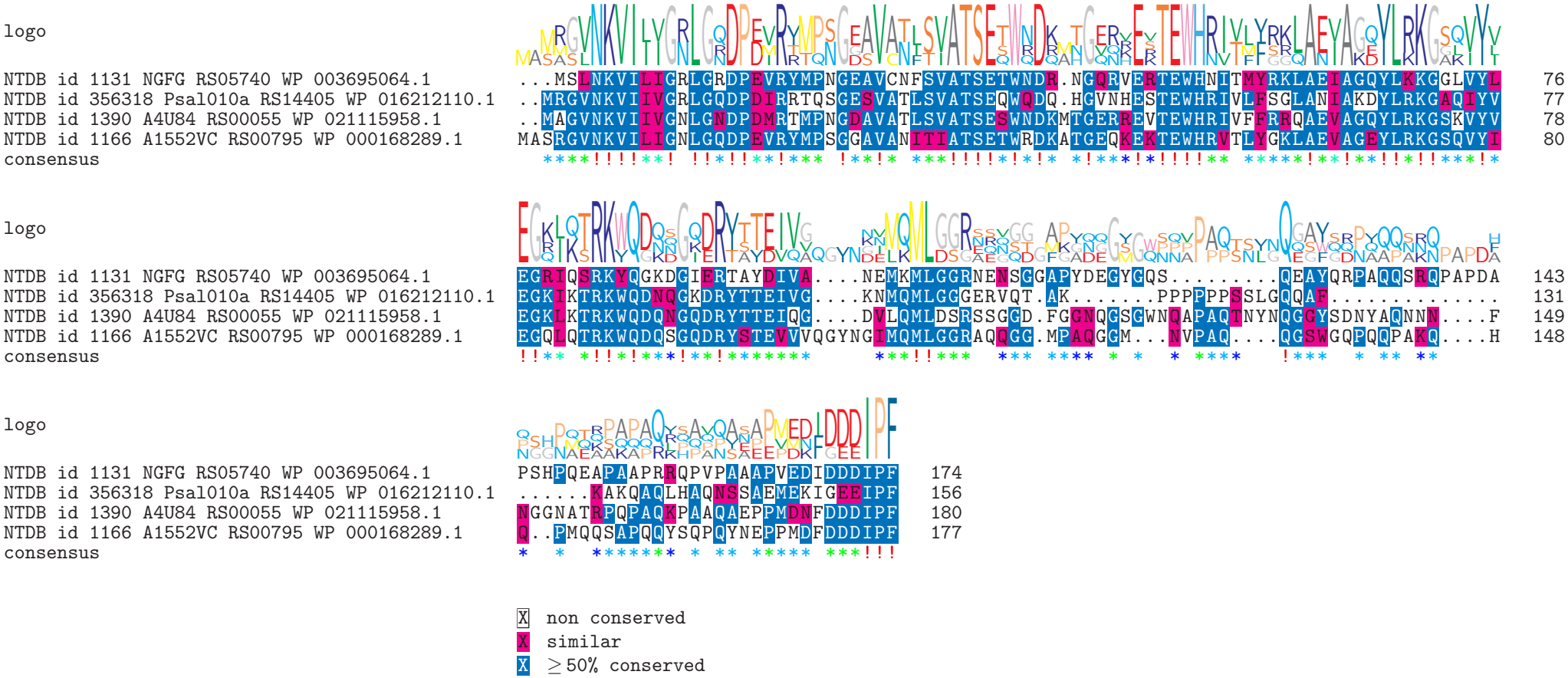

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 356318 Psal010a RS14405 WP 016212110.1 ..MRGVNKVIIVGRLGQDPDIRRTQSGESVATLSVATSEQWQDQ.HGVNHESTEWHRIVLFSGLANIAKDYLRKGAQIYV 77
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
consensus
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQS.........QEAYQRPAQQSRQPAPDA 143
NTDB id 356318 Psal010a RS14405 WP 016212110.1 EGKIKTRKWQDNQGKDRYTTEIVG....KNMQMLGGGERVQT.AK.......PPPPPPSSLGQQAF.............. 131
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGCD.FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNN....F 149
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGM...NVPAQ....QGSWGQPQQPAKQ....H 148
consensus
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1 PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 356318 Psal010a RS14405 WP 016212110.1 ......KAKQAQLHAQNSSAEMEKIGEEIPF 156
NTDB id 1390 A4U84 RS00055 WP 021115958.1 NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF 180
NTDB id 1166 A1552VC RS00795 WP 000168289.1 Q..PMQQSAPQQYSQPQYNEPPMDFDDDIPF 177
consensus
X non conserved
X similar
X ≥ 50% conserved