```
logo
T10169 WP 000269902.1 NC 019359:3621-3977      MWTVITTDLFNEWLAQQDESTQEKVLAALVVLQQQG.PSLGRPLVDTVYDSKFTNMKELRVQHRGKPLRAFFAFDPLRQA   79
T10097 WP 003120987.1 NC 016138:110952-111251  MVEVKQTATFMAWESKLKDRRAKAVIAARIFRLANGLPGDVSPVGQGV........SELR.IHYGPGYRVYFQQRGTEIV   71
T6106 WP 024431052.1 NC 006351:1437364-1437660 MFKVLTTPQFDKWLDGLRDPVGSAAINLRIERAKLGNLGQWRAVGDGV........NEMK.IDVGPGYRAYFVRREKIIV   71
T6105 WP 006027262.1 NC 006350:c179158-178862  MFKVLTTPQFDKWLDGLRDPVGSAAINLRIERAKLGNLGQWRAVGDGV........NEMK.IDVGPGYRAYFVRRGKIIV   71
consensus                                      !**!**!**!**!*********** *  *******!************!         *!** ***!***!**!********

logo
T10169 WP 000269902.1 NC 019359:3621-3977      IVLCIGDKGGKKRFYKEMLDIADQQYELHLSTLGDQSNG   118
T10097 WP 003120987.1 NC 016138:110952-111251  ILLCGGDKSSQARDIEMAKRLANEWRPQ...........    99
T6106 WP 024431052.1 NC 006351:1437364-1437660 VVLCGGDKSTQKKDIKLAKQIAGELED............    98
T6105 WP 006027262.1 NC 006350:c179158-178862  VVLCGGDKSTQKKDIKLAKQIAGELED............    98
consensus                                      **!!*!!!*************!*****
```

X non conserved

X similar

X $\geq$ 50% conserved