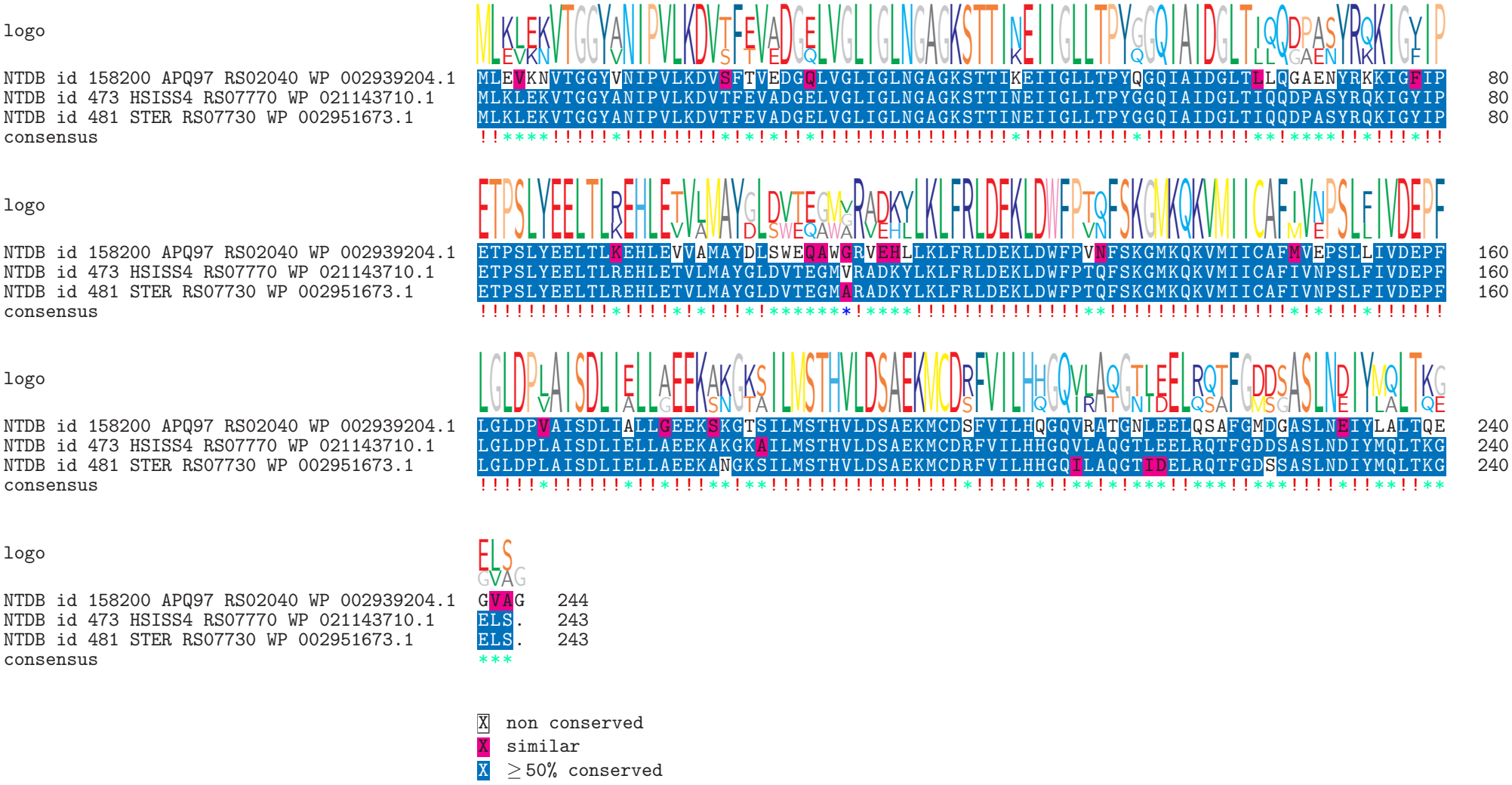
logo
NTDB id 158200 APQ97 RS02040 WP 002939204.1 MLEVKNVTGGYVNIPVLKDVSFTVEDGQLVGLIGLNGAGKSTTIKEIIGLLTPYQGQIAIDGLTLLQGAENYRKKIGFIP 80
NTDB id 473 HSISS4 RS07770 WP 021143710.1 MLKLEKVTGGYANIPVLKDVTFEVADGELVGLIGLNGAGKSTTINEIIGLLTPYGGQIAIDGLTIQQDPASYRQKIGYIP 80
NTDB id 481 STER RS07730 WP 002951673.1 MLKLEKVTGGYANIPVLKDVTFEVADGELVGLIGLNGAGKSTTINEIIGLLTPYGGQIAIDGLTIQQDPASYRQKIGYIP 80
consensus
logo
NTDB id 158200 APQ97 RS02040 WP 002939204.1 ETPSLYEELTLKEHLEVVAMAYDLSWEQAWGRVEHLLKLFRLDEKLDWFPVNFSKGMKQKVMIICAFMVEPSLLIVDEPF 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 158200 APQ97 RS02040 WP 002939204.1 LGLDPVAISDLIALLGEEKSKGTSILMSTHVLDSAEKMCDSFVILHQGQVRATGNLEELQSAFGMDGASLNEIYLALTQE 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 158200 APQ97 RS02040 WP 002939204.1 GVAG 244
NTDB id 473 HSISS4 RS07770 WP 021143710.1 ELS. 243
NTDB id 481 STER RS07730 WP 002951673.1 ELS. 243
consensus
X non conserved
X similar
X ≥ 50% conserved