```
logo
NTDB id 389157 F6X58 RS06860 WP 139453565.1  MNIFINLPTWLGDAVMASSAIYAIKEKYPKAKFTFYGSFVSTELFKRFENTQILVEHKKQRYKQILKARKSFGKFDLAFS   80
NTDB id 1241 Cj1148 YP 002344539.1           MKIFIHLPTWLGDTVMASPALYTIKEHFKDAQFILYGSFVSTALFKEFPNSKIIENKLSRYKQALSLRKELGKIDLSFA   80
consensus                                    ! !!!*!!!!!!! !!!!*!*! !!!*** ! !  !!!!!!! !!! !*!* !**!*!  !!!! !  !!  !! !! !

logo
NTDB id 389157 F6X58 RS06860 WP 139453565.1  FRSAFSSKIILNLIKAKKRFYFEKNILKEEHQVLKYLNFIEKALNFKASSNALKLPIKE........KSTQKILGINPGA  152
NTDB id 1241 Cj1148 YP 002344539.1           FRSAFSSKIILHILKTKQRYFFDKNKYKEEHQVLKYLYFIENSLSIKAHFKDLKLPFKLKFQNPLILRNGKKILGLNPGA  160
consensus                                    !!!!!!!!!!!***! ! !**!*!!  !!!!!!!!!!! !!!  !  !!*   !!!! ! *********   !!!!*!!!!

logo
NTDB id 389157 F6X58 RS06860 WP 139453565.1  HFGSAKRWEASYFARVAKEFSSTHQILIFGVE.SEREICDEIEHFLLKEGIKAKNLCGKTSIYTLCKNISMLDLLITNDS  231
NTDB id 1241 Cj1148 YP 002344539.1           SFGSAKRWDASYFAKVALNFSQSHDILIFGAGKAEQELCNEIYGILKEQNIKVKNLCNKTTIKTLCQNIAFCDLFITNDS  240
consensus                                    *!!!!!!!*!!!!!*!!  !! *! !!!!!  * ! !*! !! * !    !! !!!! !!*! !!! !!  *!! !!!!!

logo
NTDB id 389157 F6X58 RS06860 WP 139453565.1  GPMHIGAVYGVKTVAVFGSTKFSQTSPWQE.NAKIAHLNLACMPCMQKVCPLKHHKCMKDLKPELVINLAKNLLGV...  306
NTDB id 1241 Cj1148 YP 002344539.1           GPMHLSAVYKVKTVAIFGPTKFTQTSPWQNENAKLVHLDLACMPCMQKTCPLKHHKCMKDLKPEKVIEQAKNLLKNSHL  319
consensus                                    !!!!* !!! !!!!!*!!*!!!*!!!!!! *!!!* !! !!!!!!!!! !!!!!!!!!!!!!!! !!  !!!!!  ***
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X non conserved

X similar

X $\geq$ 50% conserved