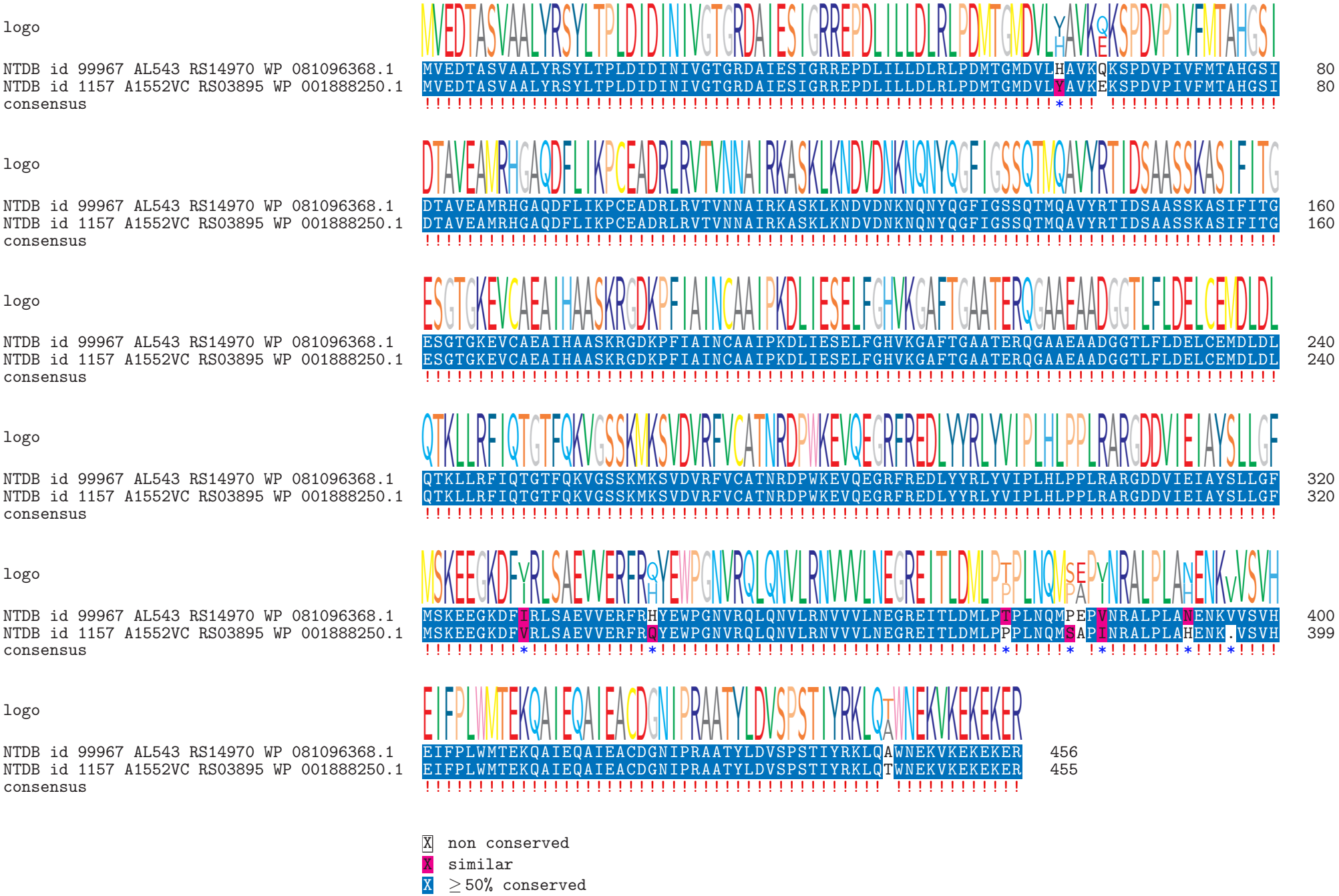

logo
NTDB id 99967 AL543 RS14970 WP 081096368.1 MVEDTASVAALYRSYLTPLDIDINIVGTGRDAIESIGRREPDLILLDLRLPDMTGMDVLHAVKQKSPDVPIVFMTAHGSI 80
NTDB id 1157 A1552VC RS03895 WP 001888250.1 MVEDTASVAALYRSYLTPLDIDINIVGTGRDAIESIGRREPDLILLDLRLPDMTGMDVLYAVKEKSPDVPIVFMTAHGSI 80
consensus
logo
NTDB id 99967 AL543 RS14970 WP 081096368.1 DTAVEAMRHGAQDFLIKPCEADRLRVTVNNAIRKASKLKNDVDNKNQNYQGFIGSSQTMQAVYRTIDSAASSKASIFITG 160
NTDB id 1157 A1552VC RS03895 WP 001888250.1 DTAVEAMRHGAQDFLIKPCEADRLRVTVNNAIRKASKLKNDVDNKNQNYQGFIGSSQTMQAVYRTIDSAASSKASIFITG 160
consensus
logo
NTDB id 99967 AL543 RS14970 WP 081096368.1 ESGTGKEVCAEAIHAASKRGDKPFIAINCAAIPKDLIESELFGHVKGAFTGAATERQGAAEAADGGTLFLDELCEMDLDL 240
NTDB id 1157 A1552VC RS03895 WP 001888250.1 ESGTGKEVCAEAIHAASKRGDKPFIAINCAAIPKDLIESELFGHVKGAFTGAATERQGAAEAADGGTLFLDELCEMDLDL 240
consensus
logo
NTDB id 99967 AL543 RS14970 WP 081096368.1 QTKLLRFIQTGTFQKVGSSKMKSVDVRFVCATNRDPWKEVQEGRFREDLYYRLYVIPLHLPPLRARGDDVIEIAYSLLGF 320
NTDB id 1157 A1552VC RS03895 WP 001888250.1 QTKLLRFIQTGTFQKVGSSKMKSVDVRFVCATNRDPWKEVQEGRFREDLYYRLYVIPLHLPPLRARGDDVIEIAYSLLGF 320
consensus
logo
NTDB id 99967 AL543 RS14970 WP 081096368.1 MSKEEGKDFIRLSAEVVERFRHYEWPGNVRQLQNVLRNVVVLNEGREITLDMLPTPLNQMPEPVNRALPLANENKVVSVH 400
NTDB id 1157 A1552VC RS03895 WP 001888250.1 MSKEEGKDFVRLSAEVVERFRQYEWPGNVRQLQNVLRNVVVLNEGREITLDMLPPPLNQMSAPINRALPLAHENK.VSVH 399
consensus
logo
NTDB id 99967 AL543 RS14970 WP 081096368.1 EIFPLWMTEKQAIEQAIEACDGNIPRAATYLDVSPSTIYRKLQAWNEKVKEKEKER 456
NTDB id 1157 A1552VC RS03895 WP 001888250.1 EIFPLWMTEKQAIEQAIEACDGNIPRAATYLDVSPSTIYRKLQTWNEKVKEKEKER 455
consensus
X non conserved
X similar
X ≥ 50% conserved