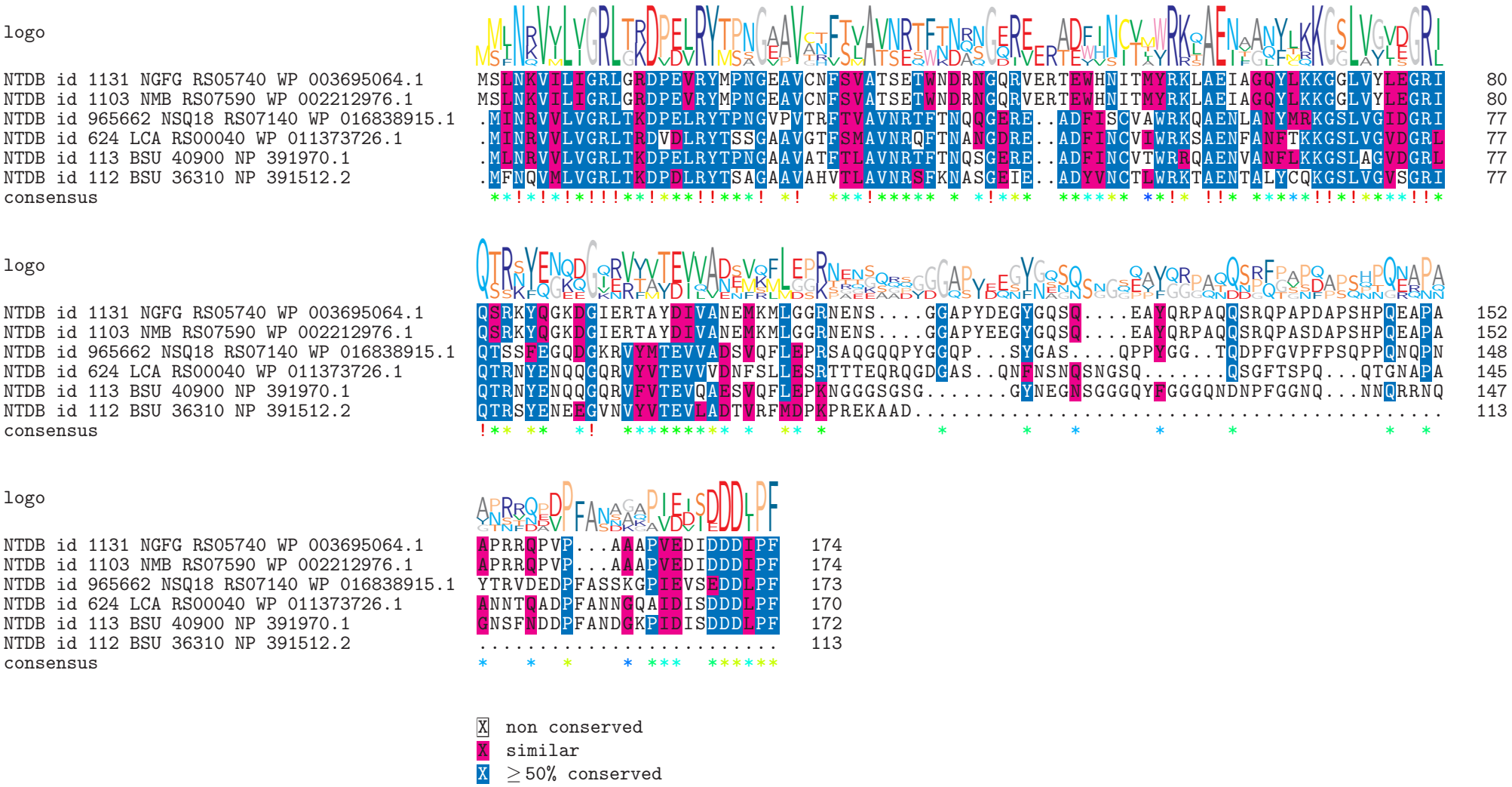
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRNGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYLEGRI 80
NTDB id 1103 NMB RS07590 WP 002212976.1 MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRNGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYLEGRI 80
NTDB id 965662 NSQ18 RS07140 WP 016838915.1 .MINRVVLVGRLTKDPELRYTPNGVPVTRFTVAVNRTFTNQQGERE..ADFISCVAWRKQAENLANYMRKGSLVGIDGRI 77
NTDB id 624 LCA RS00040 WP 011373726.1 .MINRVVLVGRLTRDVDLRYTSSGAAVGTFSMAVNRQFTNANGDRE..ADFINCVIWRKSAENFANFTKKGSLVGVDGRL 77
NTDB id 113 BSU 40900 NP 391970.1 .MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTNQSGERE..ADFINCVTWRRQAENVANFLKKGSLAGVDGRL 77
NTDB id 112 BSU 36310 NP 391512.2 .MFNQVMLVGRLTKDPDLRYTSAGAAVAHVTLAVNRSFKNASGEIE..ADYVNCTLWRKTAENTALYCQKGSLVGVSGRI 77
consensus
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 QSRKYQGKDGIERTAYDIVANEMKMLGGRNENS....GGAPYDEGYGQSQ....EAYQRPAQQSRQPAPDAPSHPQEAPA 152
NTDB id 1103 NMB RS07590 WP 002212976.1 QSRKYQGKDGIERTAYDIVANEMKMLGGRNENS....GGAPYEEGYGQSQ....EAYQRPAQQSRQPASDAPSHPQEAPA 152
NTDB id 965662 NSQ18 RS07140 WP 016838915.1 QTSSFEGQDGKRVYMTEVVADSVQFLEPRSAQGQQPYGGQP...SYGAS....QPPYGG..TQDPFGVPFPSQPPQNQPN 148
NTDB id 624 LCA RS00040 WP 011373726.1 QTRNYENQQGQRVYVTEVVVDNFSLLESRTTTEQRQGDGAS..QNFNSNQSNGSQ.......QSGFTSPQ...QTGNAPA 145
NTDB id 113 BSU 40900 NP 391970.1 QTRNYENQQGQRVFVTEVQAESVQFLEPKNGGGSGSG.......GYNEGNSGGGQYFGGGQNDNPFGGNQ...NNQRRNQ 147
NTDB id 112 BSU 36310 NP 391512.2 QTRSYENEEGVNVYVTEVLADTVRFMDPKPREKAAD..... 113
consensus
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 APRRQPVP...AAAPVEDIDDDIPF 174
NTDB id 1103 NMB RS07590 WP 002212976.1 APRRQPVP...AAAPVEDIDDDIPF 174
NTDB id 965662 NSQ18 RS07140 WP 016838915.1 YTRVDEDPFASSKGPIEVSEDDLPF 173
NTDB id 624 LCA RS00040 WP 011373726.1 ANNTQADPFANNGQAIDISDDDLPF 170
NTDB id 113 BSU 40900 NP 391970.1 GNSFNDDPFANDGKPIDISDDDLPF 172
NTDB id 112 BSU 36310 NP 391512.2 113
consensus
X non conserved
X similar
X ≥ 50% conserved