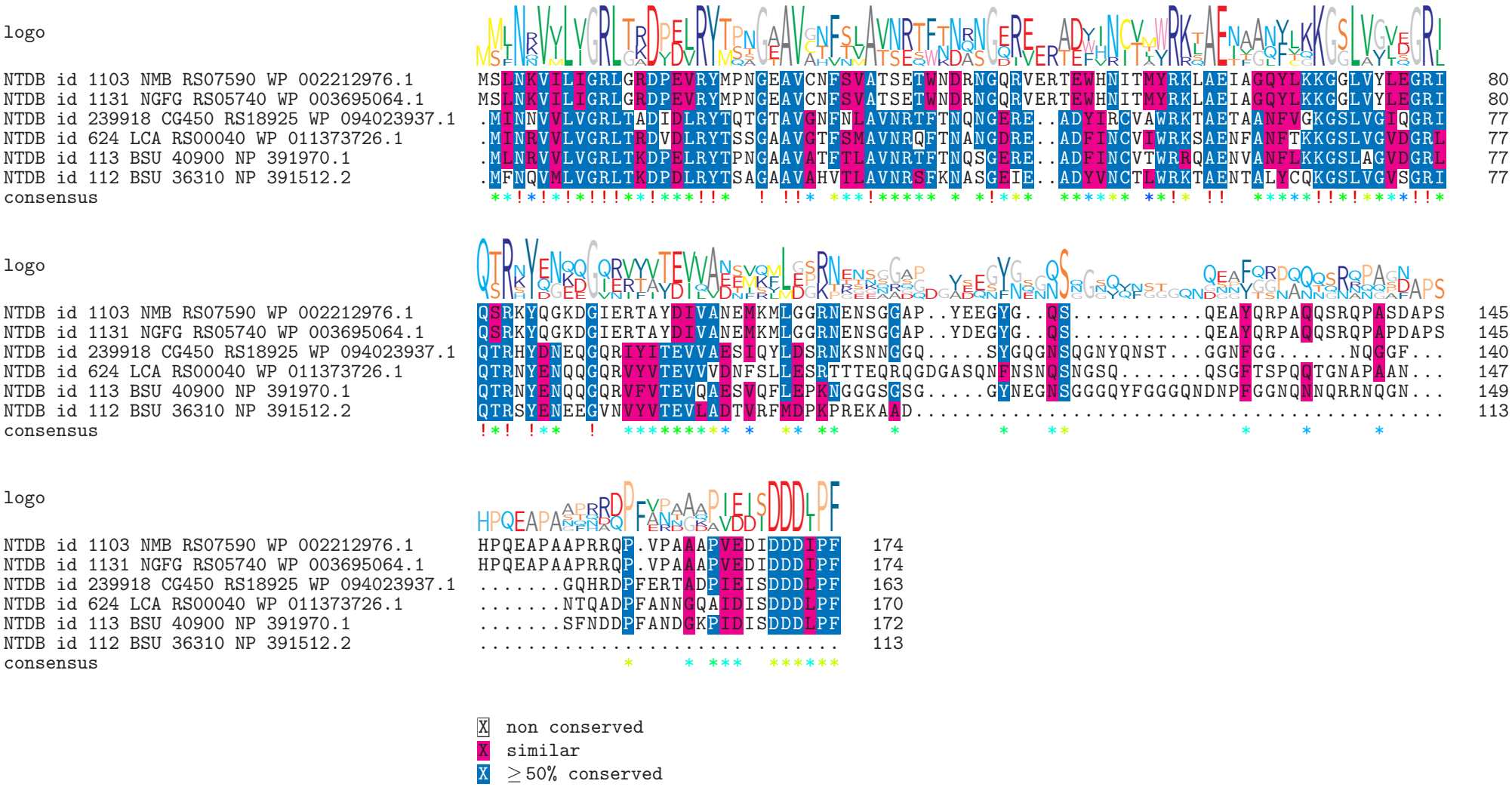
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRNGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYLEGRI 80
NTDB id 1131 NGFG RS05740 WP 003695064.1 MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRNGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYLEGRI 80
NTDB id 239918 CG450 RS18925 WP 094023937.1 .MINNVVLVGRLTADIDLRYTQTGTAVGNFNLAVNRTFTNQNGERE..ADYIRCVAWRKTAETAANFVGKGSLVGIQGRI 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 1103 NMB RS07590 WP 002212976.1 QSRKYQGKDGIERTAYDIVANEMKMLGGRNENSGGAP..YEEGYG..QS.....QEAYQRPAQQSRQPASDAPS 145
NTDB id 1131 NGFG RS05740 WP 003695064.1 QSRKYQGKDGIERTAYDIVANEMKMLGGRNENSGGAP..YDEGYG..QS.....QEAYQRPAQQSRQPAPDAPS 145
NTDB id 239918 CG450 RS18925 WP 094023937.1 QTRHYDNEQGQRIYITEVVAESIQYLDSRNKSNNGGQ.....SYGQGNSQGNYQNST...GGNFGG.....NQGGF... 140
NTDB id 624 LCA RS00040 WP 011373726.1 QTRNYENQQGQRVYVTEVVVDNFSLLESRTTTEQRQGDGASQNFNSNQSNGSQ.....QSGFTSPQQTGNAPAAN... 147
NTDB id 113 BSU 40900 NP 391970.1 QTRNYENQQGQRVFVTEVQAESVQFLEPKNGGGSGSG.....GYNEGNSGGGQYFGGGQNDNPFGGNQNNQRRNQGN... 149
NTDB id 112 BSU 36310 NP 391512.2 QTRSYENEEGVNVYVTEVLADTVRFMDPKPREKAAD..... 113
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 HPQEAPAAPRRQP.VPAAAPVEDIDDDIPF 174
NTDB id 1131 NGFG RS05740 WP 003695064.1 HPQEAPAAPRRQP.VPAAAPVEDIDDDIPF 174
NTDB id 239918 CG450 RS18925 WP 094023937.1 .......GQHRDPFERTADPIEISDDDLPF 163
NTDB id 624 LCA RS00040 WP 011373726.1 .......NTQADPFANNGQAIDISDDDLPF 170
NTDB id 113 BSU 40900 NP 391970.1 .......SFNDDPFANDGKPIDISDDDLPF 172
NTDB id 112 BSU 36310 NP 391512.2 113
consensus
X non conserved
X similar
X ≥ 50% conserved