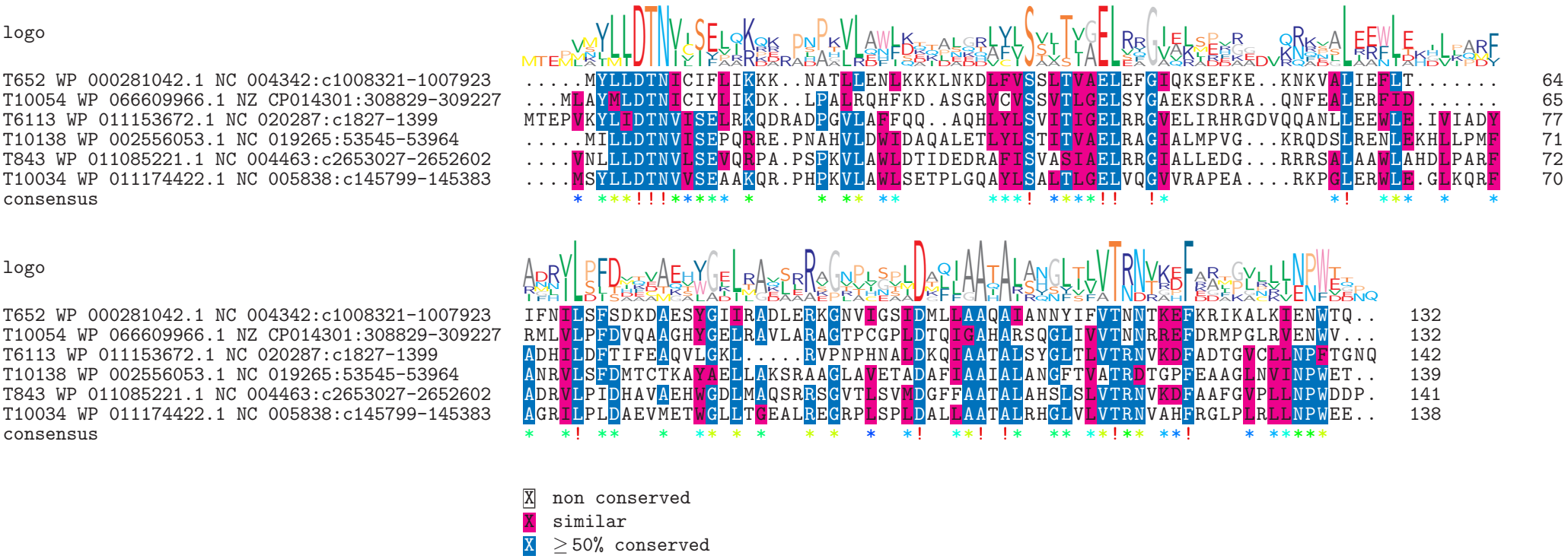
logo
T652 WP 000281042.1 NC 004342:c1008321-1007923 .....MYLLDTNICIFLIKKK..NATLLENLKKKLNKDLFVSSLTVAELEFGIQKSEFKE..KNKVALIEFLT....... 64
T10054 WP 066609966.1 NZ CP014301:308829-309227 ...MLAYMLDTNICIYLIKDK..LPALRQHFKD.ASGRVCVSSVTLGELSYGAEKSDRRA..QNFEALERFID....... 65
T6113 WP 011153672.1 NC 020287:c1827-1399 MTEPVKYLIDTNVISELRKQDRADPGVLAFFQQ..AQHLYLSVITIGELRRGVELIRHRGDVQQANLLEEWLE.IVIADY 77
T10138 WP 002556053.1 NC 019265:53545-53964 .....MILLDTNVISEPQRRE.PNAHVLDWIDAQALETLYLSTITVAELRAGIALMPVG...KRQDSLRENLEKHLLPMF 71
T843 WP 011085221.1 NC 004463:c2653027-2652602 ....VNLLLDTNVLSEVQRPA.PSPKVLAWLDTIDEDRAFISVASIAELRRGIALLEDG...RRRSALAAWLAHDLPARF 72
T10034 WP 011174422.1 NC 005838:c145799-145383 ....MSYLLDTNVVSEAAKQR.PHPKVLAWLSETPLGQAYLSALTLGELVQGVVRAPEA....RKPGLERWLE.GLKQRF 70
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T652 WP 000281042.1 NC 004342:c1008321-1007923 IFNILSFSDKDAESYGIIRADLERKGNVIGSIDMLLAAQAIANNYIFVTNNTKEFKRIKALKIENWTQ.. 132
T10054 WP 066609966.1 NZ CP014301:308829-309227 RMLVLPFDVQAAGHYGELRAVLARAGTPCGPLDTQIGAHARSQGLIVVTNNRREFDRMPGLRVENWV... 132
T6113 WP 011153672.1 NC 020287:c1827-1399 ADHILDFTIFEAQVLGKL.....RVPNPHNALDKQIAATALSYGLTLVTRNVKDFADTGVCLLNPFTGNQ 142
T10138 WP 002556053.1 NC 019265:53545-53964 ANRVLSFDMTCTKAYAELLAKSRAAGLAVETADAFIAAIALANGFTVATRDTGPFEAAGLNVINPWET.. 139
T843 WP 011085221.1 NC 004463:c2653027-2652602 ADRVLPIDHAVAEHWGDLMAQSRRSGVTLSVMDGFFAATALAHSLSLVTRNVKDFAAFGVPLLNPWDDP. 141
T10034 WP 011174422.1 NC 005838:c145799-145383 AGRILPLDAEVMETWGLLTGEALREGRPLSPLDALLAATALRHGLVLVTRNVAHFRGLPLRLLNPWEE.. 138
consensus
X non conserved
X similar
X ≥ 50% conserved