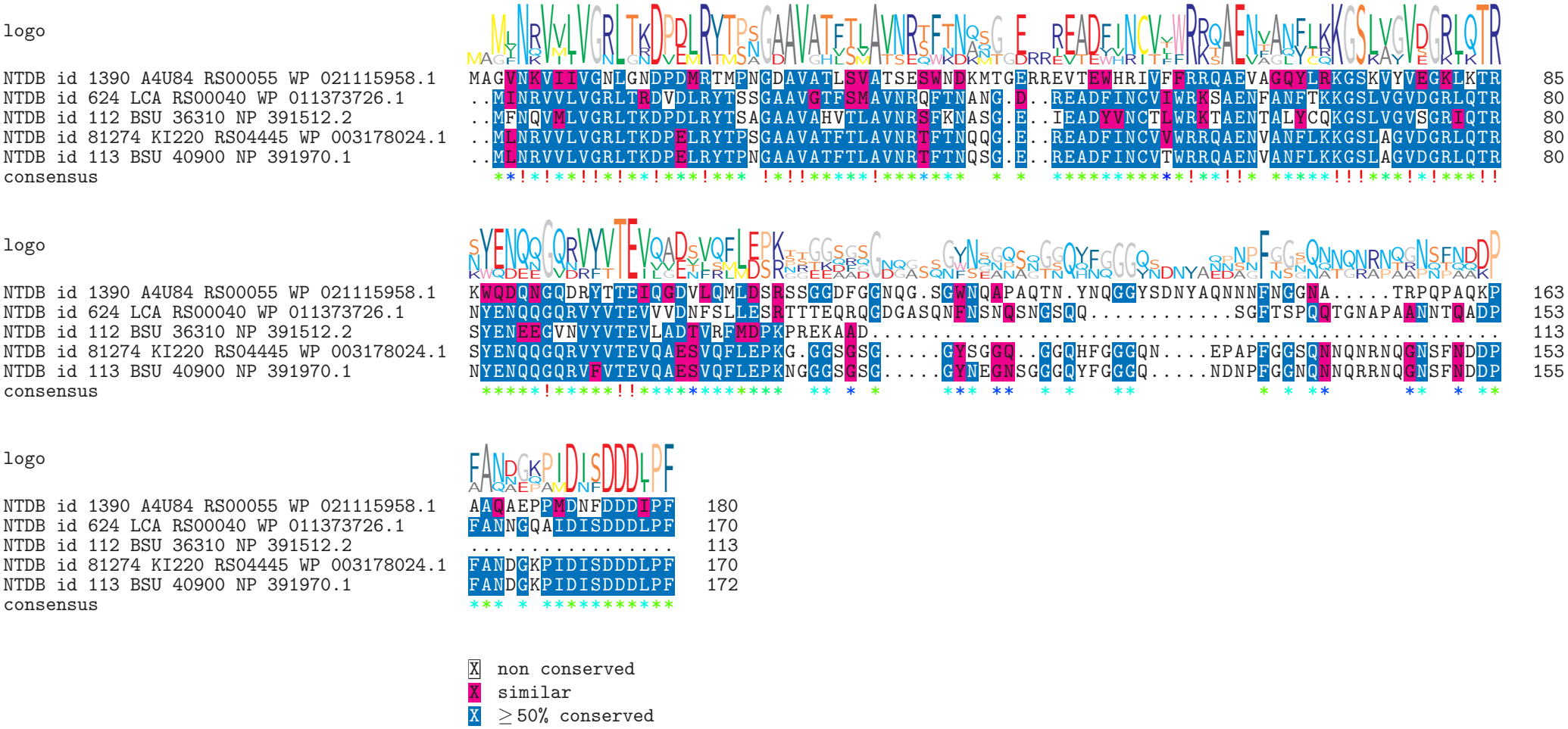
logo
NTDB id 1390 A4U84 RS00055 WP 021115958.1 MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYVEGKLKTR 85
NTDB id 624 LCA RS00040 WP 011373726.1 ..MINRVVLVGRLTRDVDLRYTSSGAAVGTFSMAVNRQFTNANG.D..READFINCVIWRKSAENFANFTKKGSLVGVDGRLQTR 80
NTDB id 112 BSU 36310 NP 391512.2 ..MFNQVMLVGRLTKDPDLRYTSAGAAVAHVTLAVNRSFKNASG.E..IEADYVNCTLWRKTAENTALYCQKGSLVGVSGRIQTR 80
NTDB id 81274 KI220 RS04445 WP 003178024.1 ..MLNRVVLVGRLTKDPELRYTPSGAAVATFTLAVNRTFTNQQG.E..READFINCVVWRRQAENVANFLKKGSLAGVDGRLQTR 80
NTDB id 113 BSU 40900 NP 391970.1 ..MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTNQSG.E..READFINCVTWRRQAENVANFLKKGSLAGVDGRLQTR 80
consensus
logo
NTDB id 1390 A4U84 RS00055 WP 021115958.1 KWQDQNGQDRYTTEIQGDVLQMLDSRSSGGDFGGNQG.SGWNQAPAQTN.YNQGGYSDNYAQNNNFNGGNA.....TRPQPAQKP 163
NTDB id 624 LCA RS00040 WP 011373726.1 NYENQQGQRVYVTEVVVDNFSLLESRTTTEQRQGDGASQNFNSNQSNGSQQ.....SGFTSPQQTGNAPAANNTQADP 153
NTDB id 112 BSU 36310 NP 391512.2 SYENEEGVNVYVTEVLADTVRFMDPKPREKAAD..... 113
NTDB id 81274 KI220 RS04445 WP 003178024.1 SYENQQGQRVYVTEVQAESVQFLEPKG.GGSGSG.....GYSGGQ..GGQHFGGGQN....EPAPFGGSQNNQNRNQGNSFNDDP 153
NTDB id 113 BSU 40900 NP 391970.1 NYENQQGQRVFVTEVQAESVQFLEPKNGGGSGSG.....GYNEGNSGGQQYFGGGQ.....NDNPFGGNQNNQRRNQGNSFNDDP 155
consensus
logo
NTDB id 1390 A4U84 RS00055 WP 021115958.1 AAQAEPPMDNFDDDIPF 180
NTDB id 624 LCA RS00040 WP 011373726.1 FANNGQAIDISDDDLPF 170
NTDB id 112 BSU 36310 NP 391512.2 113
NTDB id 81274 KI220 RS04445 WP 003178024.1 FANDGKPIDISDDDLPF 170
NTDB id 113 BSU 40900 NP 391970.1 FANDGKPIDISDDDLPF 172
consensus
X non conserved
X similar
X ≥ 50% conserved