```
logo                                          VKYWTPLALSKLNDAAEYIALENWAAKQWVQTVFTKVERLGSMPESGRWPEMPHTNYREYYSPYRYIVSYSHEKV
                                              MMEIIWSELALQDLGDAAEFIALDNPSAAEKLVNEVFDKTELLADFTPEMGRMVPEMPHTLEHLNYREIIFGHCRIFIKLDDATR
T172 Protein 396 NC 002506:387942-388193      .MKVVWSPLALQKLGDAAEFIALDNPSAAEKWVNEVFDKTELLGSMPEMGRMVPEMPHTNYREIIFGHYRIIYSLSHEIR    79
T178 WP 000229317.1 NC 002506:425872-426174   MVEIIWTELALSDLNDIAEYIALENVVAAKQLVQTVFTKVERLADFPESGRVPPELEHLNYREVVVSPCRVFYKYDDAKV    80
consensus                                     ** **!**!!!  ! ! !!*!!!*!* !!   !* !! ! ! !*  !! !!**!!**! !!!!**   *!* !   *
```

```
logo                                          VLTLV
                                              RILFRMRAERDLRRLMLTKQ
T172 Protein 396 NC 002506:387942-388193      VLTLR...............    84
T178 WP 000229317.1 NC 002506:425872-426174   RILFVMRAERDLRRLMLTKQ   100
consensus                                      *   ***************
```

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