```
logo                                                  MVLLLIFSMFLQFYLQRQVLTARQLRSEKEQLTAELMVSVALKTGLKSQGQLNFDSGNLTYNLLTDSSASSTSTGQTVSN
                                                      ..... DDAQQLKIEKERLTAELMVSLALKKDLKTSQQYHFDCGDLSYKLLTDLSADSKIGDHSSDE
NTDB id 1165719 AB3Y84 RS11165 WP 367294010.1         .....MFSMFLQFYLQRQIDDARQLRSEKEQLTAELMVSVALKTGLKSQGQIHFDSGDLSYNLLTDSSASSKITDHSSDE    75
NTDB id 592 KW2 RS10825 WP 021037947.1                MVLLLIFSLFLQFYLQKQVLTAQQLKIEKERLTAELMVSLALKKDLKTSGQLNFDCGNLTYKLLTDLSADSTSGGQTVSN    80
consensus                                             ******!!*!!!!!!!*!*  ! !!* !!! !!!!!!!!*!!!  !!* !!**!!*! !*! !!!! !! !    **

logo                                                  KTYQFSVRLKDGRNFQIKN
                                                      CIDIHA         AT
NTDB id 1165719 AB3Y84 RS11165 WP 367294010.1         KTYQFSIHLKDGANFQIKN    94
NTDB id 592 KW2 RS10825 WP 021037947.1                KTYCFDVRLKDGRIFQIKK    99
consensus                                             !!!*! **!!!!  !!!!
```

X non conserved

X similar

X $\geq$ 50% conserved