```
logo

NTDB id 421657 G3M69 RS00430 WP 121066139.1  MKEKPFNSEQLIYLEELLSHQEKHLESKLSGFSVNDLDMQSVFRLERNRLKIAYRLLGLMSFITLVLAIVLISILPLQKT   80
NTDB id 1216 CAA10655.1 697..1440( )         MREKPFNSEQLIYLEELLSHQEKHLENKLSGFSVSDLDMQSVFRLERNRLKIAYKLLGLMSFIALILAIVLISLLPLQKT   80
consensus                                    !*!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!! !!!!!!!!!!!!!!!!!!!*!!!!!!!! !*!!!!!!!*!!!!!!

logo

NTDB id 421657 G3M69 RS00430 WP 121066139.1  EHHFVDFLNQDKHYAIIQRADKSISSNEALARSLIGAYVLNRESINRIDDKSRYELVRLQSSSKVWQRFEDLIKAQNSIY  160
NTDB id 1216 CAA10655.1 697..1440( )         EHHFVDFLNQDKHYAIIQRADKSISSNEALARSLIGAYVLNRESINRIDDKSRYELVRLQSSSKVWQRFEDLIKTQNSIY  160
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 421657 G3M69 RS00430 WP 121066139.1  AQSHLEREIHIVNIAIYQQDNNPIASVSIAAKLTNENKLVYEKRYKIVLSYLFDTPDFDYASMPKNPTGFKITRYSITEI  240
NTDB id 1216 CAA10655.1 697..1440( )         VQSHLEREVHIVNIAIYQQDNNPIASVSIAAKLLNENKLVYEKRYKIVLSYLFDTPDFDYASMPKNPTGFKITRYSITEI  240
consensus                                     !!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

   <0

logo

NTDB id 421657 G3M69 RS00430 WP 121066139.1  VPTNRGD.   247
NTDB id 1216 CAA10655.1 697..1440( )         APTNRGD*   247
consensus                                     !!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved