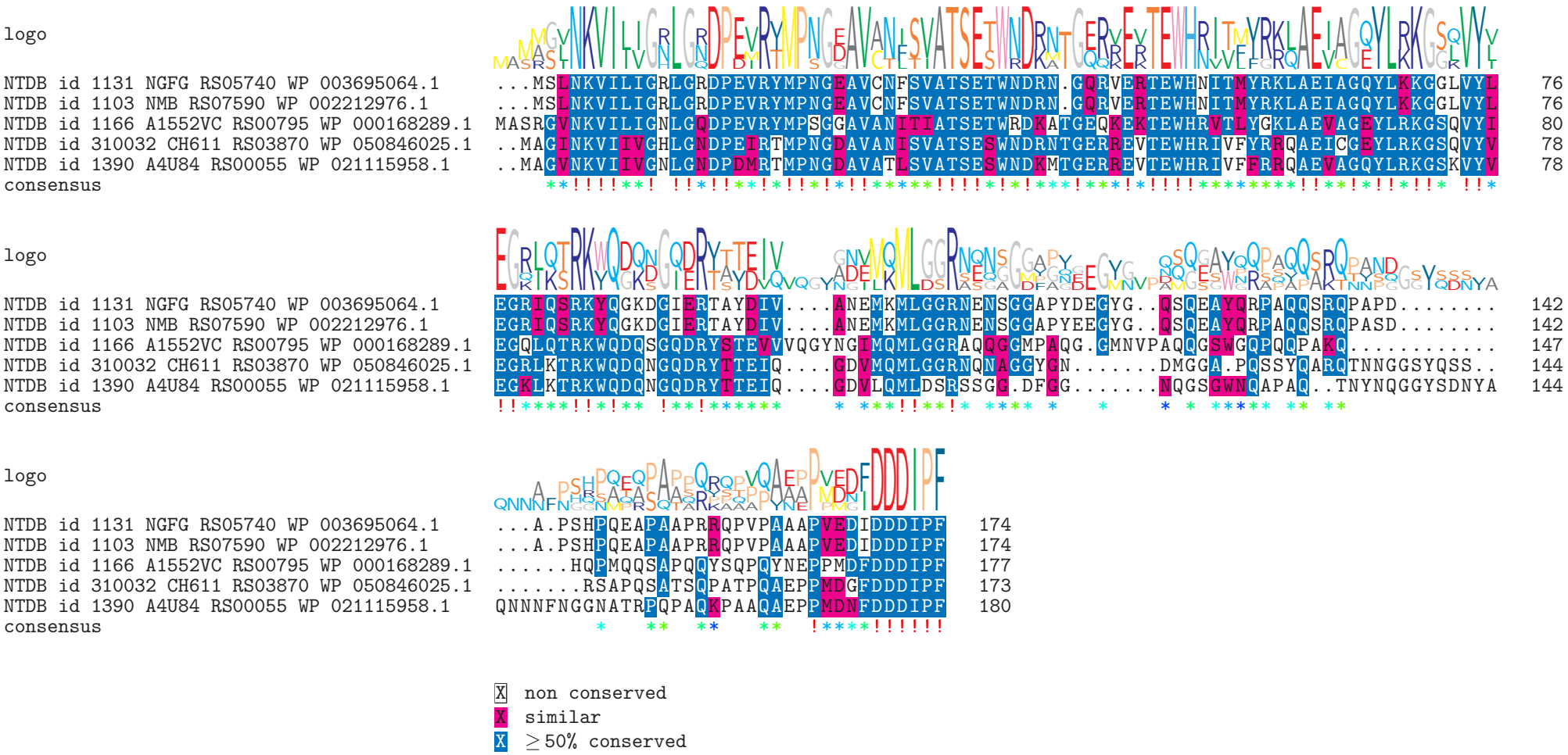

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
NTDB id 310032 CH611 RS03870 WP 050846025.1 ..MAGINKVIIVGHLGNDPEIRTMPNGDAVANISVATSESWNDRNTGERREVTEWHRIVFYRRQAEICGEYLRKGSQVYV 78
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
consensus
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYDEGYG..QSQEAYQRPAQQSRQPAPD........ 142
NTDB id 1103 NMB RS07590 WP 002212976.1 EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYEEGYG..QSQEAYQRPAQQSRQPASD........ 142
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQG.GMNVPAQQGSWGQPQQPAKQ............ 147
NTDB id 310032 CH611 RS03870 WP 050846025.1 EGRLKTRKWQDQNGQDRYTTEIQ....GDVMQMLGGRNQNAGGYGN.......DMGGA.PQSSYQARQTNNGGSYQSS.. 144
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGG.DFGG.......NQGSGWNQAPAQ..TNYNQGGYSDNYA 144
consensus
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...A.PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1103 NMB RS07590 WP 002212976.1 ...A.PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ......HQPMQQSAPQQYSQPQYNEPPMDFDDDIPF 177
NTDB id 310032 CH611 RS03870 WP 050846025.1 .......RSAPQSATSQPATPQAEPPMDGFDDDIPF 173
NTDB id 1390 A4U84 RS00055 WP 021115958.1 QNNNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF 180
consensus
X non conserved
X similar
X ≥ 50% conserved