```
logo
NTDB id 133031 OK18 RS08455 WP 053327723.1  .MISEEYFYAIALRECSQIGDINFYKLVRTFGSAQEAWKRAKKEYRKIEGLGQKTVADIGNKEHLTFAEKELKYCEKNNIRITIR   84
NTDB id 1342 RA0C RS05130 WP 004918085.1    MVNAEEILYSIALRRSPLVGDVIFASLVSEVGSAKEAWGLSKLELRKISGVGKQIASNIGDKNLLKFAEKEIEFCIKNEIKIHLR   85
consensus                                   **  !!  ! !!!! ** *!!* !  !!   !!! !!!   ! ! !!! !*!      !! ! *! !!!!!* *! !! !*!**!

logo
NTDB id 133031 OK18 RS08455 WP 053327723.1  HLNELPALLNECADAPAILYQKGNYENSTPKISIVGTRNMTAYGRQFIEDFFEAAQSSRFISVSGLALGIDKEVHEKSIQYRKPT  169
NTDB id 1342 RA0C RS05130 WP 004918085.1    HLNQLPRLLGECYDAPAILFQKGDYEDTANNISIVGTRNATAYGKSFLEDLIVQFKNKNIQVISGLALGTDGIAHQEALKNNIKT  170
consensus                                   !!! !! !! !! !!!!!!*!!! !! * * !!!!!!!! !!!!* !*!!            *!!!!!! !   !   *    *!

logo
NTDB id 133031 OK18 RS08455 WP 053327723.1  IAVLAHGFQFLYPAKNRKLSERILQEGGCLITEFNSSRKPDRENFIQRNRIVAGISPSTIVVETGFGGGSVSTAAFANDYNREVF  254
NTDB id 1342 RA0C RS05130 WP 004918085.1    SAVLAHGLHIIYPSKHKILAEEILNNGGALFSEFCSDEKPDREHFLQRNRIVAGLSANTIVVESAYAGGSISTASFANQYNREVY  255
consensus                                    !!!!!! * *!! !** ! ! !!* !!*! *!!*!  !!!!!*!*!!!!!!!!*!* !!!!!****!!!*!!! !!! !!!!!*

logo
NTDB id 133031 OK18 RS08455 WP 053327723.1  ALPGKITDPYSQGCNQLIFHNKAIAISTIKDLVNMLGFNHPKEKTGELFPYYETSAQLTENQHIIYKIIKENPQITLDDLSQEAA  339
NTDB id 1342 RA0C RS05130 WP 004918085.1    ALAGRLTDKYSQGCNHLIFQNKAKIISSIKGLLKDLDLDNTS.KVLELFPSVE..VSLTPVQEPVYDIIKEVNDISLDDLSEKLN  337
consensus                                   !!*!**!!*!!!!!!*!!!*!!!  !!*!! !*  !   ** *!  !!!!  !**  !!* !***! !!!! * !*!!!!!

logo
NTDB id 133031 OK18 RS08455 WP 053327723.1  VSSYKILPIILELELLGKVKSFSGRQFMAI   369
NTDB id 1342 RA0C RS05130 WP 004918085.1    QPISKVFPVLLELELLGLIRANSGRRYSII   367
consensus                                    *  !* !**!!!!!!! **  !!! *  !
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X non conserved

X similar

X ≥ 50% conserved