```
logo
NTDB id 144630 SAH002 RS03910 WP 001289497.1  MSEIGFKYSILASGSTGNCFYIETPQKRLLIDAGLTGKKVTSLLAEINRKPEDLDAILVTHEHSDHIKGVGVLARKYHLD   80
NTDB id 383 SMU RS06875 WP 002262928.1        MTETGFRYSILASGSSGNSFYLETPQKRILIDAGLSGKKISQLMAEIDRKVDDVDAILVTHEHKDHIHGVGVLARKYGLD   80
consensus                                     !*! !!*!!!!!!!!*!!*!!*!!!!!!*!!!!!!*!!!** !*!!! !!**!*!!!!!!!!! !!!*!!!!!!!!!*!!

logo
NTDB id 144630 SAH002 RS03910 WP 001289497.1  IYANEQTWKVMDERNMLGKVDVSQKHVFGRGKTLTFGDLDIESFGVSHDAVDPQFYRMMKDDKSFVMLTDTGYVSDRMAG   160
NTDB id 383 SMU RS06875 WP 002262928.1        VYANAATWQAMD..GMIGKIDVSQKHIFDLGKTITFGDIDIESFGVSHDAAQPQFYRLMKDGKSFVILTDTGYVSDRMAD   158
consensus                                     *!!!  !!  !!** !*!!*!!!!!!*!  !!!*!!!!*!!!!!!!!!!!  !!!!!*!!! !!!!*!!!!!!!!!!!!!

logo
NTDB id 144630 SAH002 RS03910 WP 001289497.1  LIENADGYLIESNHDIEILRSGSYPWTLKQRILSDKGHLSNEDGSETMIRTIGNRTKHIYLGHLSKENNIKELAHMTMEN   240
NTDB id 383 SMU RS06875 WP 002262928.1        LIENADGYLIESNHDIEILRSGIYPWSLKQRILSDKGHLSNDDGAETMVRTLGNKTKKIYLGHLSKENNVKELAHMTMKN   238
consensus                                     !!!!!!!!!!!!!!!!!!!!!!! !!!*!!!!!!!!!!!!!!!*!! !!!*!!*!!*!!*!!!!!!!!!!!!*!!!!!!!! !

logo
NTDB id 144630 SAH002 RS03910 WP 001289497.1  NLMRADFGVGTDFSVHDTSPDSATPLTRI   269
NTDB id 383 SMU RS06875 WP 002262928.1        ALMQADLAVDHDFKIYDTSPDTALPLAKI   267
consensus                                      !! !! *! *!! **!!!!!*! !! *!
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X non conserved

X similar

X ≥ 50% conserved