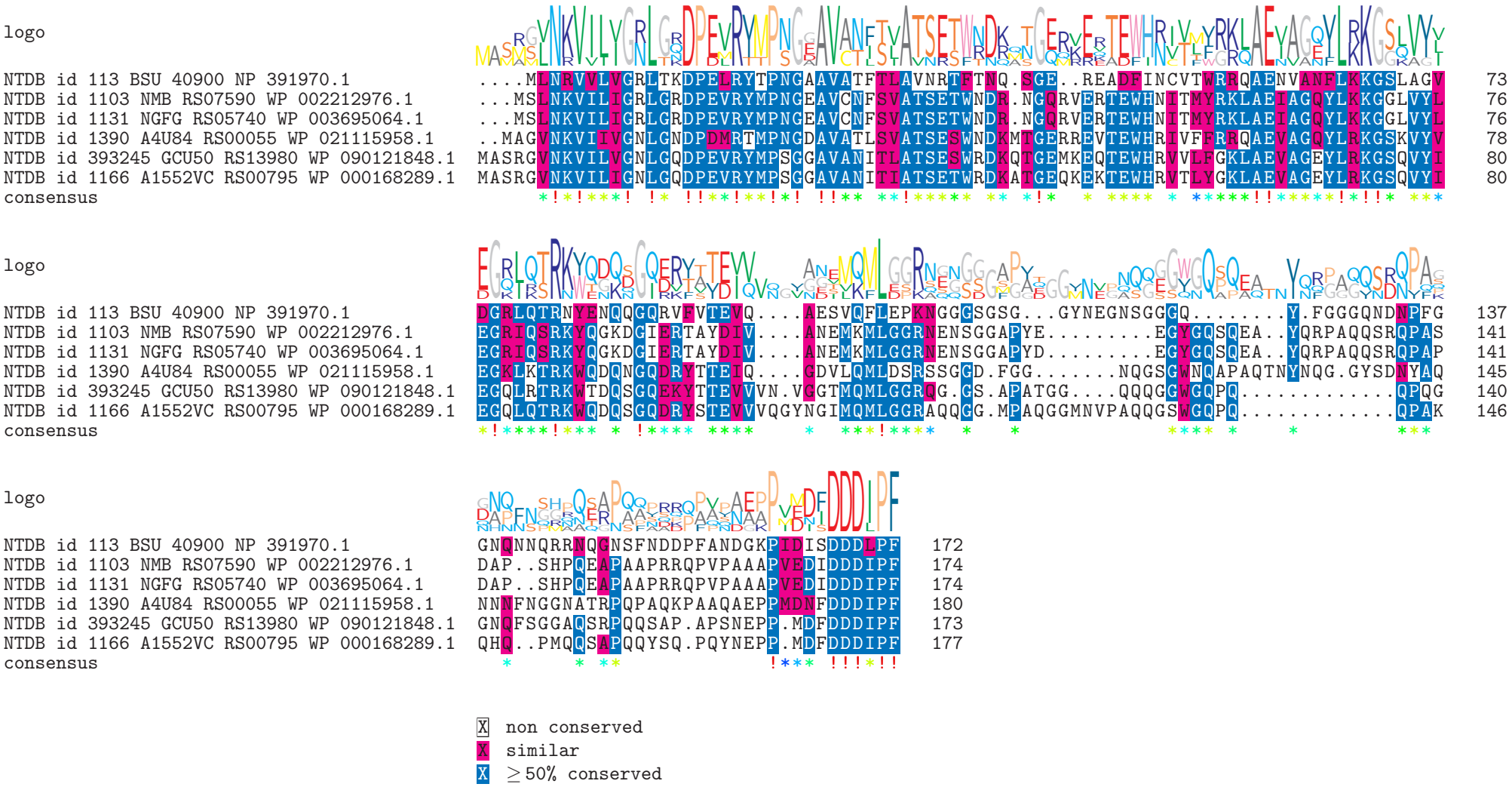
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NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
NTDB id 393245 GCU50 RS13980 WP 090121848.1 MASRGVNKVILVGNLGQDPEVRYMPSGGAVANITLATSESWRDKQTGEMKEQTEWHRVVLFGKLAEVAGEYLRKGSQVYI 80
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consensus
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NTDB id 113 BSU 40900 NP 391970.1 DGRLQTRNYENQQGQRVFVTEVQ....AESVQFLEPKNGGGSGSG...GYNEGNSGGGQ.....Y.FGGGQNDNPFG 137
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consensus
X non conserved
X similar
X ≥ 50% conserved