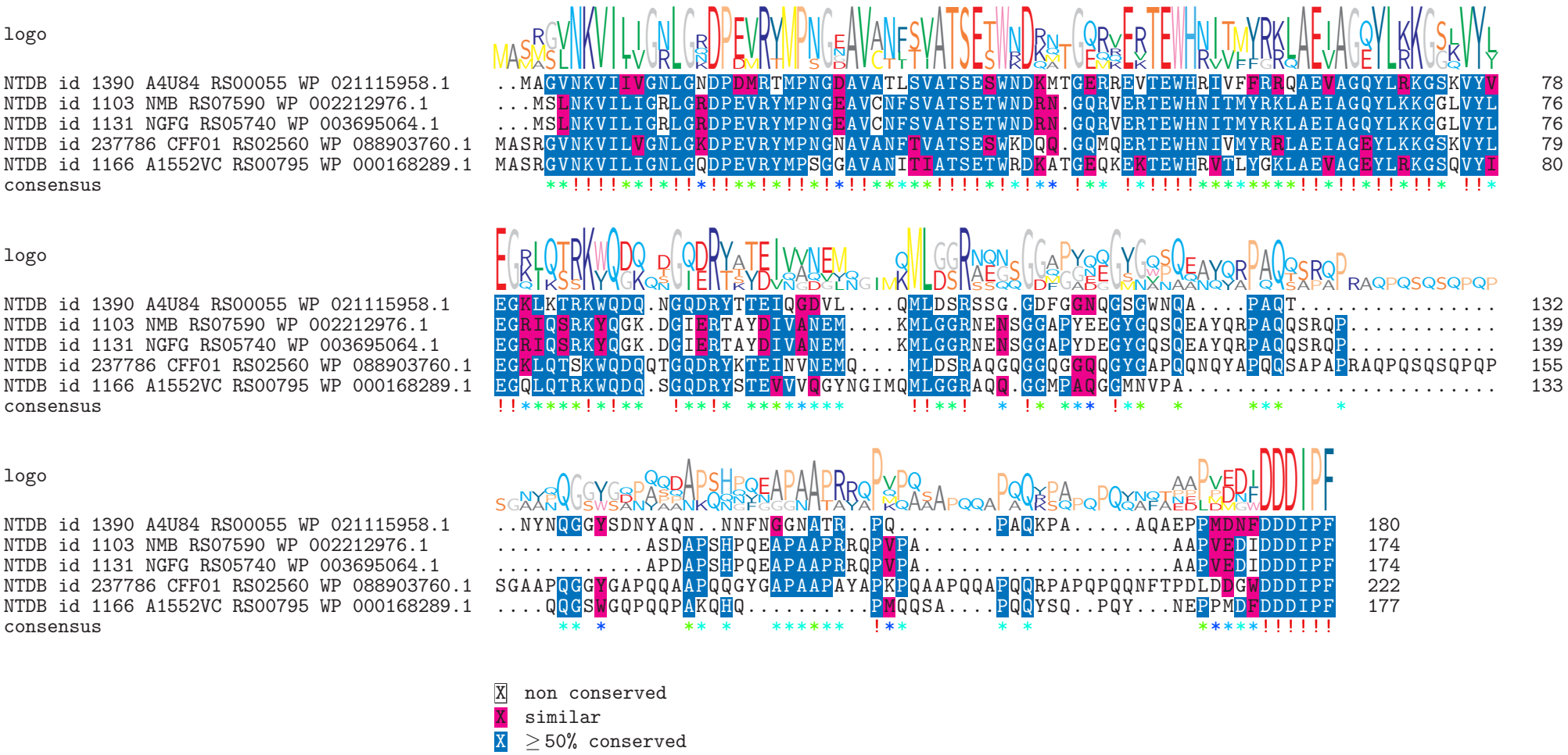
logo
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 237786 CFF01 RS02560 WP 088903760.1 MASRGVNKVILVGNLGKDPEVRYMPNGNAVANFTVATSESWKDQQ.GQMQERTEWHNIVMYRRLAEIAGEYLKKGSKVYL 79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
consensus
logo
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQ.NGQDRYTTEIQGDVL....QMLDSRSSG.GDFGGNQGSGWNQA....PAQT..... 132
NTDB id 1103 NMB RS07590 WP 002212976.1 EGRIQSRKYQGK.DGIERTAYDIVANEM....KMLGGRNENSGGAPYEEGYGQSQEAYQRPAQQSRQP..... 139
NTDB id 1131 NGFG RS05740 WP 003695064.1 EGRIQSRKYQGK.DGIERTAYDIVANEM....KMLGGRNENSGGAPYDEGYGQSQEAYQRPAQQSRQP..... 139
NTDB id 237786 CFF01 RS02560 WP 088903760.1 EGKLQTSKWQDQQTGQDRYKTEINVNEMQ....MLDSRAQGQGGQGGQQGYGAPQQNQYAPQQSAPAPRAQPQSQSQPQP 155
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQ.SGQDRYSTEVVVQGYNGIMQMLGGRAQQ.GGMPAQQGGMNVPA..... 133
consensus
logo
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..NYNQGGYSDNYAQN..NNFNGGNATR..PQ.....PAQKPA.....AQAEPPMDNFDDDIPF 180
NTDB id 1103 NMB RS07590 WP 002212976.1 .....ASDAPSHPQEAPAAPRRQPVPA.....AAPVEDIDDDIPF 174
NTDB id 1131 NGFG RS05740 WP 003695064.1 .....APDAPSHPQEAPAAPRRQPVPA.....AAPVEDIDDDIPF 174
NTDB id 237786 CFF01 RS02560 WP 088903760.1 SGAAPQGGYGAPQQAAPQQGYGAPAAPAYAPKPQAAPQQAPQQRPAPQPQQNFTPDLDDGWDDDIPF 222
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ....QQGSWGQPQQPAKQHQ.....PMQQSA....PQQYSQ..PQY...NEPPMDFDDDIPF 177
consensus
X non conserved
X similar
X ≥ 50% conserved