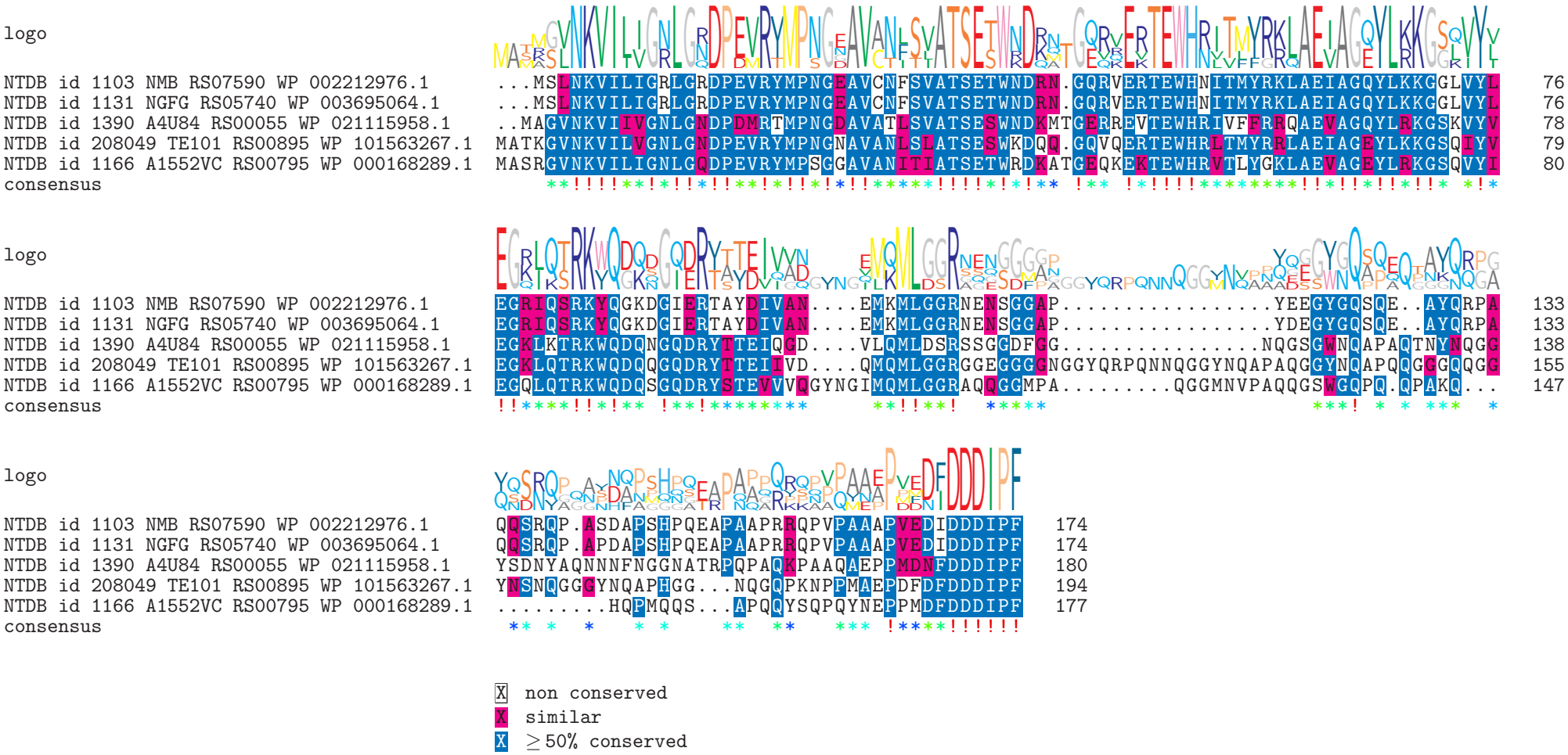
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
NTDB id 208049 TE101 RS00895 WP 101563267.1 MATKGVNKVILVGNLGNDPEVRYMPNGNAVANLSLATSESWKDQQ.GQVQERTEWHRLTMYRRLAEIAGEYLKKGSQIYV 79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAP.....YEEGYGQSQE..AYQRPA 133
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAP.....YDEGYGQSQE..AYQRPA 133
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQGD....VLQMLDSRSSGGDFGG.....NQGSGWNQAPAQTNYNQGG 138
NTDB id 208049 TE101 RS00895 WP 101563267.1 EGKLQTRKWQDQQGQDRYTTEIIVD....QMQMLGGRGGEGGGGNGGYQRPQNNQGGYNQAPAQGGYNQAPQQGGQQGG 155
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPA.....QGGMNVPAQQGSWGQPQ.QPAKQ... 147
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 QQSRQP.ASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1131 NGFG RS05740 WP 003695064.1 QQSRQP.APDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1390 A4U84 RS00055 WP 021115958.1 YSDNYAQNNNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF 180
NTDB id 208049 TE101 RS00895 WP 101563267.1 YNSNQGGGYNQAPHGG...NQGQPKNPPMAEPDFDFDDDIPF 194
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ....HQPMQQS...APQQYSQPQYNEPPMDFDDDIPF 177
consensus
X non conserved
X similar
X ≥ 50% conserved