```
logo
NTDB id 1138657 DQM55 RS02515 WP 172454704.1  MLEIKELTGGYVNIPVLKDVSFEVGNGQLVGLIGLNGAGKSTTINEIIGLLTPYKGQIQIDGLELRTNPSDYRKKIGFIP   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 1138657 DQM55 RS02515 WP 172454704.1  ETPSLYEELTLREHIETVAMAYDIEQETAFKRVDKLLEMFRLKEKLDWFPVHFSKGMKQKVMIICAFVVDPSLFIVDEPF  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 1138657 DQM55 RS02515 WP 172454704.1  LGLDPVAIADLIQLLEEEKKKGKSILMSTHVLDSAEKMCDSFVILHKGQVRAKGSLTELRAQFQMPDASLNDIYLALTEE  240
NTDB id 473 HSISS4 RS07770 WP 021143710.1     LGLDPLAISDLIELLAEEKAKGKAILMSTHVLDSAEKMCDRFVILHHGQVLAQGTLEELRQTFGDDSASLNDIYMQLTKG  240
NTDB id 481 STER RS07730 WP 002951673.1       LGLDPLAISDLIELLAEEKANGKSILMSTHVLDSAEKMCDRFVILHHGQILAQGTIDELRQTFGDSSASLNDIYMQLTKG  240
consensus                                     !!!!!*!!*!!!!*!!*!!!!**!!*!!!!!!!!!!!!!!!!!!!*!!!!!*!!**!*!***!!!**!!** *!!!!!!!**!!**

logo
NTDB id 1138657 DQM55 RS02515 WP 172454704.1  AAL   243
NTDB id 473 HSISS4 RS07770 WP 021143710.1     ELS   243
NTDB id 481 STER RS07730 WP 002951673.1       ELS   243
consensus                                     ***
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X non conserved

X similar

X ≥ 50% conserved