```
logo
NTDB id 315491 D5R92 RS03455 WP 003101905.1  MLKIEHLTGGYLNIPVLKDVSFSVEDGQLVGLIGLNGAGKSTTINEIIGLLKPYQGKISIDGLTLDENMAEYRKKIGFIP   80
NTDB id 473 HSISS4 RS07770 WP 021143710.1    MLKLEKVTGGYANIPVLKDVTFEVADGELVGLIGLNGAGKSTTINEIIGLLTPYGGQIAIDGLTIQQDPASYRQKIGYIP   80
NTDB id 481 STER RS07730 WP 002951673.1      MLKLEKVTGGYANIPVLKDVTFEVADGELVGLIGLNGAGKSTTINEIIGLLTPYGGQIAIDGLTIQQDPASYRQKIGYIP   80
consensus                                    !!!*!**!!!!*!!!!!!!!!*!*!*!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!*!*!*!!!!!*****!*!!*!!!*!!

logo
NTDB id 315491 D5R92 RS03455 WP 003101905.1  ETPSLYEELTLSEHLETIAMAYDIDVKLAMERAKPLLESYRLTDKLDWYPVNFSKGMKQKVMIICAFIIDPSLFILDEPF  160
NTDB id 473 HSISS4 RS07770 WP 021143710.1    ETPSLYEELTLREHLETVLMAYGLDVTEGMVRADKYLKLFRLDEKLDWFPTQFSKGMKQKVMIICAFIVNPSLFIVDEPF  160
NTDB id 481 STER RS07730 WP 002951673.1      ETPSLYEELTLREHLETVLMAYGLDVTEGMARADKYLKLFRLDEKLDWFPTQFSKGMKQKVMIICAFIVNPSLFIVDEPF  160
consensus                                    !!!!!!!!!!!*!!!!!**!!!**!!***! !!***!***!!**!!!!*!**!!!!!!!!!!!!!!!!!**!!!!!*!!!!

logo
NTDB id 315491 D5R92 RS03455 WP 003101905.1  LGLDPLAISDLISYLESEKVKGKSILMSTHVLDSAEKMCDRFVILHQGQVRAQGTLEELRQEFAQTGASLNDIYLALTRE  240
NTDB id 473 HSISS4 RS07770 WP 021143710.1    LGLDPLAISDLIELLAEEKAKGKAILMSTHVLDSAEKMCDRFVILHHGQVLAQGTLEELRQTFGDDSASLNDIYMQLTKG  240
NTDB id 481 STER RS07730 WP 002951673.1      LGLDPLAISDLIELLAEEKANGKSILMSTHVLDSAEKMCDRFVILHHGQILAQGTIDELRQTFGDSSASLNDIYMQLTKG  240
consensus                                    !!!!!!!!!!!!**!**!**!!*!!!!!!!!!!!!!!!!!!!
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