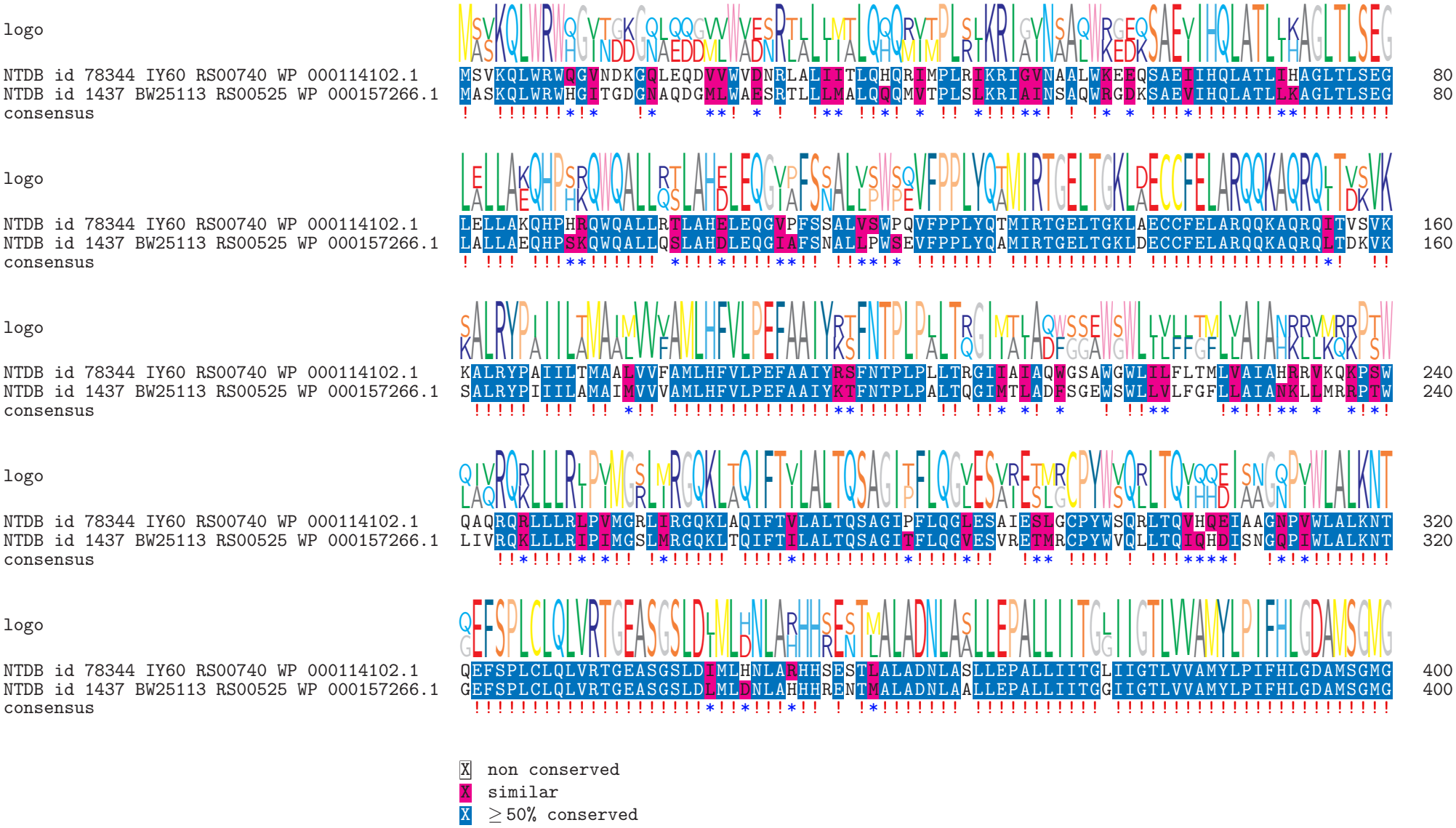

logo
NTDB id 78344 IY60 RS00740 WP 000114102.1 MSVKQLWRWQGVNDKGQLEQDVVWVDNRLALIITLQHQRIMPLRIKRIGVNAALWKEEQSAEIIHQLATLIHAGLTLSEG 80
NTDB id 1437 BW25113 RS00525 WP 000157266.1 MASKQLWRWHGITGDGNAQDGMLWAESRTLLLMALQQQMVTPLSLKRIAINSAQWRGDKSAEVIHQLATLLKAGLTLSEG 80
consensus
logo
NTDB id 78344 IY60 RS00740 WP 000114102.1 LELLAKQHPHRQWQALLRTLAHELEQGVPFSSALVSWPQVFPPLYQTMIRTGELTGKLAECCFELARQQKAQRQITVSVK 160
NTDB id 1437 BW25113 RS00525 WP 000157266.1 LALLAEQHPSKQWQALLQSLAHDLEQGIAFSNALLPWSEVFPPLYQAMIRTGELTGKLDECCFELARQQKAQRQLTDKVK 160
consensus
logo
NTDB id 78344 IY60 RS00740 WP 000114102.1 KALRYPAIILTMAALVVFAMLHFVLPEFAAIYRSFNTPLPLLTRGIIAIAQWGSAWGWLILFLTMLVAIAHRRVKQKPSW 240
NTDB id 1437 BW25113 RS00525 WP 000157266.1 SALRYPIIILAMAIMVVVAMLHFVLPEFAAIYKTFNTPLPALTQGIMTLADFSGEWSWLLVLFGFLLAIANKLLMRRPTW 240
consensus
logo
NTDB id 78344 IY60 RS00740 WP 000114102.1 QAQRQRLLLRLPVMGRLIRGQKLAQIFTVLALTQSAGIPFLQGLESAIESLGCPYWSQRLTQVHQEIAAGNPVWLALKNT 320
NTDB id 1437 BW25113 RS00525 WP 000157266.1 LIVRQKLLLRIPIMGSLMRGQKLTQIFTILALTQSAGITFLQGVESVRETMRCPYWVQLLTQIQHDISNGQPIWLALKNT 320
consensus
logo
NTDB id 78344 IY60 RS00740 WP 000114102.1 QEFSPLCLQLVRTGEASGSLDIMLHNLARHHSESTLALADNLASLLEPALLIITGLIIGTLVVAMYLPIFHLGDAMSGMG 400
NTDB id 1437 BW25113 RS00525 WP 000157266.1 GEFSPLCLQLVRTGEASGSLDLMLDNLAHHHRENTMALADNLAALLEPALLIITGGIIGTLVVAMYLPIFHLGDAMSGMG 400
consensus
X non conserved
X similar
X ≥ 50% conserved