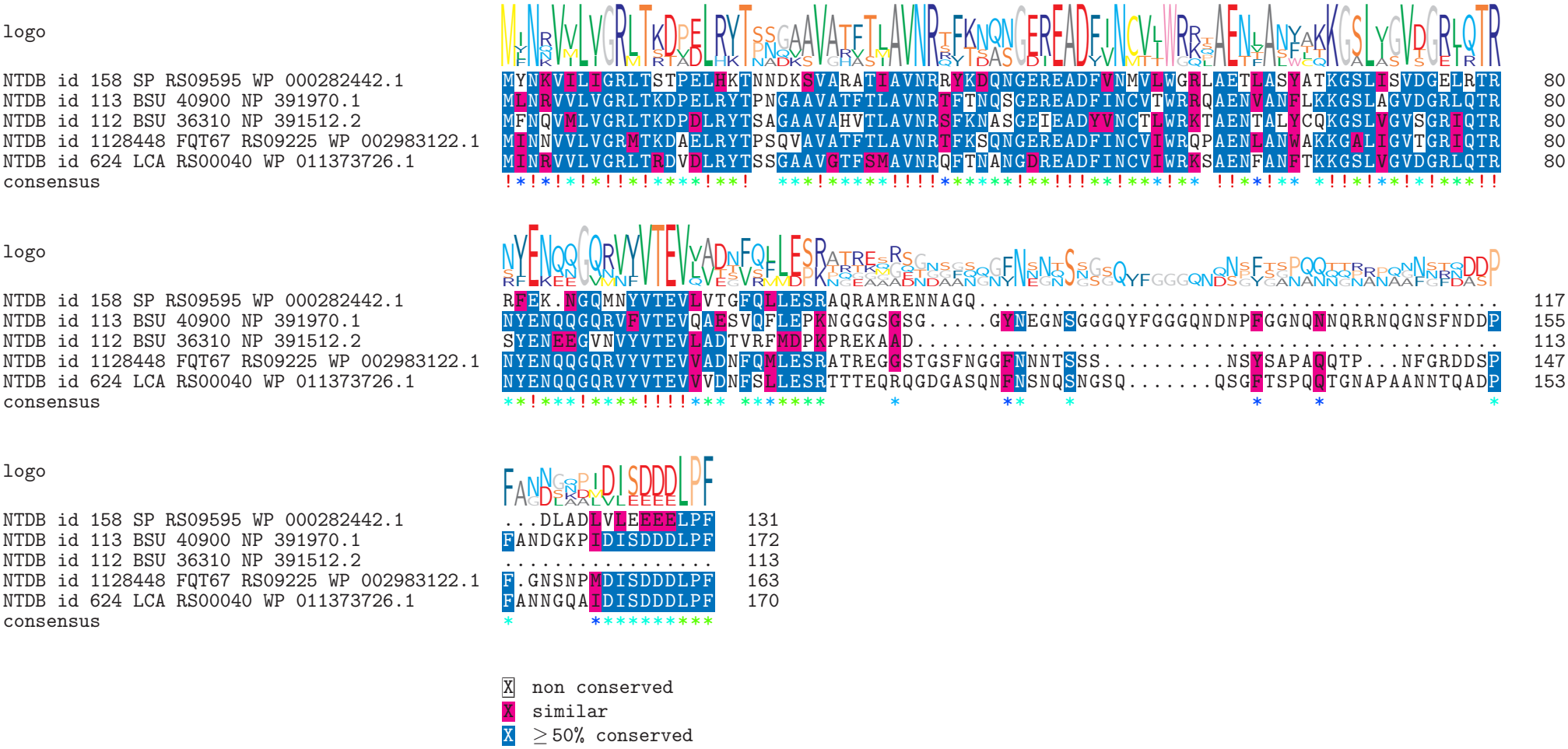
logo
NTDB id 158 SP RS09595 WP 000282442.1 MYNKVILIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR 80
NTDB id 113 BSU 40900 NP 391970.1 MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTNQSGEREADFINCVTWRRQAENVANFLKKGSLAGVDGRLQTR 80
NTDB id 112 BSU 36310 NP 391512.2 MFNQVMLVGRLTKDPDLRYTSAGAAVAHVTLAVNRSFKNASGEIEADYVNCTLWRKTAENTALYCQKGSLVGVSGRIQTR 80
NTDB id 1128448 FQT67 RS09225 WP 002983122.1 MINNVVLVGRMTKDAELRYTPSQVAVATFTLAVNRTFKSQNGEREADFINCVIWRQPAENLANWAKKGALIGVTGRIQTR 80
NTDB id 624 LCA RS00040 WP 011373726.1 MINRVVLVGRLTRDVDLRYTSSGAAVGTFSMAVNRQFTNANGDREADFINCVIWRKSAENFANFTKKGSLVGVDGRLQTR 80
consensus
logo
NTDB id 158 SP RS09595 WP 000282442.1 RFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQ 117
NTDB id 113 BSU 40900 NP 391970.1 NYENQQGQRVFVTEVQAESVQFLEPKNGGGSGSG.....GYNEGNSGGGQYFGGGQNDNPFGGNQNNQRRNQGNSFNDDP 155
NTDB id 112 BSU 36310 NP 391512.2 SYENEEGVNVYVTEVLADTVRFMDPKPREKAAD 113
NTDB id 1128448 FQT67 RS09225 WP 002983122.1 NYENQQGQRVYVTEVVADNFQMLESRATREGGSTGSFNGGFNNNTSSS.....NSYSAPAQQTP...NFGRDDSP 147
NTDB id 624 LCA RS00040 WP 011373726.1 NYENQQGQRVYVTEVVVDNFSLLESRTTTEQRQGDGASQNFNSNQSNGSQ.....QSGFTSPQQTGNAPAANNTQADP 153
consensus
logo
NTDB id 158 SP RS09595 WP 000282442.1 ...DLADLVLEEEELPF 131
NTDB id 113 BSU 40900 NP 391970.1 FANDGKPIDISDDDLPF 172
NTDB id 112 BSU 36310 NP 391512.2 113
NTDB id 1128448 FQT67 RS09225 WP 002983122.1 F.GNSNPMDISDDDLPF 163
NTDB id 624 LCA RS00040 WP 011373726.1 FANNGQAIDISDDDLPF 170
consensus
non conserved
similar
≥ 50% conserved