```
logo
NTDB id 999402 ABEG41 RS12880 WP 074794436.1  MMNSRLLLPMAAASATAGITAAAYFPAIFLFILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYQF   80
NTDB id 107 BSU 25570 NP 390435.1             MRNSRLLLPMAAASATAGITAAAYFPAIFLFILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYQF   80
consensus                                     ! !!!!
```

```
logo

NTDB id 999402 ABEG41 RS12880 WP 074794436.1  IKQLDALILTHADQDHIGEAETLLKHHKVKRLVIPKGFVSEPKDEKVLQTAREEGVTIEEVKRGDVLQIKDLQFHVLSPE  640
NTDB id 107 BSU 25570 NP 390435.1             IKQLDALILTHADQDHIGEAEILLKHHKVKRLVIPKGFVSEPKDEKVLQAAREEGVAIEEVKRGDVLQIKDLQFHVLSPE  640
consensus                                     !!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!! !!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 999402 ABEG41 RS12880 WP 074794436.1  TPDPASKNNSSLVLWMETGVLSWILTGDLEKEGEQEVMDVFPNIKADVLKVGHHGSKGSTGEEFIQQLQPKTAIISAGKN  720
NTDB id 107 BSU 25570 NP 390435.1             APDPASKNNSSLVLWMETGGMSWILTGDLEKEGEQEVMNVFPNIKADVLKVGHHGSKGSTGEEFIQQLQPKTAIISAGKN  720
consensus                                      !!!!!!!!!!!!!!!!!! *!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 999402 ABEG41 RS12880 WP 074794436.1  NRYHHPHQKVLQLLQRHSIRVLRTDQNGTIQYRYKNRVGTFSVYPPYDTSDITETN  776
NTDB id 107 BSU 25570 NP 390435.1             NRYHHPHQKVLQLLQRHSIRVLRTDQNGTIQYRYKNRVGTFSVYPPYDTSDITETN  776
consensus                                     !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved