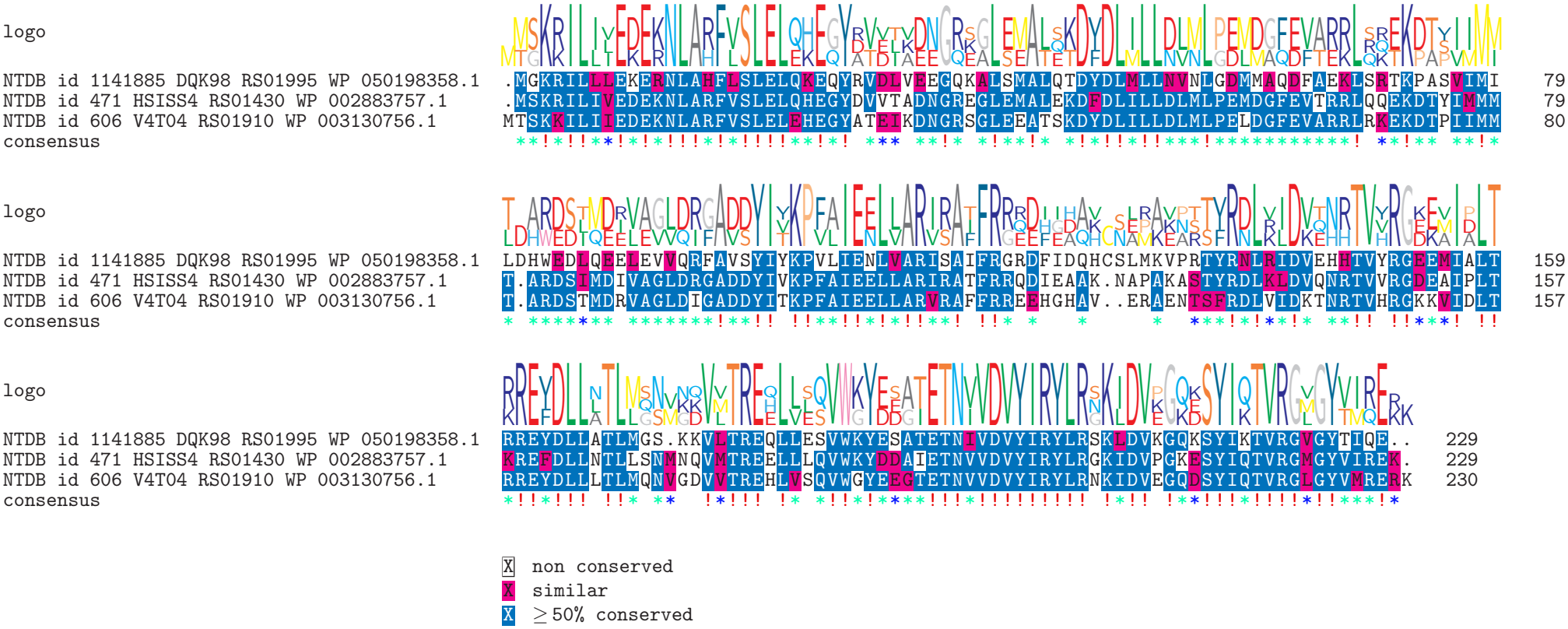
logo
NTDB id 1141885 DQK98 RS01995 WP 050198358.1 .MGKRILLLEKERNLAHFLSLELQKEQYRVDLVEEGQKALSMALQTDYDLMLLNVNLGDMMAQDFAEKLSRTKPASVIMI 79
NTDB id 471 HSISS4 RS01430 WP 002883757.1 .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDTYIMMM 79
NTDB id 606 V4T04 RS01910 WP 003130756.1 MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDTPIIMM 80
consensus
logo
NTDB id 1141885 DQK98 RS01995 WP 050198358.1 LDHWEDLQEELEVVQRFAVSYIYKPVLIENLVARISAIFRGRDFIDQHCSLMKVPRTYRNLRIDVEHHTVYRGEEMIALT 159
NTDB id 471 HSISS4 RS01430 WP 002883757.1 T.ARDSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAAK.NAPAKASTYRDLKLDVQNRTVVRGDEAIPLT 157
NTDB id 606 V4T04 RS01910 WP 003130756.1 T.ARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAV..ERAENTSFRDLVIDKTNRTVHRGKKVIDLT 157
consensus
logo
NTDB id 1141885 DQK98 RS01995 WP 050198358.1 RREYDLLATLMGS.KKVLTREQLLESVWKYESATETNIVDVYIRYLRSKLDVKGQKSYIKTVRGVGYTIQE.. 229
NTDB id 471 HSISS4 RS01430 WP 002883757.1 KREFDLLNTLLSNMNQVMTREELLLQVWKYDDAIETNVVDVYIRYLRGKIDVPGKESYIQTVRGMGYVIREK. 229
NTDB id 606 V4T04 RS01910 WP 003130756.1 RREYDLLLTLMQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK 230
consensus
X non conserved
X similar
X ≥ 50% conserved