```
logo
NTDB id 302615 DS740 RS13810 WP 074794436.1  MMNSRLLLPMAAASATAGITAAAYFPAIFLFILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYQF   80
NTDB id 107 BSU 25570 NP 390435.1            MRNSRLLLPMAAASATAGITAAAYFPAIFLFILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYQF   80
consensus                                    ! !!!!
```

```
logo
NTDB id 302615 DS740 RS13810 WP 074794436.1  IKQLDALILTHADQDHIGEAETLLKHHKVKRLVIPKGFVSEPKDEKVLQTAREEGVTIEEVKRGDVLQIKDLQFHVLSPE  640
NTDB id 107 BSU 25570 NP 390435.1            IKQLDALILTHADQDHIGEAEILLKHHKVKRLVIPKGFVSEPKDEKVLQAAREEGVAIEEVKRGDVLQIKDLQFHVLSPE  640
consensus                                    !!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!! !!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 302615 DS740 RS13810 WP 074794436.1  TPDPASKNNSSLVLWMETGVLSWILTGDLEKEGEQEVMDVFPNIKADVLKVGHHGSKGSTGEEFIQQLQPKTAIISAGKN  720
NTDB id 107 BSU 25570 NP 390435.1            APDPASKNNSSLVLWMETGGMSWILTGDLEKEGEQEVMNVFPNIKADVLKVGHHGSKGSTGEEFIQQLQPKTAIISAGKN  720
consensus                                     !!!!!!!!!!!!!!!!!!! *!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 302615 DS740 RS13810 WP 074794436.1  NRYHHPHQKVLQLLQRHSIRVLRTDQNGTIQYRYKNRVGTFSVYPPYDTSDITETN  776
NTDB id 107 BSU 25570 NP 390435.1            NRYHHPHQKVLQLLQRHSIRVLRTDQNGTIQYRYKNRVGTFSVYPPYDTSDITETN  776
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved