```
logo                                                  MNEKGFTLVEMLIVLFIYSILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQVTAYEIDHEGKMPDMNDLQSEGYYKKNTV
                                                      MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQTPSLADLQSEGYVKKDAV

NTDB id 1058695 ACFXAA RS19185 WP 050515801.1  ..........MLIVLFIVSILLLITIPNVTKHNQSIQHKGCEGLQNMVKAQVTAYEIDHEGKMPDMNDLQSEGYIKKNTV    70
NTDB id 98 BSU 24710 NP 390351.1               MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQTPSLADLQSEGYVKKDAV    80
consensus                                      **********!!!!!!!*!!!!!!!!!!!!!!!!*!!*!!!!!!!!!!!!!!*!!*!*!!!!  ! * !!!!!!!*!!  !
```

```
logo                                                  CPNGKRQILITSGGEVTKVEQH

NTDB id 1058695 ACFXAA RS19185 WP 050515801.1  CPNGKQILISGGEVTVEQ    88
NTDB id 98 BSU 24710 NP 390351.1               CPNGKRIIITGGEVKVEH    98
consensus                                      !!!!! !*!*!!!! !!*
```

X non conserved
X similar
X $\geq$50% conserved