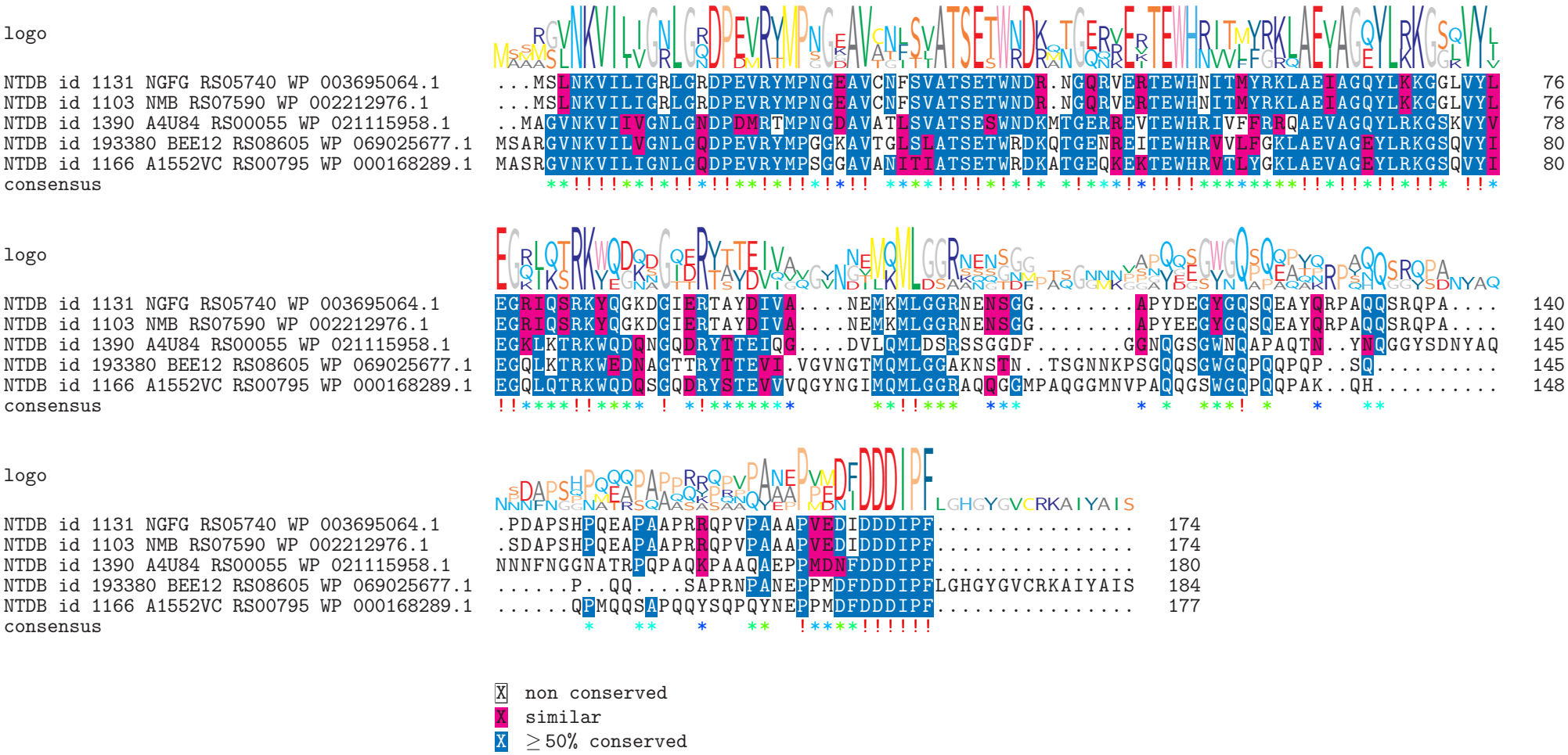
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
NTDB id 193380 BEE12 RS08605 WP 069025677.1 MSARGVNKVILVGNLGQDPEVRYMPGGKAVTGLSLATSETWRDKQTGENREITEWHRVVLFGKLAEVAGEYLRKGSQVYI 80
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
consensus
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGG........APYDEGYGQSQEAYQRPAQQSRQPA.... 140
NTDB id 1103 NMB RS07590 WP 002212976.1 EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGG........APYEEGYGQSQEAYQRPAQQSRQPA.... 140
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDF.......GGNQGSGWNQAPAQTN..YNQGGYSDNYAQ 145
NTDB id 193380 BEE12 RS08605 WP 069025677.1 EGQLKTRKWEDNAGTTRYTTEVI.VGVNGTMQMLGGAKNSTN..TSGNNKPSGQQSGWGQPQQPQP..SQ.......... 145
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVPAQQGSWGQPQQPAK..QH.......... 148
consensus
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 .PDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF.................. 174
NTDB id 1103 NMB RS07590 WP 002212976.1 .SDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF.................. 174
NTDB id 1390 A4U84 RS00055 WP 021115958.1 NNNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF.................. 180
NTDB id 193380 BEE12 RS08605 WP 069025677.1 ......P..QQ....SAPRNPANEPPMDFDDDIPFLGHGYGVCRKAIYAIS 184
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ......QPMQQSAPQQYSQPQYNEPPMDFDDDIPF.................. 177
consensus
X non conserved
X similar
X ≥ 50% conserved