```
logo
```

```
NTDB id 1119341 EL068 RS08575 WP 002465629.1  MTTQNIYVIRKGDMVIRPNYDNQNQPNGSEIIRFDKTVIQSPFKVQKIIERSCKFYGNNYLSKKAETNRITGISSKPPIL   80
NTDB id 39 MW RS04875 WP 000287265.1          MYSQNIYVIRKGDMVIRPAFDDDDQRNGSEIIRFDKTRIQNPFKVQKIIERSCKFYGNTYLGKKAETNRITGISSKPPIL   80
NTDB id 23 SA RS04995 WP 000287265.1          MYSQNIYVIRKGDMVIRPAFDDDDQRNGSEIIRFDKTRIQNPFKVQKIIERSCKFYGNTYLGKKAETNRITGISSKPPIL   80
consensus                                     !**!!!!!!!!!!!!!!!!**!***!*!!!!!!!!!!!*!!*!!!!!!!!!!!!!!!!!!*!!*!!!!!!!!!!!!!!!!!!
```

```
logo
```

```
NTDB id 1119341 EL068 RS08575 WP 002465629.1  LTPLFPTYFFPTHSDRQEENLWINMHYVESIKELKHRKCKIIFANNETITLNVSFHSLWHQYTNAIFFYYMVDKQSRMKS   160
NTDB id 39 MW RS04875 WP 000287265.1          LTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESIILHVSYHSLWHQYNNSIFYYYMVDKQSRMIS   160
NTDB id 23 SA RS04995 WP 000287265.1          LTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESIILHVSYHSLWHQYNNSIFYYYMVDKQSRMIS   160
consensus                                     !!!!!!!!!!!!!!!!!!*!!*!*!!!!*!!!!!!!*!!!!**!*!!!*!*!*!!*!!!!!!!*!*!!*!!!!!!!!!!*!
```

```
logo
```

```
NTDB id 1119341 EL068 RS08575 WP 002465629.1  NDPDKPIDYEKSSLNIFEALSRYSLFVDK   189
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 ≥ 50% conserved