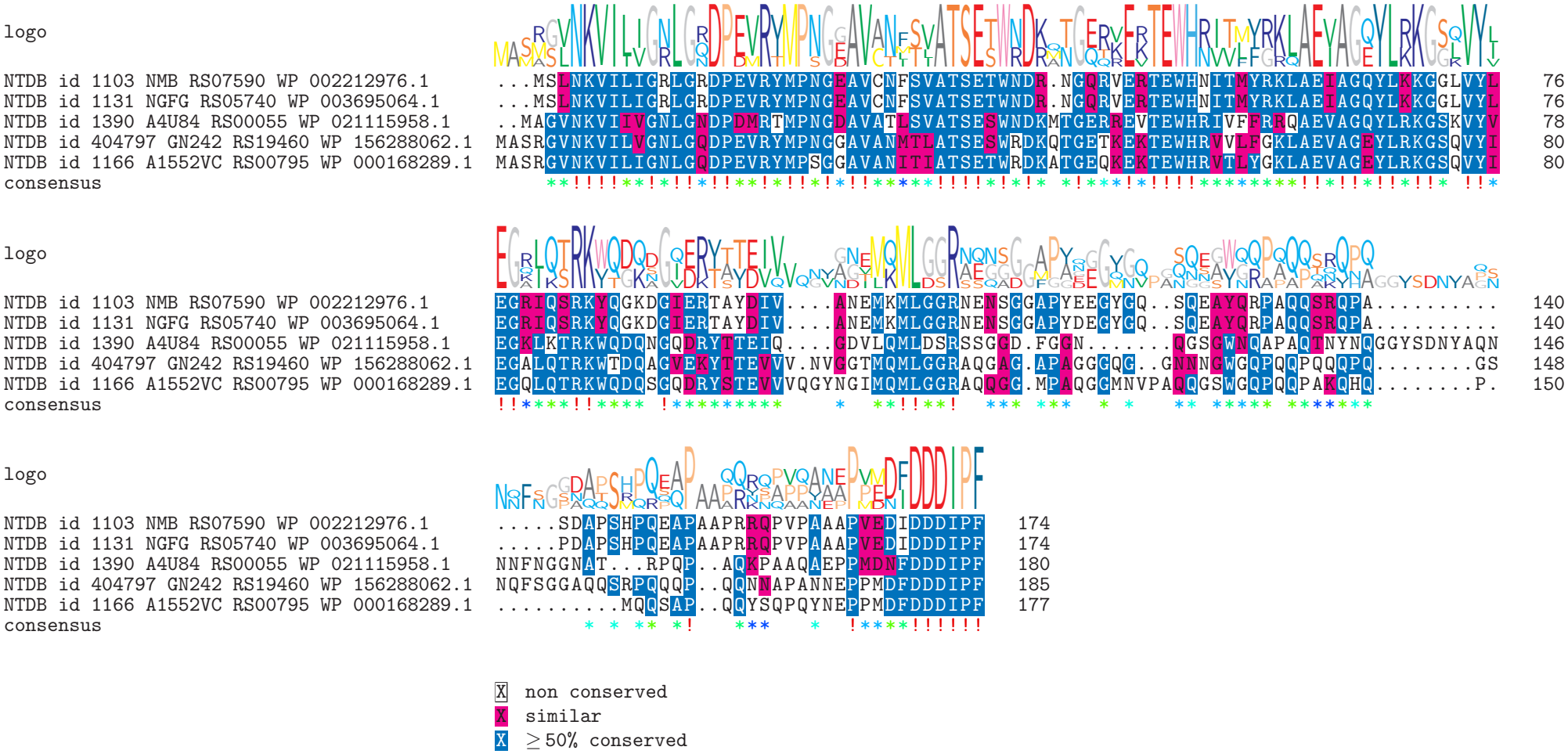
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
NTDB id 404797 GN242 RS19460 WP 156288062.1 MASRGVNKVILVGNLGQDPEVRYMPNGGAVANMTLATSESWRDKQTGETKEKTEWHRVVLFGKLAEVAGEYLRKGSQVYI 80
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYEEGYGQ..SQEAYQRPAQQSRQPA.......... 140
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYDEGYGQ..SQEAYQRPAQQSRQPA.......... 140
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGGD.FGGN.......QGSGWNQAPAQTNYNQGGYSDNYAQN 146
NTDB id 404797 GN242 RS19460 WP 156288062.1 EGALQTRKWTDQAGVEKYTTEVVV.NVGGTMQMLGGRAQGAG.APAGGGQG..GNNNGWGQPQQPQQQPQ........GS 148
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMNVPAQQGSWGQPQQPAKQHQ........P. 150
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 .....SDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1131 NGFG RS05740 WP 003695064.1 .....PDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1390 A4U84 RS00055 WP 021115958.1 NNFNGGNAT...RPQP..AQKPAAQAEPPMDNFDDDIPF 180
NTDB id 404797 GN242 RS19460 WP 156288062.1 NQFSGGAQQSRPQQQP..QQNNAPANNEPPMDFDDDIPF 185
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ....MQQSAP..QQYSQPQYNEPPMDFDDDIPF 177
consensus
non conserved
similar
≥ 50% conserved