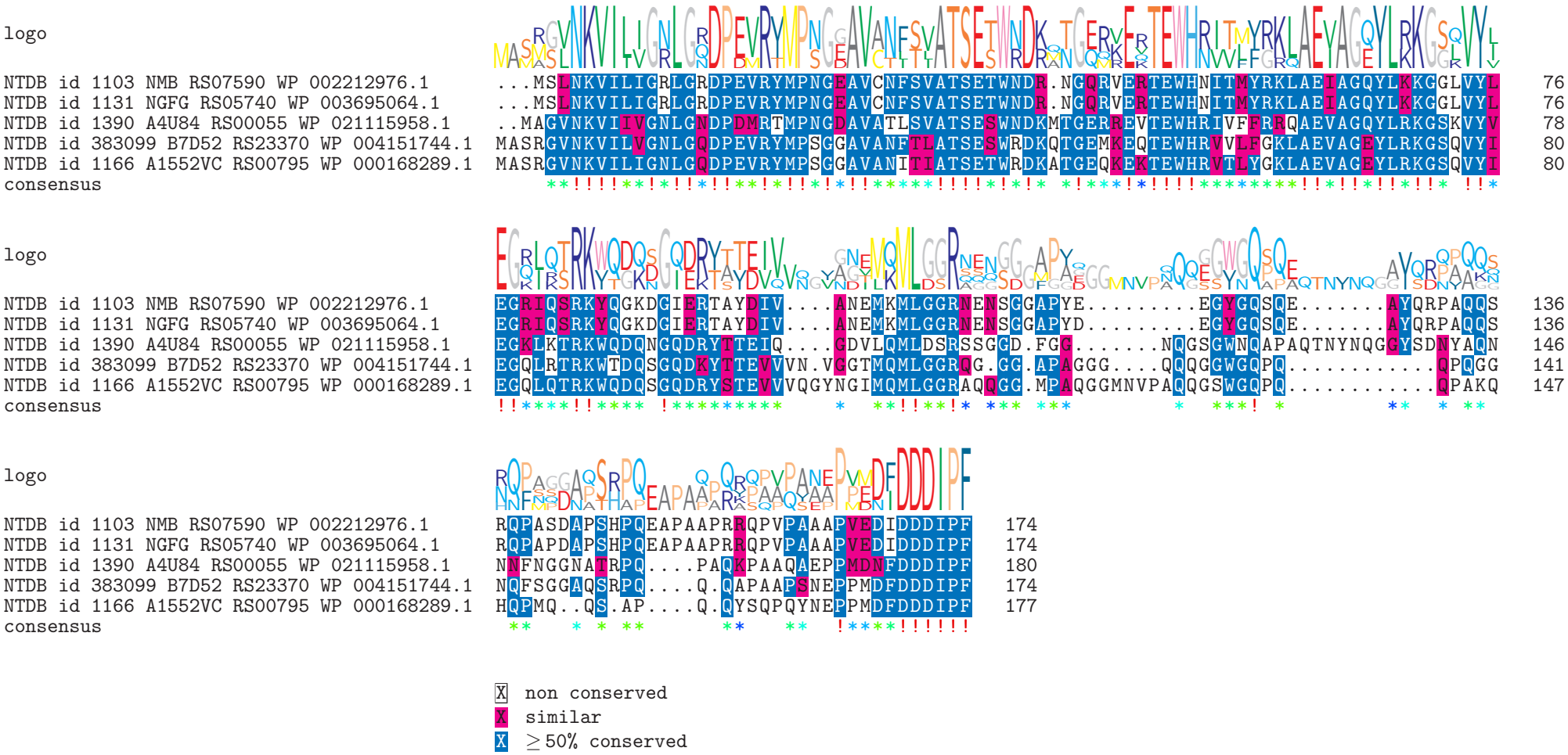

logo
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
NTDB id 383099 B7D52 RS23370 WP 004151744.1 MASRGVNKVILVGNLGQDPEVRYMPSGGAVANFTLATSESWRDKQTGEMKEQTEWHRVVLFGKLAEVAGEYLRKGSQVYI 80
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYE.........EGYGQSQE.......AYQRPAQQS 136
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYD.........EGYGQSQE.......AYQRPAQQS 136
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGGD.FGG.......NQGSGWNQAPAQTNYNQGGYSDNYAQN 146
NTDB id 383099 B7D52 RS23370 WP 004151744.1 EGQLRTRKWTDQSGQDKYTTEVVVN.VGGTMQMLGGRQG.GG.APAGGG....QQQGGWGQPQ.............QPQGG 141
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMNVPAQQGSWGQPQ.............QPAKQ 147
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 RQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1131 NGFG RS05740 WP 003695064.1 RQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1390 A4U84 RS00055 WP 021115958.1 NNFNGGNATRPQ....PAQKPAAQAEPPMDNFDDDIPF 180
NTDB id 383099 B7D52 RS23370 WP 004151744.1 NQFSGGAQSRPQ....Q.QAPAAPSNEPPMDFDDDIPF 174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 HQPMQ..QS.AP....Q.QYSQPQYNEPPMDFDDDIPF 177
consensus
X non conserved
X similar
X ≥ 50% conserved