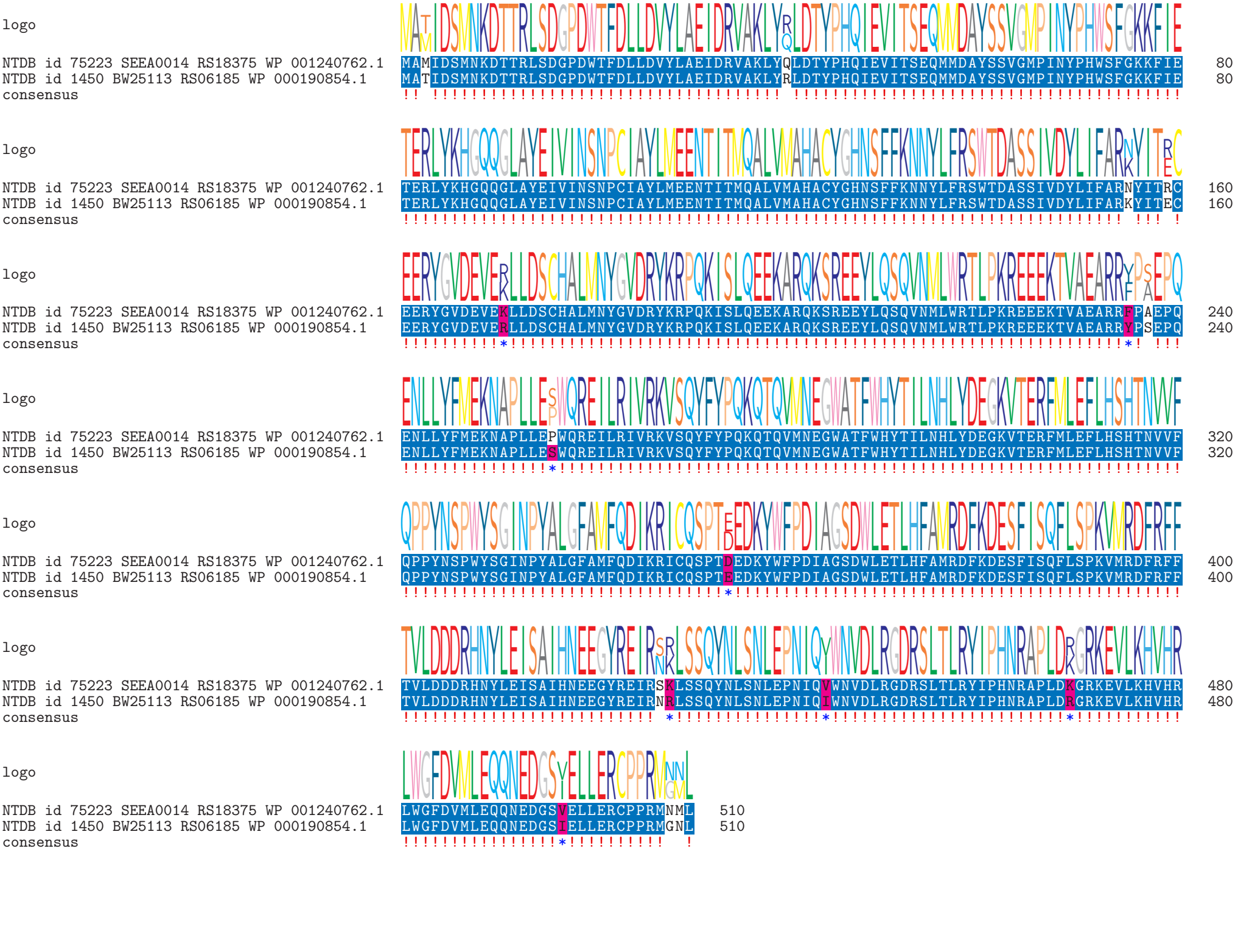
logo
NTDB id 75223 SEEA0014 RS18375 WP 001240762.1 MAMIDSMNKDTTRLSDGPDWTFDLLDVYLAEIDRVAKLYQLDTYPHQIEVITSEQMMDAYSSVGMPINYPHWSFGKKFIE 80
NTDB id 1450 BW25113 RS06185 WP 000190854.1 MATIDSMNKDTTRLSDGPDWTFDLLDVYLAEIDRVAKLYRLDTYPHQIEVITSEQMMDAYSSVGMPINYPHWSFGKKFIE 80
consensus
logo
NTDB id 75223 SEEA0014 RS18375 WP 001240762.1 TERLYKHGQQGLAYEIVINSNPCIAYLMEENTITMQALVMAHACYGHNSFFKNNYLFRSWTDASSIVDYLIFARNYITRC 160
NTDB id 1450 BW25113 RS06185 WP 000190854.1 TERLYKHGQQGLAYEIVINSNPCIAYLMEENTITMQALVMAHACYGHNSFFKNNYLFRSWTDASSIVDYLIFARKYITEC 160
consensus
logo
NTDB id 75223 SEEA0014 RS18375 WP 001240762.1 EERYGVDEVEKLLDSCHALMNYGVDRYKRPQKISLQEEKARQKSREEYLQSQVNMLWRTLPKREEEKTVAEARRFPAEPQ 240
NTDB id 1450 BW25113 RS06185 WP 000190854.1 EERYGVDEVERLLDSCHALMNYGVDRYKRPQKISLQEEKARQKSREEYLQSQVNMLWRTLPKREEEKTVAEARRYPSEPQ 240
consensus
logo
NTDB id 75223 SEEA0014 RS18375 WP 001240762.1 ENLLYFMEKNAPLLEPWQREILRIVRKVSQYFYPQKQTQVMNEGWATFWHYTILNHLYDEGKVTERFMLEFLHSHTNVVF 320
NTDB id 1450 BW25113 RS06185 WP 000190854.1 ENLLYFMEKNAPLLESWQREILRIVRKVSQYFYPQKQTQVMNEGWATFWHYTILNHLYDEGKVTERFMLEFLHSHTNVVF 320
consensus
logo
NTDB id 75223 SEEA0014 RS18375 WP 001240762.1 QPPYNSPWYSGINPYALGFAMFQDIKRICQSPTDEDKYWFPDIAGSDWLETLHFAMRDFKDESFISQFLSPKVMRDFRFF 400
NTDB id 1450 BW25113 RS06185 WP 000190854.1 QPPYNSPWYSGINPYALGFAMFQDIKRICQSPTEEDKYWFPDIAGSDWLETLHFAMRDFKDESFISQFLSPKVMRDFRFF 400
consensus
logo
NTDB id 75223 SEEA0014 RS18375 WP 001240762.1 TVLDDDRHNYLEISAIHNEEGYREIRSKLSSQYNLSNLEPNIQVWNVDLRGDRSLTLRYIPHNRAPLDKGRKEVLKHVHR 480
NTDB id 1450 BW25113 RS06185 WP 000190854.1 TVLDDDRHNYLEISAIHNEEGYREIRNRLSSQYNLSNLEPNIQIWNVDLRGDRSLTLRYIPHNRAPLDRGRKEVLKHVHR 480
consensus
logo
NTDB id 75223 SEEA0014 RS18375 WP 001240762.1 LWGFDVMLEQQNEDGSVELLERCPPRMNML 510
NTDB id 1450 BW25113 RS06185 WP 000190854.1 LWGFDVMLEQQNEDGSIELLERCPPRMGNL 510
consensus

X non conserved
X similar
X ≥ 50% conserved