```
logo
NTDB id 396061 GFB47 RS01105 WP 153445877.1  MIDFRDLDFEEMPEWPVLPQVIVIGVLVLILQVFGAWFYLSPQHDELDQIRQQEVTLKSSIRIKAAKVAALPQLQAQLDE   80
NTDB id 1406 DSB67 RS13990 WP 010643497.1    MVNWQDLEMDEIAEWPLVPQLVAILLMVVLIQGFGYWFYLKPEEARLNTLIHEEQTLKVALRIKANKVAALPQLQSQLDE   80
NTDB id 1173 A1552VC RS12115 WP 000157690.1  MASLQELELDEISEWPLLPQLVVILLIMLLIQGAGYWFYLMPKQDEIALLKQEEETVKATLRIKANKVAVLPQIQAQLDE   80
consensus                                    !* ***!***!* !!!**!!***!*******!**!*!!!! !  ***  ****! !*! **!!!!*!!!*!!!*!*!!!!

logo
NTDB id 396061 GFB47 RS01105 WP 153445877.1  LNTRYEFLLKQLPVQKELASMLSSVNDLGLKNDLSFTRVDWGEKQVQEFLYRLPLNIELTGAYDNIGQFSEAIAKLPRII  160
NTDB id 1406 DSB67 RS13990 WP 010643497.1    LTTRYELLLQQLPAQKELATMLASVNELGIDNKLTFTRIDWGDKESEHFLYRLPLNIELTGDFHDIGRFSQAIATLPRII  160
NTDB id 1173 A1552VC RS12115 WP 000157690.1  LKERYDFLLRQLPVQKELASMLASVNQLGLDSSLTFTRIDWGERESQEFLYRLPLNIELTGSYHDIGDFSQAIAKLPRII  160
consensus                                    ! *!!**!!*!!!*!!!!!*!!*!!!*!!*** !*!!!*!!!******!!!!!!!!!!!!! ***!! !!*!!!*!!!!!

logo
NTDB id 396061 GFB47 RS01105 WP 153445877.1  SLHDVEWNRVSQESGTLHFRVRAYTYQFKPEETKK.   195
NTDB id 1406 DSB67 RS13990 WP 010643497.1    TFQDVTWQRVSQESETLHFRVRANTYQFKP......   190
NTDB id 1173 A1552VC RS12115 WP 000157690.1  NFDDVDWQRVSQESSTLHFRVRAYTYQFKQEVDDEK   196
consensus                                    ** !!*!*!!!!!! !!!!!!!!*!!!!!**
```

X non conserved

X similar

X $\geq$ 50% conserved