```
logo
NTDB id 1081416 CLC19168 RS06015 WP 257402623.1  MNIFINLPTWLGDSVMASAAIYAIKEKYPQAKFTFYGSFVSTELFKHFENAQVLVENKKQRYKQILKARKNFGKFDLAFS   80
NTDB id 1241 Cj1148 YP 002344539.1               MKIFIHLPTWLGDTVMASPALYTIKEHFKDAQFILYGSFVSTALFKEFPNSKIIIENKLSRYKQALSLRKELGKIDLSFA   80
consensus                                        ! !!!*!!!!!!!*!!!!*!*! !!!*** ! !  !!!!!!! !!!*!*!  ***!!!  !!!! ! !!  !! !! !

logo
NTDB id 1081416 CLC19168 RS06015 WP 257402623.1  FRSAFSSKIILNLIKAKKRFYFDKNILKEEHQVLKYLNFIEKALNFKATSNSLKLPIKA........KSTQKLLGINAGA  152
NTDB id 1241 Cj1148 YP 002344539.1               FRSAFSSKIILHILKTKQRYFFDKNKYKEEHQVLKYLYFIENSLSIKAHFKDLKLPFKLKFQNPLILRNGKKILGLNPGA  160
consensus                                        !!!!!!!!!!!***! ! !**!!!!  !!!!!!!!!! !!!  !  !!*   !!!! ! *********   !*!!*!*!!

logo
NTDB id 1081416 CLC19168 RS06015 WP 257402623.1  HFGSAKRWEASYFARVAKEFSFTHKILIFGVE.SEREICDEIEHYLLKEGIKAKNLCGKTSIYTLCKNISMLDLLITNDS  231
NTDB id 1241 Cj1148 YP 002344539.1               SFGSAKRWDASYFAKVALNFSQSHDILIFGAGKAEQELCNEIYGILKEQNIKVKNLCNKTTIKTLCQNIAFCDLFITNDS  240
consensus                                        *!!!!!!!*!!!!!*!!  !! *! !!!!!  * ! !*! !! * !   !! !!!! !!*! !!! !!  *!! !!!!!

logo
NTDB id 1081416 CLC19168 RS06015 WP 257402623.1  GPMHIGAVYGVKTVAIFGSTKFGQTSPWQK.NAKIAHLNLACMPCMQKVCPLKHHKCMKDLKPEVVINLAKTFF.....  304
NTDB id 1241 Cj1148 YP 002344539.1               GPMHLSAVYKVKTVAIFGPTKFTQTSPWQNENAKLVHLDLACMPCMQKTCPLKHHKCMKDLKPEKVIEQAKNLLKNSHL  319
consensus                                        !!!!* !!! !!!!!!!!*!!! !!!!!! *!!!* !! !!!!!!!!! !!!!!!!!!!!!!!! !!  !!   *****
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X non conserved

X similar

X ≥ 50% conserved