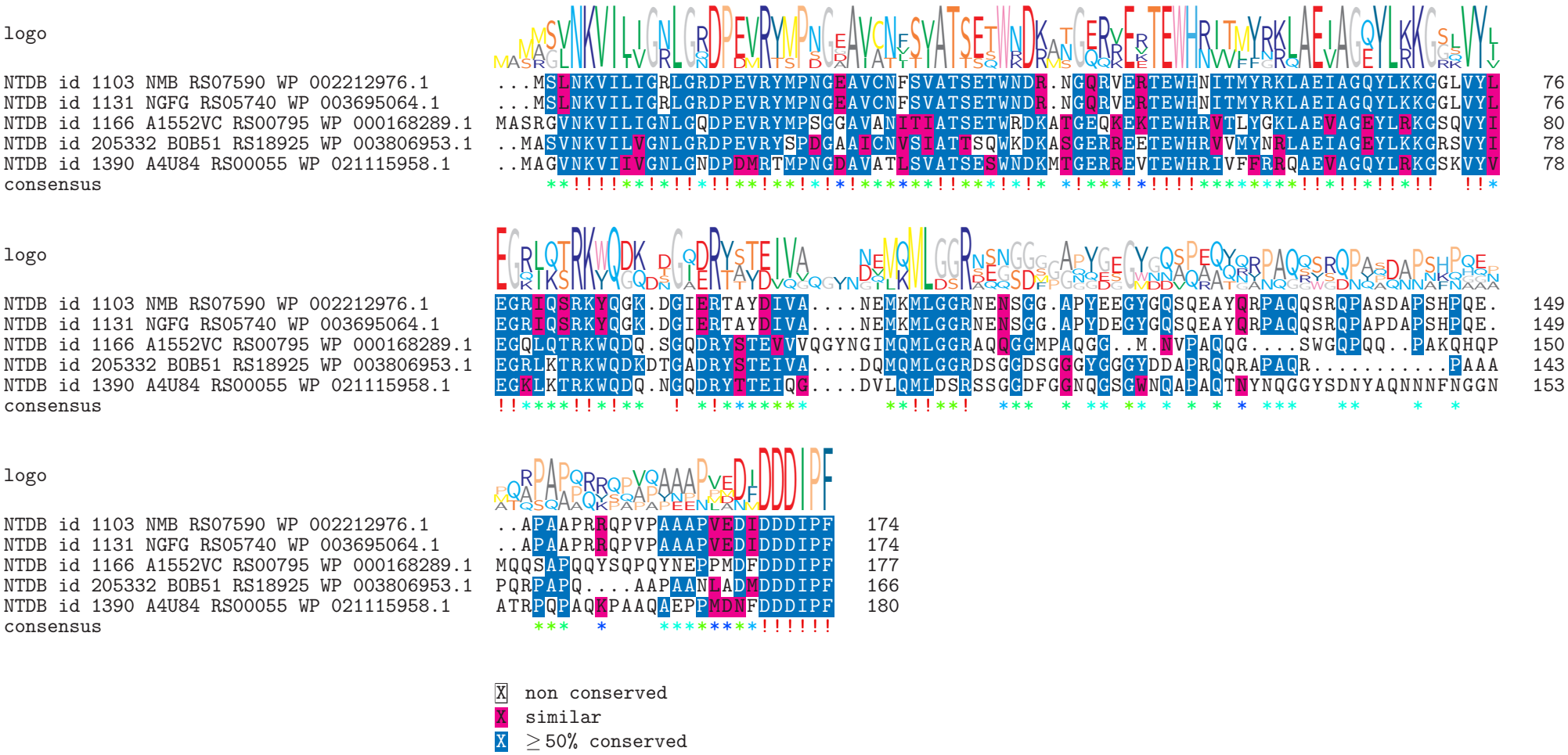

logo
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
NTDB id 205332 BOB51 RS18925 WP 003806953.1 ..MASVNKVILVGNLGRDPEVRYSPDGAAICNVSIATTSQWKDKASGERREETEWHRVVMYNRLAEIAGEYLKKGRSVYI 78
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVITVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 EGRLQSRKYQGK.DGIERTAYDIVA....NEMKMLGGRNENSGG.APYEEGYGQSQEAYQRPAQQSRQPASDAPSHPQE. 149
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGK.DGIERTAYDIVA....NEMKMLGGRNENSGG.APYDEGYGQSQEAYQRPAQQSRQPAPDAPSHPQE. 149
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQ.SGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGG..M.NVPAQQG....SWGQPQQ..PAKQHQP 150
NTDB id 205332 BOB51 RS18925 WP 003806953.1 EGRLKTRKWQDKDTGADRYSTEIVA....DQMQMLGGRDSGGDSGGGYGGGYDDAPRQQRAPAQR...........PAAA 143
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQ.NGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGGN 153
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 ..APAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1131 NGFG RS05740 WP 003695064.1 ..APAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MQQSAPQQYSQPQYNEPPMDFDDDIPF 177
NTDB id 205332 BOB51 RS18925 WP 003806953.1 PQRPAPQ....AAPAANLADMDDDIPF 166
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ATRPQPAQKPAAQAEPPMDNFDDDIPF 180
consensus
X non conserved
X similar
X ≥ 50% conserved