```
logo
NTDB id 1130939 H0506 RS09315 WP 011226714.1  .MSEQLKTLSISEIYPNPFQPRLKFSDEELVELSQSISENGLIQPIIVRKSDIIGYELIAGERRLRACKRLGMTEIPAVV   79
NTDB id 382 SMU RS09895 WP 011074697.1        MMTEQLKYLNTKDINPNPYQPRLQFKTKELEELAQSIKENGLIQPIIVRKSDIFGYDLVAGERRLKAAKLAGLNKIPVII   80
consensus                                     *!*!!!! !   *! !!!*!!!! !   !! !! !!! !!!!!!!!!!!!!!! !!*!*!!!!!!*!*!  !*  !! **

logo
NTDB id 1130939 H0506 RS09315 WP 011226714.1  KEVTDQESRKQAIIENLQRSNLNPIEEAKAYRNLIDELTYSHEELAKAMGKSRPYISNALRLLQLPREIQTSIENGSLSQ   159
NTDB id 382 SMU RS09895 WP 011074697.1        KKISDDDSMKQAIIENLQRSDLNPIEEAKAYQNLINRNHMTHDDIAKVIGKSRPYITNSIRLLNLPLHISQALEKGLISQ   160
consensus                                     ! **! *! !!!!!!!!!!! !!!!!!!!!! !!!   * *!***!! *!!!!!!!*! *!!!*!! *!   *! ! *!!

logo
NTDB id 1130939 H0506 RS09315 WP 011226714.1  GHARALLAIEDSRKQLTIFQQVVAERWSVRTLEKKLQELPR.KQKSKKDIHVKDKEKELERLLGLPITLRYHKNHSGTIQ   238
NTDB id 382 SMU RS09895 WP 011074697.1        GHARLLLSIEKQDLQDKWFQKILTEQLSVHQIERALKSQTKKEKKTSKDIFLAEKEKELSQSLGLPVVIHYNKKHQGQLK   240
consensus                                     !!!! !! !!    !   !! ** !  !!* *!* !   ***  !* !!!** *!!!!!   !!!!* **!*! ! ! *

logo
NTDB id 1130939 H0506 RS09315 WP 011226714.1  IHFSTEEDFNRIINKLI   255
NTDB id 382 SMU RS09895 WP 011074697.1        ISFSSEEDFNRLMNKLN   257
consensus                                     !*!!*!!!!!!**!!!
```

X non conserved

X similar

X $\geq$ 50% conserved