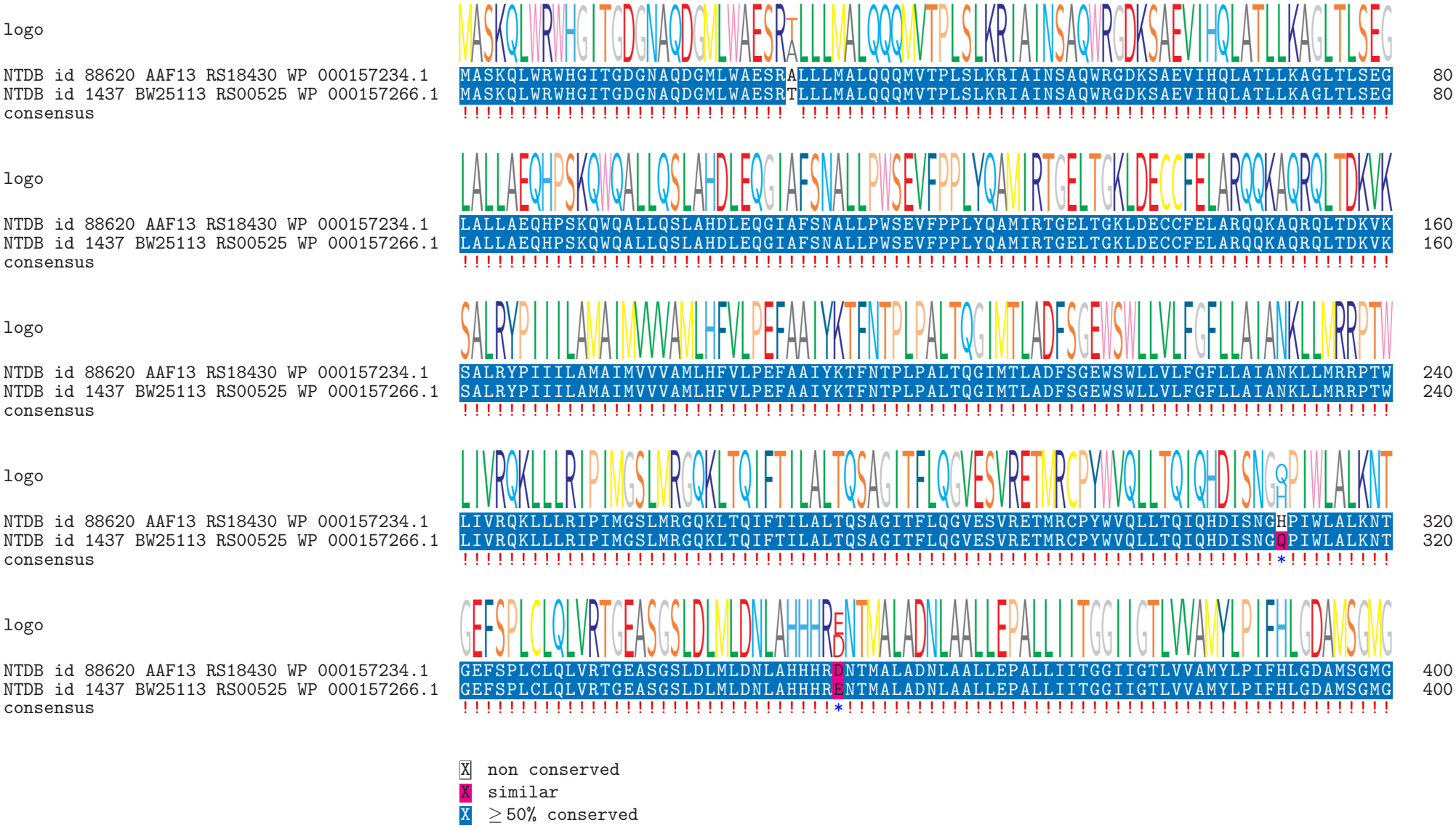

logo
NTDB id 88620 AAF13 RS18430 WP 000157234.1 MASKQLWRWHGITGDGNAQDGMLWAESRALLLMALQQQMVTPLSLKRIAINSAQWRGDKSAEVIHQLATLLKAGLTLSEG 80
NTDB id 1437 BW25113 RS00525 WP 000157266.1 MASKQLWRWHGITGDGNAQDGMLWAESRTLLLMALQQQMVTPLSLKRIAINSAQWRGDKSAEVIHQLATLLKAGLTLSEG 80
consensus
logo
NTDB id 88620 AAF13 RS18430 WP 000157234.1 LALLAEQHPSKQWQALLQSLAHDLEQGIAFSNALLPWSEVFPPLYQAMIRTGELTGKLDECCFELARQQKAQRQLTDKVK 160
NTDB id 1437 BW25113 RS00525 WP 000157266.1 LALLAEQHPSKQWQALLQSLAHDLEQGIAFSNALLPWSEVFPPLYQAMIRTGELTGKLDECCFELARQQKAQRQLTDKVK 160
consensus
logo
NTDB id 88620 AAF13 RS18430 WP 000157234.1 SALRYPIIILAMAIMVVVAMLHFVLPEFAAIYKTFNTPLPALTQGIMTLADFSGEWSWLLVLFGFLLAIANKLLMRRPTW 240
NTDB id 1437 BW25113 RS00525 WP 000157266.1 SALRYPIIILAMAIMVVVAMLHFVLPEFAAIYKTFNTPLPALTQGIMTLADFSGEWSWLLVLFGFLLAIANKLLMRRPTW 240
consensus
logo
NTDB id 88620 AAF13 RS18430 WP 000157234.1 LIVRQKLLLRIPIMGSLMRGQKLTQIFTILALTQSAGITFLQGVESVRETMRCPYWVQLLTQIQHDISNGHPIWLALKNT 320
NTDB id 1437 BW25113 RS00525 WP 000157266.1 LIVRQKLLLRIPIMGSLMRGQKLTQIFTILALTQSAGITFLQGVESVRETMRCPYWVQLLTQIQHDISNGQPIWLALKNT 320
consensus
logo
NTDB id 88620 AAF13 RS18430 WP 000157234.1 GEFSPLCLQLVRTGEASGSLDLMLDNLAHHHRDNTMALADNLAALLEPALLIITGGIIGTLVVAMYLPIFHLGDAMSGMG 400
NTDB id 1437 BW25113 RS00525 WP 000157266.1 GEFSPLCLQLVRTGEASGSLDLMLDNLAHHHRENTMALADNLAALLEPALLIITGGIIGTLVVAMYLPIFHLGDAMSGMG 400
consensus
X non conserved
X similar
X ≥ 50% conserved