```
logo
NTDB id 261189 SPSF3K RS09985 WP 003105648.1  MLKIENLTGGYINIPVLKDISFSVDAGQLVGLIGLNGAGKSTTINEIIGLLRPYKGSITIDGISLIDDKTAYRKKIGFIP   80
NTDB id 481 STER RS07730 WP 002951673.1       MLKLEKVTGGYANIPVLKDVTFEVADGELVGLIGLNGAGKSTTINEIIGLLTPYGGQIAIDGLTIQQDPASYRQKIGYIP   80
NTDB id 473 HSISS4 RS07770 WP 021143710.1     MLKLEKVTGGYANIPVLKDVTFEVADGELVGLIGLNGAGKSTTINEIIGLLTPYGGQIAIDGLTIQQDPASYRQKIGYIP   80
consensus                                     !!!*!**!!!!*!!!!!!!**!*!**!*!!!!!!!!!!!!!!!!!!!!!!!!*!!*!*!*!!!*****!***!!*!!!*!!

logo
NTDB id 261189 SPSF3K RS09985 WP 003105648.1  ETPSLYEELTLAEHIETIAMAYDIPVKEAHDRAKPLLKTYRLEDKLDWYPVHFSKGMKQKVMIICAFIIDPRLFILDEPF  160
NTDB id 481 STER RS07730 WP 002951673.1       ETPSLYEELTLREHLETVLMAYGLDVTEGMARADKYLKLFRLDEKLDWFPTQFSKGMKQKVMIICAFIVNPSLFIVDEPF  160
NTDB id 473 HSISS4 RS07770 WP 021143710.1     ETPSLYEELTLREHLETVLMAYGLDVTEGMVRADKYLKLFRLDEKLDWFPTQFSKGMKQKVMIICAFIVNPSLFIVDEPF  160
consensus                                     !!!!!!!!!!!*!!*!!**!!!***!*!** !!***!!**!!**!!!!*!**!!!!!!!!!!!!!!!!!**!*!!!*!!!!

logo
NTDB id 261189 SPSF3K RS09985 WP 003105648.1  LGLDPLAIADLIAYLDEEKKKGKSILMSTHVLDSAEKMCDRFVILHHGQIRAQGTLDDLRTAFGDNQASLNDIYLALTKE  240
NTDB id 481 STER RS07730 WP 002951673.1       LGLDPLAISDLIELLAEEKANGKSILMSTHVLDSAEKMCDRFVILHHGQILAQGTIDELRQTFGDSSASLNDIYMQLTKG  240
NTDB id 473 HSISS4 RS07770 WP 021143710.1     LGLDPLAISDLIELLAEEKAKGKAILMSTHVLDSAEKMCDRFVILHHGQVLAQGTLEELRQTFGDDSASLNDIYMQLTKG  240
consensus                                     !!!!!!!!*!!!**!*!!!**!!*!!!!!!!!!!!!!!!!!!!
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