```
logo

NTDB id 1063001 ACG4I4 RS01305 WP 000157690.1  MASLQELELDEISEWPLLPQLVVILLIMLLIQGAGYWFYLMPKQDEIALLKQEEETVKATLRIKANKVAVLPQIQAQLDE   80
NTDB id 1173 A1552VC RS12115 WP 000157690.1    MASLQELELDEISEWPLLPQLVVILLIMLLIQGAGYWFYLMPKQDEIALLKQEEETVKATLRIKANKVAVLPQIQAQLDE   80
NTDB id 1406 DSB67 RS13990 WP 010643497.1      MVNWQDLEMDEIAEWPLVPQLVAILLMVVLIQGFGYWFYLKPEEARLNTLIHEEQTLKVALRIKANKVAALPQLQSQLDE   80
consensus                                      !***!*!!*!!!*!!!!*!!!!!*!!!***!!!!*!!!!!!!*!*******!**!!*!*!**!!!!!!!!!!*!!!*!*!!!!

logo

NTDB id 1063001 ACG4I4 RS01305 WP 000157690.1  LKERYDFLLRQLPVQKELASMLASVNQLGLDSSLTFTRIDWGERESQEFLYRLPLNIELTGSYHDIGDFSQAIAKLPRII   160
NTDB id 1173 A1552VC RS12115 WP 000157690.1    LKERYDFLLRQLPVQKELASMLASVNQLGLDSSLTFTRIDWGERESQEFLYRLPLNIELTGSYHDIGDFSQAIAKLPRII   160
NTDB id 1406 DSB67 RS13990 WP 010643497.1      LTTRYELLLQQLPAQKELATMLASVNELGIDNKLTFTRIDWGDKESEHFLYRLPLNIELTGDFHDIGRFSQAIATLPRII   160
consensus                                      !**!!**!!*!!!*!!!!!*!!!!!!*!!*!**!!!!!!!!!!**!!**!!!!!!!!!!!!!!**!!!!*!!!!!!*!!!!!

logo

NTDB id 1063001 ACG4I4 RS01305 WP 000157690.1  NFDDVDWQRVSQESSTLHFRVRAYTYQFKQEVDDEK   196
NTDB id 1173 A1552VC RS12115 WP 000157690.1    NFDDVDWQRVSQESSTLHFRVRAYTYQFKQEVDDEK   196
NTDB id 1406 DSB67 RS13990 WP 010643497.1      TFQDVTWQRVSQESETLHFRVRANTYQFKP......   190
consensus                                      *!*!!*!!!!!!!!!*!!!!!!!!*!!!!!*******
```

X non conserved
X similar
X $\geq$ 50% conserved