```
logo
NTDB id 327067 EG348 RS02580 WP 317126994.1  MKDIFLDLLFPNRCLDCNKIIESNLIVCNICFGKIQFTHFDYFEENTLKEICKTLFPVENAYALIQFEKESLSRKIIHEL   80
NTDB id 1333 RA0C RS05105 WP 004918078.1     ...MFLDFFLPNRCLKCNLIISSDEIVCWACKSHINFTYNDFGGKGDFYQKVVLKFPVNKAFSLMKYEKKELSQDIIHAL   77
consensus                                    ****!!!   !!!!! !! !! !  !!!  !  *!*!!* !*         *   !!!  !* !* *!!  !!  !!! !

logo
NTDB id 327067 EG348 RS02580 WP 317126994.1  KYKNREKTGKILADWITERVDFKNEKPDLIVSIPLHKKKLKERGYNQLHLFAETLSEFYNIPFDHEVILRSYYSKAQALK  160
NTDB id 1333 RA0C RS05105 WP 004918078.1     KYHSMERVGKVLAHWVSENMDLSHQGIDILTNIPLHPKKQRERGYNQLHLFTKTLSQLQRIPYSHYLLKRNFYSKPQALK  157
consensus                                    !!*  !* !!*!!*!**! *!  *  *!**  !!!!*!! *!!!!!!!!!!  !!!    !!* ! ** ! *!!!*!!!!

logo
NTDB id 327067 EG348 RS02580 WP 317126994.1  DKKHRLSTENLFAINKNISGKHVLLVDDVFTTGNTIATAAWEILKTGNNKVSVLVMAIDK  220
NTDB id 1333 RA0C RS05105 WP 004918078.1     QKLDREKMQNIFSVDEKIEGKHILLIDDVYTTGNTMSKAVWELLKGNGNQVSVLVLAIDE  217
consensus                                     ! *!    !*! *   ! !!!*!!*!!!!*!!!!!*  ! !!*!!   ! !!!!!*!!!
```

X non conserved

X similar

X ≥ 50% conserved