```
logo
NTDB id 1095084 ACMVYP RS07120 WP 017175793.1  MTQFNKPPNTIYVIRKGDMVIRPTFDDHQQPNGSEILKFDTSREISPFKTQKIIERSCRFYGNNYLSKKAETQRIAGITS   80
NTDB id 39 MW RS04875 WP 000287265.1           .....MYSQNIYVIRKGDMVIRPAFDDDDQRNGSEIIRFDKTRIQNPFKVQKIIERSCKFYGNTYLGKKAETNRITGISS   75
NTDB id 23 SA RS04995 WP 000287265.1           .....MYSQNIYVIRKGDMVIRPAFDDDDQRNGSEIIRFDKTRIQNPFKVQKIIERSCKFYGNTYLGKKAETNRITGISS   75
consensus                                            *****!!!!!!!!!!!!!!*!!!**!*!!!!!**!!**!***!!!*!!!!!!!!*!!!!*!!*!!!!*!!*!!*!

logo
NTDB id 1095084 ACMVYP RS07120 WP 017175793.1  KPPILLTPLFPTYFFPTHSDRLEENIWINMHYIESIKELKNRKSKIIFANQEALTLNVSFHSLWHQYSNSIFYYYMIDKQ  160
NTDB id 39 MW RS04875 WP 000287265.1           KPPILLTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESIILHVSYHSLWHQYNNSIFYYYMVDKQ  155
NTDB id 23 SA RS04995 WP 000287265.1           KPPILLTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESIILHVSYHSLWHQYNNSIFYYYMVDKQ  155
consensus                                      !!!!!!!!!!!!!!!!!!!!!!**!!!!*!!!!!**!!*!!!!!!*!**!*!*!***!*!!*!!!!!!!!*!!!!!!!!!*!!!

logo
NTDB id 1095084 ACMVYP RS07120 WP 017175793.1  ARMKSNNPDMPIDYEKTSLNIFEALSHYSLLEDK  194
NTDB id 39 MW RS04875 WP 000287265.1           SRMISKNPDQPIDYNKATLNVFEALTRYSLFEDK  189
NTDB id 23 SA RS04995 WP 000287265.1           SRMISKNPDQPIDYNKATLNVFEALTRYSLFEDK  189
consensus                                      *!!*!*!!!*!!!!*!**!!*!!!!**!!!*!!!
```

X non conserved

X similar

X $\geq$ 50% conserved