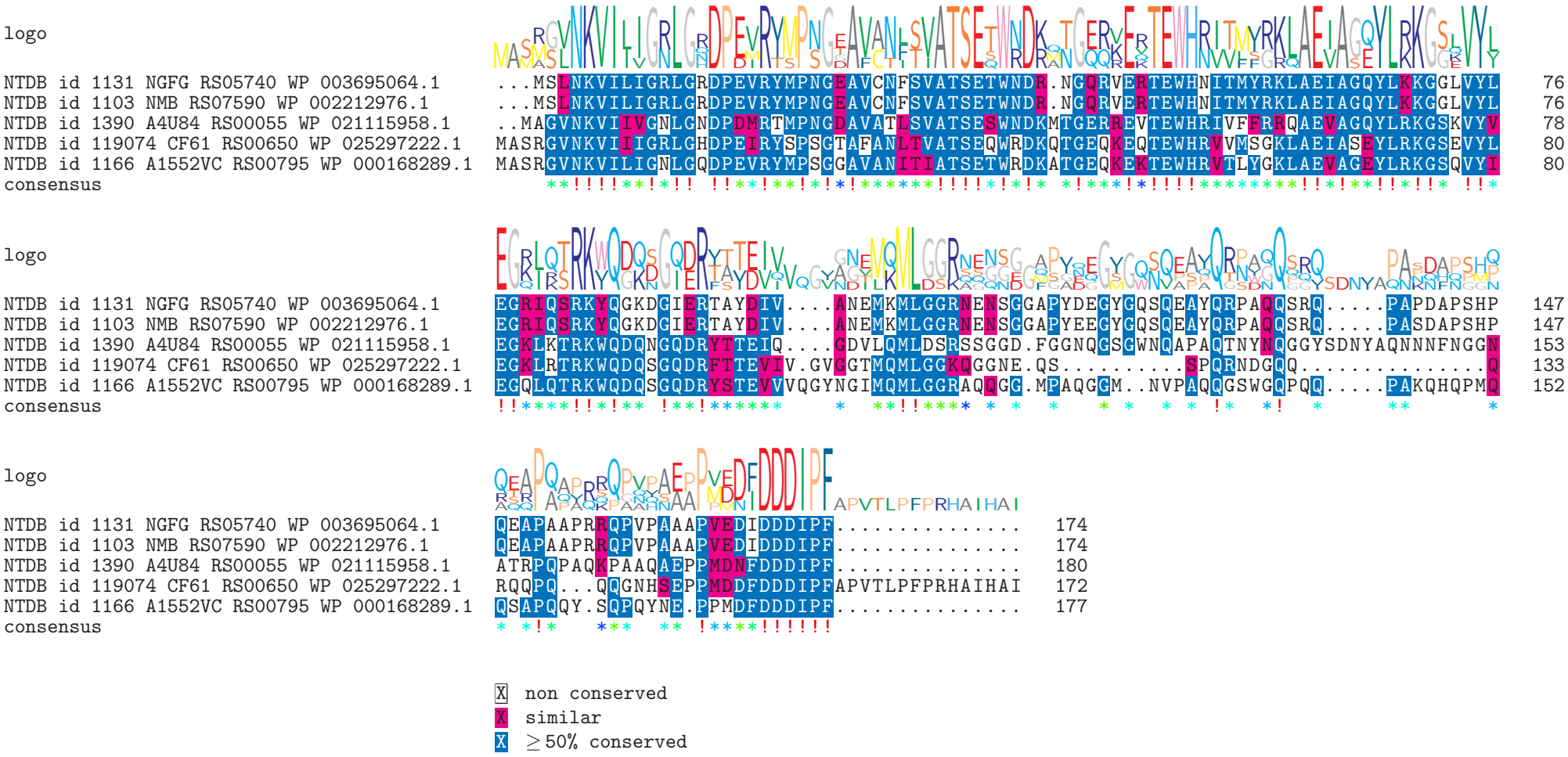
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
NTDB id 119074 CF61 RS00650 WP 025297222.1 MASRGVNKVIIIGRLGHDPEIRYSPSGTAFANLTVATSEQWRDKQTGEQKEQTEWHRVVMSGKLAEIASEYLRKGSEVYL 80
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
consensus
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYDEGYGQSQEAYQRPAQQSRQ.....PAPDAPSHP 147
NTDB id 1103 NMB RS07590 WP 002212976.1 EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYEEGYGQSQEAYQRPAQQSRQ.....PASDAPSHP 147
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGGN 153
NTDB id 119074 CF61 RS00650 WP 025297222.1 EGKLRTRKWQDQSGQDRFTTEVIV.GVGGTMQMLGGKQGGNE.QS..........SPQRNDGQQ..............Q 133
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGGM..NVPAQQGSWGQPQQ.....PAKQHQPMQ 152
consensus
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 QEAPAAPRRQPVPAAAPVEDIDDDIPF.............. 174
NTDB id 1103 NMB RS07590 WP 002212976.1 QEAPAAPRRQPVPAAAPVEDIDDDIPF.............. 174
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ATRPQPAQKPAAQAEPPMDNFDDDIPF.............. 180
NTDB id 119074 CF61 RS00650 WP 025297222.1 RQQPQ...QQGNHSEPPMDDFDDDIPFAPVTLPFPRHAIHAI 172
NTDB id 1166 A1552VC RS00795 WP 000168289.1 QSAPQQY.SQPQYNE.PPMDFDDDIPF.............. 177
consensus
X non conserved
X similar
X ≥ 50% conserved