```
logo
                                          MIDIYVLENNVNQQFRIEKMIGDLLETNQWETRHFDLFSEPDCLTNLVEESFPQIFSWILIIMGINTKGLKWLRR
                                          MIDIYVLEDNVNQQFRIEKMIGDLLETNQWETRHFDLFSEPDCLTNLVEESFPQIFFLDIDFHEDIPKETEVAKAIREKSRTATI

NTDB id 60851 KE3 RS11005 WP 228380495.1  MIDIYVLENNVNQQFRIEKMIGDLLETNQWETRHFDLFSEPDCLINLVEEPFPQFFSWILIFMEIIPKELKWLRR..........  75
NTDB id 545 H702 RS02385 WP 052071091.1   MIDIYVLEDNVNQQFRIEKMIGDLLETNQWETRHFDLFSEPDCLTNLVEESFPQIFFLDIDIHGDNTKGIEVAKAIREKSRTATI  85
consensus                                 !!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!*!!! !   *  *   *! *   * **********

logo
                                          VFVTTHSEFMLLTYRAQVSALDFIDKTDKDDAIKDNLKKCLQKVFEDQGFNIPKEIFVFENNKTKIRIPLEDILYFETAESHRLK

NTDB id 60851 KE3 RS11005 WP 228380495.1  ..................................................................................  75
NTDB id 545 H702 RS02385 WP 052071091.1   VFVTTHSEFMLLTYRAQVSALDFIDKTDKDDAIKDNLKKCLQKVFEDQGFNIPKEIFVFENNKTKIRIPLEDILYFETAESHRLK  170
consensus                                 ********************************************************************************

logo
                                          LITKTGQRHFYGSIKEVMQANQKLFRCHKSYLINLNNIVRLDKKEGLVYFDDDKACYVSRKYIKDLRSKMENLK

NTDB id 60851 KE3 RS11005 WP 228380495.1  ......................................................................  75
NTDB id 545 H702 RS02385 WP 052071091.1   LITKTGQRHFYGSIKEVMQANQKLFRCHKSYLINLNNIVRLDKKEGLVYFDDDKACYVSRKYIKDLRSKMENLK  244
consensus                                 ******************************************************************
```

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