consensus                                    !!!!!!!!!!!!!!!!!!!!!!! !!!*!!!!!!!!!!!!!!!*!! !!!*!!*!!*!!*!!!!!!!!!!!*!!!!!!!! !

logo
NTDB id 266948 C2E56 RS03845 WP 001289498.1  NLMRADFGVGTDFSVHDTSPDSATPLTRI   269
NTDB id 383 SMU RS06875 WP 002262928.1       ALMQADLAVDHDFKIYDTSPDTALPLAKI   267
consensus                                     !! !! *! *!! **!!!!!*! !! *!
```

X non conserved

X similar

X ≥ 50% conserved