```
logo

NTDB id 667771 MPG77 RS02235 WP 245109903.1  MKEKPFNSEQLIYLEELLSHQEKHLENKLSGFSVNDLDMQSVFRLERNRLKIAYKLLGLMSFIALVLAVVLISVLPLQKT   80
NTDB id 1216 CAA10655.1 697..1440( )         MREKPFNSEQLIYLEELLSHQEKHLENKLSGFSVSDLDMQSVFRLERNRLKIAYKLLGLMSFIALILAIVLISLLPLQKT   80
consensus                                    !*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!*!!!!*!!!!!!

logo

NTDB id 667771 MPG77 RS02235 WP 245109903.1  EHHFVDFLNQDKHYAIIQRADKSISSNEALARSLIGAYVLNRESINRIDDKSRYELVRLQSSSKVWQRFENLIKTQNSIY  160
NTDB id 1216 CAA10655.1 697..1440( )         EHHFVDFLNQDKHYAIIQRADKSISSNEALARSLIGAYVLNRESINRIDDKSRYELVRLQSSSKVWQRFEDLIKTQNSIY  160
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 667771 MPG77 RS02235 WP 245109903.1  AQSHLEREVHIVNIAIYQQDNNPIASVSIAAKLMNENKLVYEKRYKIVLSYLFDTPDFDYASMPKNPTGFKVTRYSITEI  240
NTDB id 1216 CAA10655.1 697..1440( )         VQSHLEREVHIVNIAIYQQDNNPIASVSIAAKLLNENKLVYEKRYKIVLSYLFDTPDFDYASMPKNPTGFKITRYSITEI  240
consensus                                     !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

   <0

logo

NTDB id 667771 MPG77 RS02235 WP 245109903.1  APTNRGD.   247
NTDB id 1216 CAA10655.1 697..1440( )         APTNRGD*   247
consensus                                    !!!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved