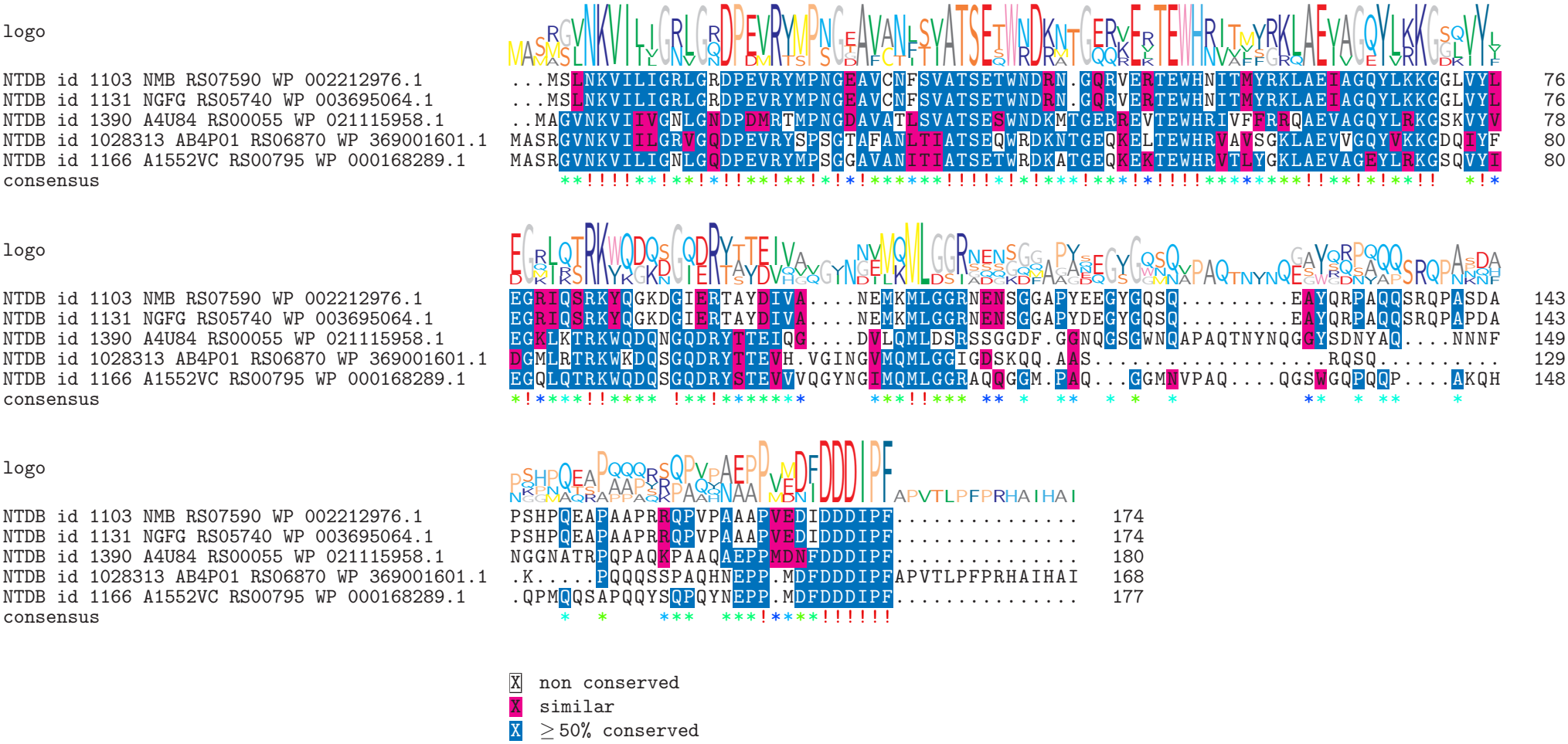

logo
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
NTDB id 1028313 AB4P01 RS06870 WP 369001601.1 MASRGVNKVILLGRVGQDPEVRYSPSGTAFANLTIATSEQWRDKNTGEQKELTEWHRVAVSGKLAEVVGQYVKKGDQIYF 80
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQ.........EAYQRPAQQSRQPASDA 143
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQ.........EAYQRPAQQSRQPAPDA 143
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDF.GGNQGSGWNQAPAQTNYNQGGYSDNYAQ....NNNF 149
NTDB id 1028313 AB4P01 RS06870 WP 369001601.1 DGMLRTRKWKDQSGQDRYTTEVH.VGINGVMQMLGGIGDSKQQ.AAS.....................RQSQ......... 129
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGM.PAQ...GGMNVPAQ....QGSWGQPQQP....AKQH 148
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF..... 174
NTDB id 1131 NGFG RS05740 WP 003695064.1 PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF..... 174
NTDB id 1390 A4U84 RS00055 WP 021115958.1 NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF..... 180
NTDB id 1028313 AB4P01 RS06870 WP 369001601.1 .K.....PQQQSSPAQHNEPP.MDFDDDIPFAPVTLPFPRHAIHAI 168
NTDB id 1166 A1552VC RS00795 WP 000168289.1 .QPMQQSAPQQYSQPQYNEPP.MDFDDDIPF..... 177
consensus
X non conserved
X similar
X ≥ 50% conserved