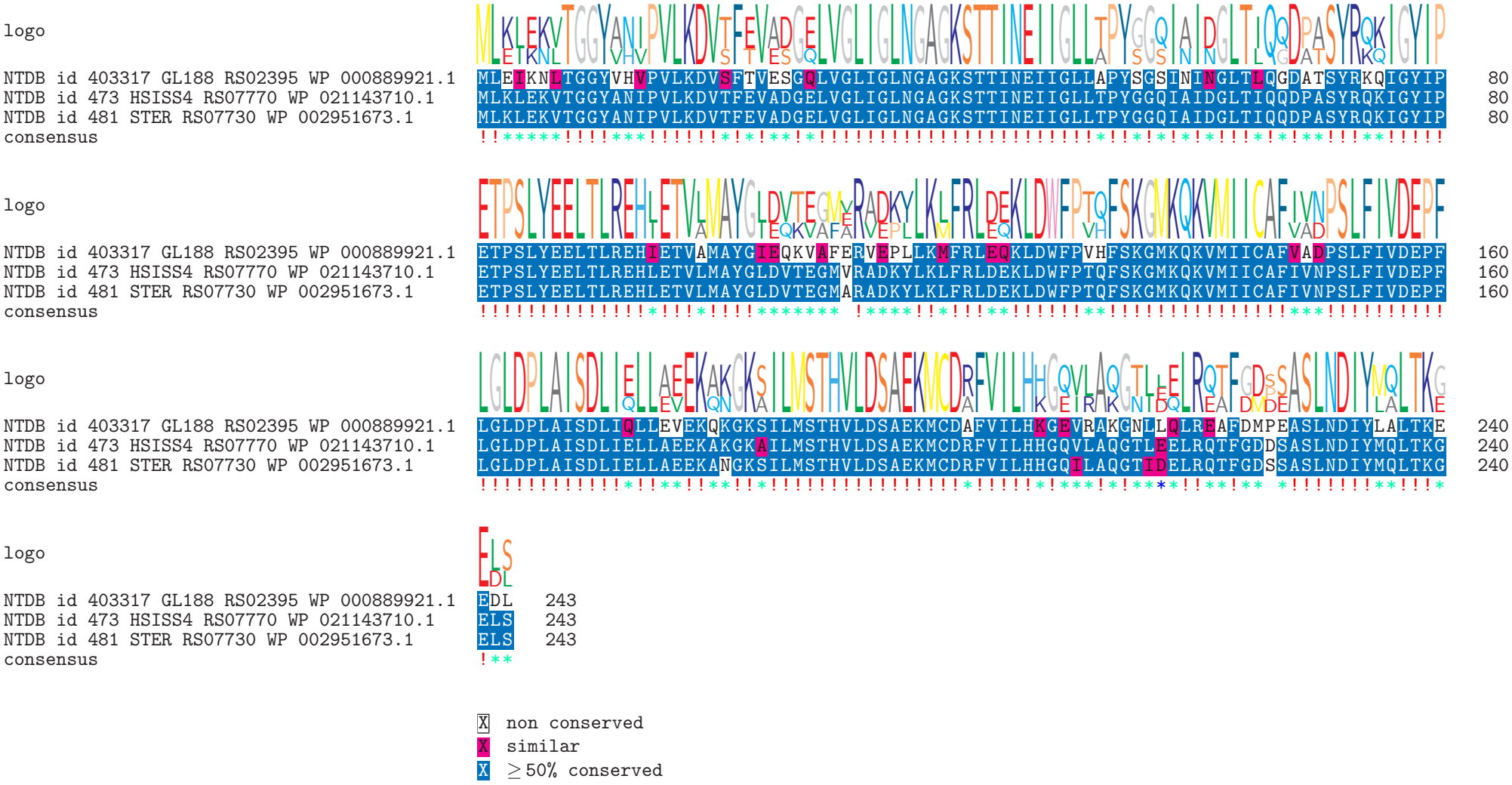
logo
NTDB id 403317 GL188 RS02395 WP 000889921.1 MLEIKNLTGGYVHVPVLKDVSFTVESGQLVGLIGLNGAGKSTTINEIIGLLAPYSGSININGLTLQGDATSYRKQIGYIP 80
NTDB id 473 HSISS4 RS07770 WP 021143710.1 MLKLEKVTGGYANIPVLKDVTFEVADGELVGLIGLNGAGKSTTINEIIGLLTPYGGQIAIDGLTIQQDPASYRQKIGYIP 80
NTDB id 481 STER RS07730 WP 002951673.1 MLKLEKVTGGYANIPVLKDVTFEVADGELVGLIGLNGAGKSTTINEIIGLLTPYGGQIAIDGLTIQQDPASYRQKIGYIP 80
consensus
logo
NTDB id 403317 GL188 RS02395 WP 000889921.1 ETPSLYEELTLREHIETVAMAYGIEQKVAFERVEPLLKMFRLEQKLDWFPVHFSKGMKQKVMIICAFVADPSLFIVDEPF 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 403317 GL188 RS02395 WP 000889921.1 LGLDPLAISDLIQLLEVEKQKGKSILMSTHVLDSAEKMCDAFVILHKGEVRAKGNLLQLREAFDMPEASLNDIYLALTKE 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 403317 GL188 RS02395 WP 000889921.1 EDL 243
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