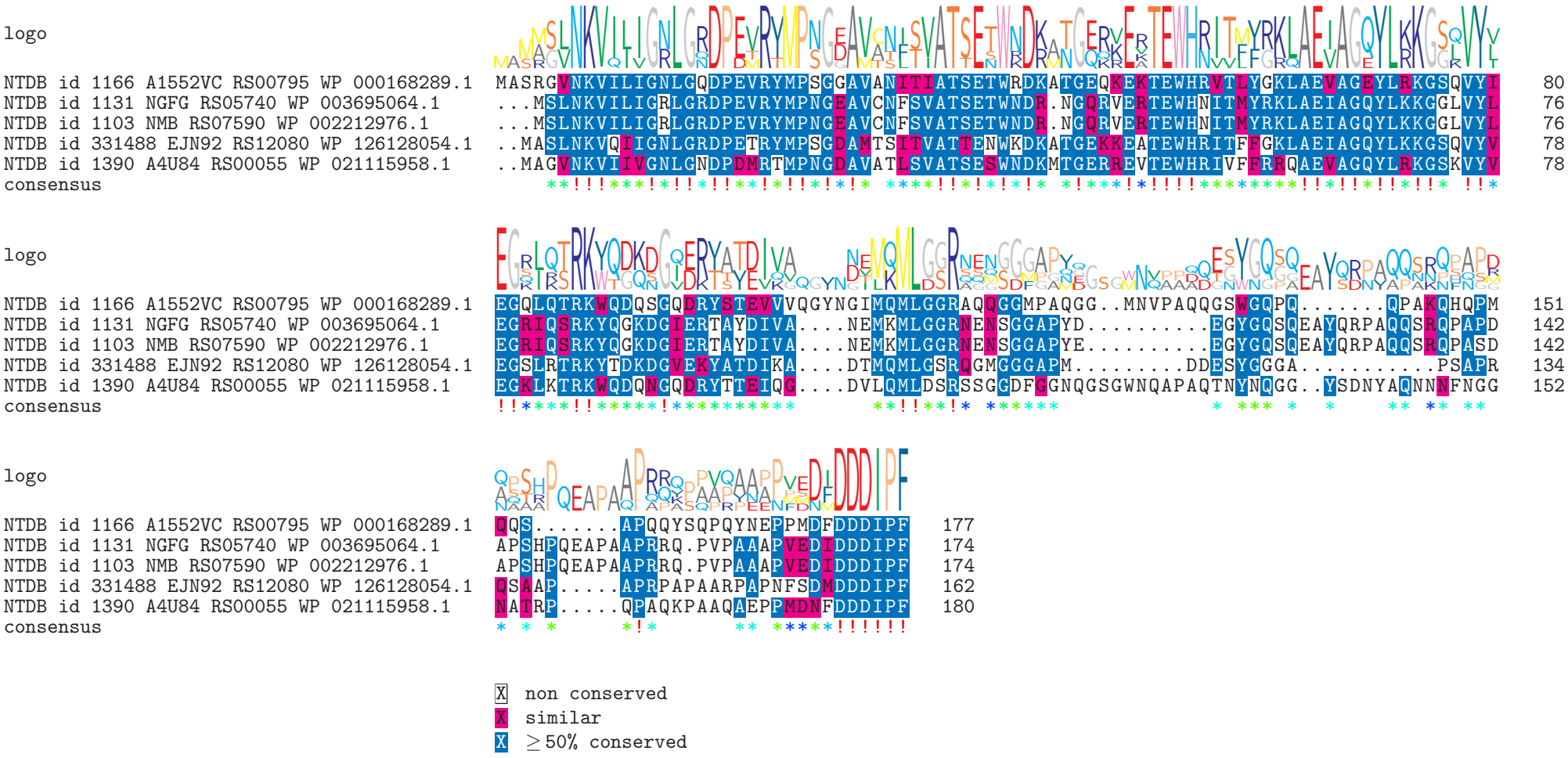
logo
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 331488 EJN92 RS12080 WP 126128054.1 ..MASLNKVQIIGNLGRDPETRYMPSGDAMTSITVATTENWKDKATGEKKEATEWHRITFFGKLAEIAGQYLKKGSQVYV 78
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
consensus
logo
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGG..MNVPAQQGSWGQPQ.......QPAKQHQPM 151
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYD..........EGYGQSQEAYQRPAQQSRQPAPD 142
NTDB id 1103 NMB RS07590 WP 002212976.1 EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYE..........EGYGQSQEAYQRPAQQSRQPASD 142
NTDB id 331488 EJN92 RS12080 WP 126128054.1 EGSLRTRKYTDKDGVEKYATDIKA....DTMQMLGSRQGMGGGAPM.........DDESYGGGA...........PSAPR 134
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGG..YSDNYAQNNNFNGG 152
consensus
logo
NTDB id 1166 A1552VC RS00795 WP 000168289.1 QQS.......APQQYSQPQYNEPPMDFDDDIPF 177
NTDB id 1131 NGFG RS05740 WP 003695064.1 APSHPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF 174
NTDB id 1103 NMB RS07590 WP 002212976.1 APSHPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF 174
NTDB id 331488 EJN92 RS12080 WP 126128054.1 QSAAP.....APRPAPAARPAPNFSDMDDDIPF 162
NTDB id 1390 A4U84 RS00055 WP 021115958.1 NATRP.....QPAQKPAAQAEPPMDNFDDDIPF 180
consensus
X non conserved
X similar
X ≥ 50% conserved