```
logo
NTDB id 326172 EGY07 RS02530 WP 027374733.1  MILDLLFPNRCIHCNRMIDAELLVCDLCFDQIHFTHYNYFENNPIREKCSLLFPVENTFALIQFEKENLSRKLIHELKYK   80
NTDB id 1333 RA0C RS05105 WP 004918078.1     MFLDFFLPNRCLKCNLIISSDEIVCWACKSHINFTYNDFGGKGDFYQKVVLKFPVNKAFSLMKYEKKELSQDIIHALKYH   80
consensus                                    ! !!   !!!!**!! *!  * *!!  !  *!*!!*  *    *   !* ! !!!   ! !* *!!  !!  *!! !!!*

logo
NTDB id 326172 EGY07 RS02530 WP 027374733.1  GREKVGKILAEWTTEVVDFKNRKPDLLVSVPLHPKKLRERGYNQLHLFTETLSEFYRIPFDHQLIKRNHYSKAQALKDKK  160
NTDB id 1333 RA0C RS05105 WP 004918078.1     SMERVGKVLAHWVSENMDLSHQGIDILTNIPLHPKKQRERGYNQLHLFTKTLSQLQRIPYSHYLLKRNFYSKPQALKQKL  160
consensus                                      !*!!!*!!*! *! *!  *  *!*!  *!!!!!! !!!!!!!!!!!! !!!   !!!* ! !*!!!*!!!*!!!! !

logo
NTDB id 326172 EGY07 RS02530 WP 027374733.1  HRLETVNTFSLTQPVSGQHILLIDDVFTTGNTISSIAWEILNSGNNRVSVLVMAIDV  217
NTDB id 1333 RA0C RS05105 WP 004918078.1     DREKMQNIFSVDEKIEGKHILLIDDVYTTGNTMSKAVWELLKGNGNQVSVLVLAIDE  217
consensus                                    *!    ! !!*  ** ! !!!!!!!!*!!!!!*!   !!*!    ! !!!!!*!!!
```

X non conserved

X similar

X ≥ 50% conserved