```
logo

NTDB id 1027729 AB3456 RS09685 WP 017722273.1  ..MIQPYIIKKGDMVLQPCNLATGQYEGSYLLRYKADSQIIKERVSKIIDRSCRFYGSSYVFKKEDTMRITGISSKPPIL   78
NTDB id 39 MW RS04875 WP 000287265.1           MYSQNIYVIRKGDMVIRPAFDDDDQRNGSEIIRFDKTRIQNPFKVQKIIERSCKFYGNTYLGKKAETNRITGISSKPPIL   80
NTDB id 23 SA RS04995 WP 000287265.1           MYSQNIYVIRKGDMVIRPAFDDDDQRNGSEIIRFDKTRIQNPFKVQKIIERSCKFYGNTYLGKKAETNRITGISSKPPIL   80
consensus                                      ******!*!*!!!!!**!******!**!!***!***********!*!!!*!!!*!!!**!**!**!*!!!!!!!!!!!!

logo

NTDB id 1027729 AB3456 RS09685 WP 017722273.1  FTPLFPTYFFPTHSDRKADNTWINIHYVHNIRPLKEKKCKVTFVDNQSIVANISAHSMHHQYLNGIYYYYMMDRAARVAT  158
NTDB id 39 MW RS04875 WP 000287265.1           LTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESIILHVSYHSLWHQYNNSIFYYYMVDKQSRMIS  160
NTDB id 23 SA RS04995 WP 000287265.1           LTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESIILHVSYHSLWHQYNNSIFYYYMVDKQSRMIS  160
consensus                                      *!!!!!!!!!!!!!!!***!*!*!*!!***!**!!**!!!!!!**!*!!****!*!!**!!!*!*!*!!!!*!***!***

logo

NTDB id 1027729 AB3456 RS09685 WP 017722273.1  FDPDSPIDYTKPQLNIYEALAKYSLFESK   187
NTDB id 39 MW RS04875 WP 000287265.1           KNPDQPIDYNKATLNVFEALTRYSLFEDK   189
NTDB id 23 SA RS04995 WP 000287265.1           KNPDQPIDYNKATLNVFEALTRYSLFEDK   189
consensus                                      **!!*!!!!*!**!!**!!!**!!!!!*!
```

X non conserved

X similar

X $\geq$ 50% conserved