```
logo
NTDB id 1081432 CLC19175 RS06475 WP 257402287.1  MNIFINLPTWLGDAVMASAAIYAIKEKYPQAKFTFYGSFVSTELFKHFENAQILVENKKQRYRQILQARKNLGKFDLAFS   80
NTDB id 1241 Cj1148 YP 002344539.1               MKIFIHLPTWLGDTVMASPALYTIKEHFKDAQFILYGSFVSTALFKEFPNSKIIIENKLSRYKQALSLRKELGKIDLSFA   80
consensus                                        ! !!!*!!!!!!! !!!!*!*! !!!*** ! !  !!!!!!! !!!*!*!  !**!!!  !!*! !  !! !!! !! !

logo
NTDB id 1081432 CLC19175 RS06475 WP 257402287.1  FRSAFSSKIILNLIKAKKRFYFDKNILKEEHQVLKYLNFIEKALNFKATSNALKLPIKA........KSTQKILGINAGA  152
NTDB id 1241 Cj1148 YP 002344539.1               FRSAFSSKIILHILKTKQRYFFDKNKYKEEHQVLKYLYFIENSLSIKAHFKDLKLPFKLKFQNPLILRNGKKILGLNPGA  160
consensus                                        !!!!!!!!!!!***! ! !**!!!!  !!!!!!!!!! !!!  !  !!*   !!!! ! *********   !!!!*!*!!

logo
NTDB id 1081432 CLC19175 RS06475 WP 257402287.1  HFGSAKRWEASYFARVAKEFSSTHQILIFGVE.SEREICDEIEQFLLKEGIKAKNLCGKTSIYTLCKNISMLDLLITNDS  231
NTDB id 1241 Cj1148 YP 002344539.1               SFGSAKRWDASYFAKVALNFSQSHDILIFGAGKAEQELCNEIYQILKEQNIKVKNLCNKTTIKTLCQNIAFCDLFITNDS  240
consensus                                        *!!!!!!!*!!!!!*!!  !! *! !!!!!  * ! !*! !! ! !   !! !!!! !!*! !!! !!  *!! !!!!!

logo
NTDB id 1081432 CLC19175 RS06475 WP 257402287.1  GPMHIGAVYGVKTVAIFGSTKFSQTSPWQK.NVKIAHLNLACMPCMQKVCPLKHHKCMKDLKPEVVINLTKTFF.....  304
NTDB id 1241 Cj1148 YP 002344539.1               GPMHLSAVYKVKTVAIFGPTKFTQTSPWQNENAKLVHLDLACMPCMQKTCPLKHHKCMKDLKPEKVIEQAKNLLKNSHL  319
consensus                                        !!!!* !!! !!!!!!!!*!!!*!!!!!! *! !* !! !!!!!!!!! !!!!!!!!!!!!!!! !!   !   *****
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X non conserved

X similar

X $\geq$ 50% conserved