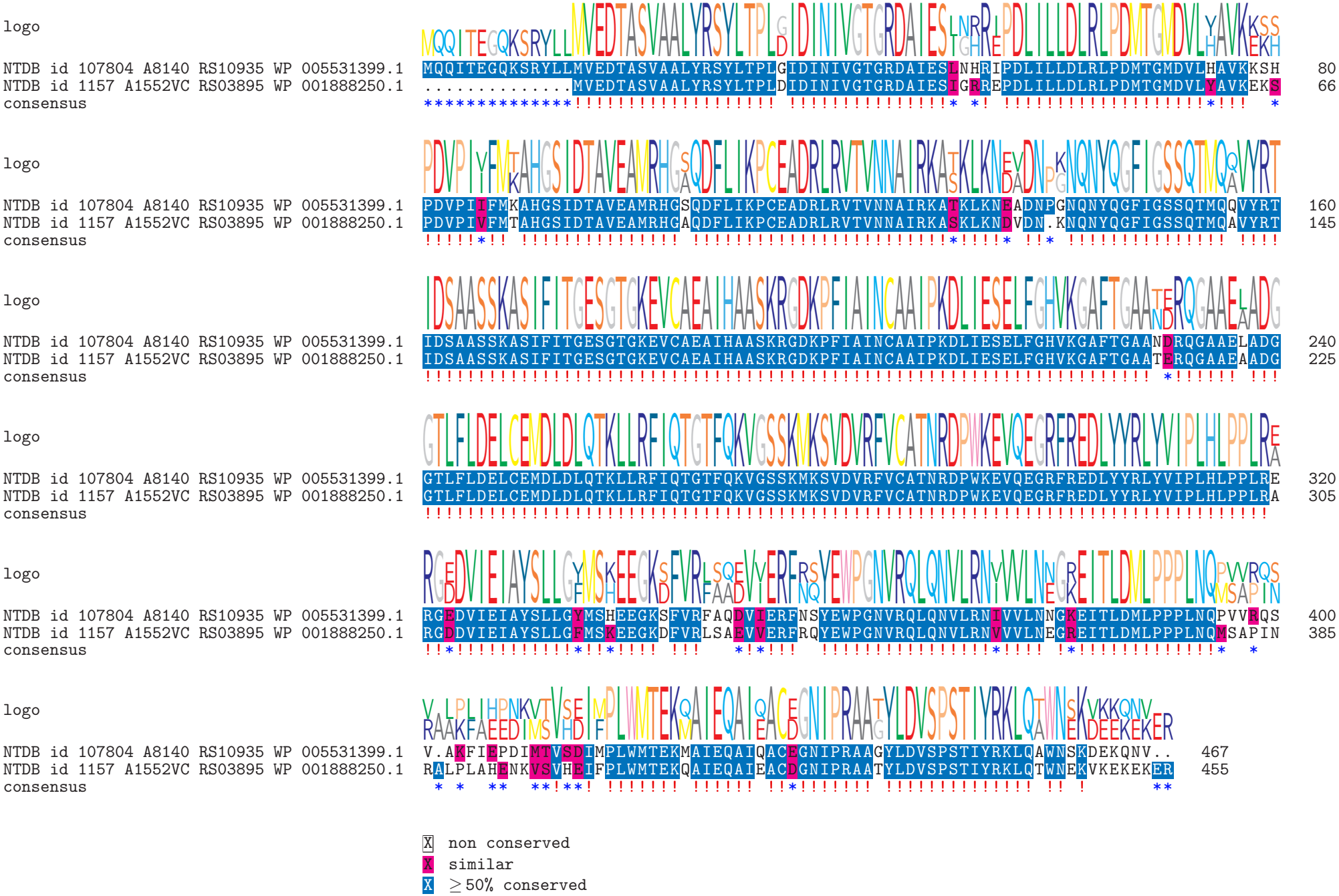

logo
NTDB id 107804 A8140 RS10935 WP 005531399.1 MQQITEGQKSRYLLMVEDTASVAALYRSYLTPLGIDINIVGTGRDAIESLNHRIPDLILLDLRLPDMTGMDVLHAVKKSH 80
NTDB id 1157 A1552VC RS03895 WP 001888250.1 .............MVEDTASVAALYRSYLTPLDIDINIVGTGRDAIESIGRREPDLILLDLRLPDMTGMDVLYAVKEKS 66
consensus
logo
NTDB id 107804 A8140 RS10935 WP 005531399.1 PDVPIIFMKAHGSIDTAVEAMRHGSQDFLIKPCEADRLRVTVNNAIRKATKLKNEADNPGNQNYQGFIGSSQTMQQVYRT 160
NTDB id 1157 A1552VC RS03895 WP 001888250.1 PDVPIVFMTAHGSIDTAVEAMRHGAQDFLIKPCEADRLRVTVNNAIRKASKLKNDVDN.KNQNYQGFIGSSQTMQAVYRT 145
consensus
logo
NTDB id 107804 A8140 RS10935 WP 005531399.1 IDSAASSKASIFITGESGTGKEVCAEAIHAASKRGDKPFIAINCAAIPKDLIESELFGHVKGAFTGAANDRQGAAELADG 240
NTDB id 1157 A1552VC RS03895 WP 001888250.1 IDSAASSKASIFITGESGTGKEVCAEAIHAASKRGDKPFIAINCAAIPKDLIESELFGHVKGAFTGAATERQGAAEAADG 225
consensus
logo
NTDB id 107804 A8140 RS10935 WP 005531399.1 GTLFLDELCEMDLDLQTKLLRFIQTGTFQKVGSSKMKSVDVRFVCATNRDPWKEVQEGRFREDLYYRLYVIPLHLPPLRE 320
NTDB id 1157 A1552VC RS03895 WP 001888250.1 GTLFLDELCEMDLDLQTKLLRFIQTGTFQKVGSSKMKSVDVRFVCATNRDPWKEVQEGRFREDLYYRLYVIPLHLPPLRA 305
consensus
logo
NTDB id 107804 A8140 RS10935 WP 005531399.1 RGEDVIEIAYSLLGYMSHEEGKSFVRFAQDVIERFNSYEWPGNVRQLQNVLRNIVVLNNGKEITLDMLPPPLNQPVVRQS 400
NTDB id 1157 A1552VC RS03895 WP 001888250.1 RGDDVIEIAYSLLGFMSKEEGKDFVRLSAEVVERFRQYEWPGNVRQLQNVLRNVVVLNEGREITLDMLPPPLNQMSAPIN 385
consensus
logo
NTDB id 107804 A8140 RS10935 WP 005531399.1 V.AKFIEPDIMTVSDIMPLWMTEKMAIEQAIQACEGNIPRAAGYLDVSPSTIYRKLQAWNSKDEKQNV.. 467
NTDB id 1157 A1552VC RS03895 WP 001888250.1 RALPLAHENKVSVHEIFPLWMTEKQAIEQAIEACDGNIPRAATYLDVSPSTIYRKLQTWNEKVKEKEKER 455
consensus
X non conserved
X similar
X ≥ 50% conserved