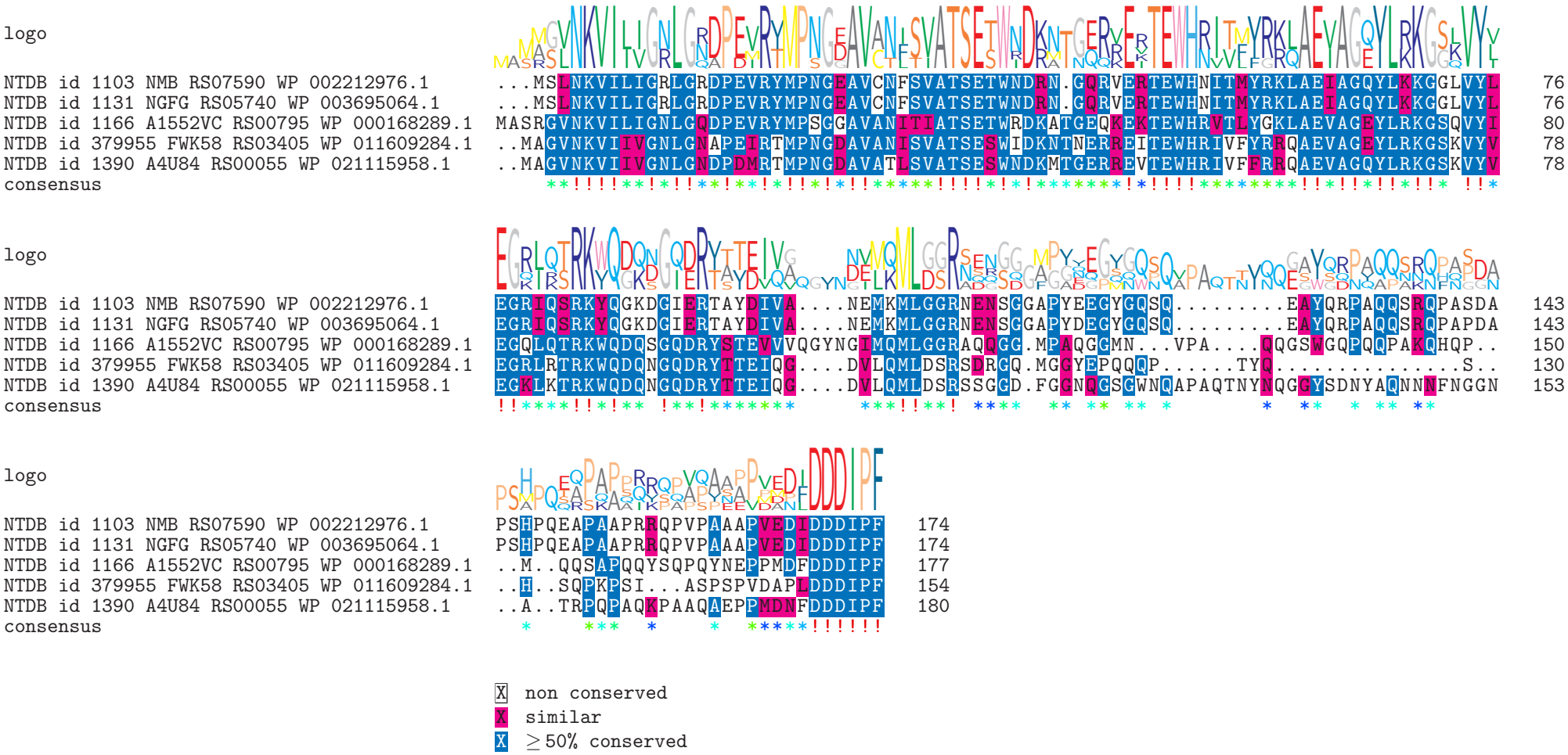

logo
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL 76
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI 80
NTDB id 379955 FWK58 RS03405 WP 011609284.1 ..MAGVNKVIIVGNLGNAPEIRTMPNGDAVANISVATSESWIDKNTNERREITEWHRIVFYRRQAEVAGEYLRKGSKVYV 78
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV 78
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 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMN...VPA....QQGSWGQPQQPAKQHQP.. 150
NTDB id 379955 FWK58 RS03405 WP 011609284.1 EGRLRTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSDRGQ.MGGYEPQQQP......TYQ..............S.. 130
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGGN 153
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1131 NGFG RS05740 WP 003695064.1 PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ..M..QQSAPQQYSQPQYNEPPMDFDDDIPF 177
NTDB id 379955 FWK58 RS03405 WP 011609284.1 ..H..SQPKPSI...ASPSPVDAPLDDDIPF 154
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..A..TRPQPAQKPAAQAEPPMDNFDDDIPF 180
consensus
X non conserved
X similar
X ≥ 50% conserved