```
logo
```

```
NTDB id 1055946 ACFDA7 RS08655 WP 001829272.1  MTHLPETIYVIRKGDMVIRPKYDEYQQTNGTEIIRFDQTRKESPFKVQRIIERSCKFYGNNYISKKAETNRITGISSKPP   80
NTDB id 39 MW RS04875 WP 000287265.1           ..MYSQNIYVIRKGDMVIRPAFDDDDQRNGSEIIRFDKTRIQNPFKVQKIIERSCKFYGNTYLGKKAETNRITGISSKPP   78
NTDB id 23 SA RS04995 WP 000287265.1           ..MYSQNIYVIRKGDMVIRPAFDDDDQRNGSEIIRFDKTRIQNPFKVQKIIERSCKFYGNTYLGKKAETNRITGISSKPP   78
consensus                                        *****!!!!!!!!!!!!!!**!***!*!!*!!!!!!*!!***!!!!!*!!!!!!!!!!!!!*!**!!!!!!!!!!!!!!!!
```

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logo
```

```
NTDB id 1055946 ACFDA7 RS08655 WP 001829272.1  ILLTPLFPTYFFPTHSDRQEENIWINMHYIENVKELKNRKSKIIFANGDSLTLNVSFHSLWHQYTNAIIYYYMVDKQSRM   160
NTDB id 39 MW RS04875 WP 000287265.1           ILLTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESIILHVSYHSLWHQYNNSIFYYYMVDKQSRM   158
NTDB id 23 SA RS04995 WP 000287265.1           ILLTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESIILHVSYHSLWHQYNNSIFYYYMVDKQSRM   158
consensus                                      !!!!!!!!!!!!!!!!!!!!*!!!!*!!!!!!**!!!!!!!*!**!*!**!**!*!!*!!!!!!!*!*!*!!!!!!!!!!
```

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logo
```

```
NTDB id 1055946 ACFDA7 RS08655 WP 001829272.1  KSNNPEQPIDYNQSSLNIFEALSRYSLFEEN   191
NTDB id 39 MW RS04875 WP 000287265.1           ISKNPDQPIDYNKATLNVFEALTRYSLFEDK   189
NTDB id 23 SA RS04995 WP 000287265.1           ISKNPDQPIDYNKATLNVFEALTRYSLFEDK   189
consensus                                      *!*!!*!!!!!!!***!!*!!!!*!!!!!!**
```

X non conserved
X similar
X ≥ 50% conserved