```
logo
NTDB id 35730 SDSE RS19355 WP 003057032.1  MLNIENLTGGYLNIPVLKEVSFSVGNGELVGLIGLNGAGKSTTINEIIGFLKPYQGEISIDGLTLTADPAAYRKKIGFIPETPSL   85
NTDB id 473 HSISS4 RS07770 WP 021143710.1  MLKLEKVTGGYANIPVLKDVTFEVADGELVGLIGLNGAGKSTTINEIIGLLTPYGGQIAIDGLTIQQDPASYRQKIGYIPETPSL   85
NTDB id 481 STER RS07730 WP 002951673.1    MLKLEKVTGGYANIPVLKDVTFEVADGELVGLIGLNGAGKSTTINEIIGLLTPYGGQIAIDGLTIQQDPASYRQKIGYIPETPSL   85
consensus                                  !!**!**!!!!*!!!!!!*!*!*!**!!!!!!!!!!!!!!!!!!!!!!!!!!*!*!!*!*!*!!!!!***!!!*!!*!!!*!!!!!!!

logo
NTDB id 35730 SDSE RS19355 WP 003057032.1  YEELTLAEHINTVAMAYDIDVELAQKRAEPFLKMFRLADKLDWFPVHFSKGMKQKVMIICAFVIDPSLFILDEPFLGLDPLAISD  170
NTDB id 473 HSISS4 RS07770 WP 021143710.1  YEELTLREHLETVLMAYGLDVTEGMVRADKYLKLFRLDEKLDWFPTQFSKGMKQKVMIICAFIVNPSLFIVDEPFLGLDPLAISD  170
NTDB id 481 STER RS07730 WP 002951673.1    YEELTLREHLETVLMAYGLDVTEGMARADKYLKLFRLDEKLDWFPTQFSKGMKQKVMIICAFIVNPSLFIVDEPFLGLDPLAISD  170
consensus                                  !!!!!!*!!**!!*!!!**!!**** !!***!!*!!!**!!!!!!**!!!!!!!!!!!!!!!!!***!!!!!*!!!!!!!!!!!!!!

logo
NTDB id 35730 SDSE RS19355 WP 003057032.1  LIQTLELEKGKGKSILMSTHVLDSAEKMCDRFVILHHGQVRAQGTLADLQDAFGDASASLNDIYLALTKEG..   241
NTDB id 473 HSISS4 RS07770 WP 021143710.1  LIELLAEEKAKGKAILMSTHVLDSAEKMCDRFVILHHGQVLAQGTLEELRQTFGDDSASLNDIYMQLTKGELS   243
NTDB id 481 STER RS07730 WP 002951673.1    LIELLAEEKANGKSILMSTHVLDSAEKMCDRFVILHHGQILAQGTIDELRQTFGDSSASLNDIYMQLTKGELS   243
consensus                                  !!**!**!!**!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!**!!!!**!***!!! !!!!!!!!**!!!****
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X non conserved

X similar

X ≥ 50% conserved