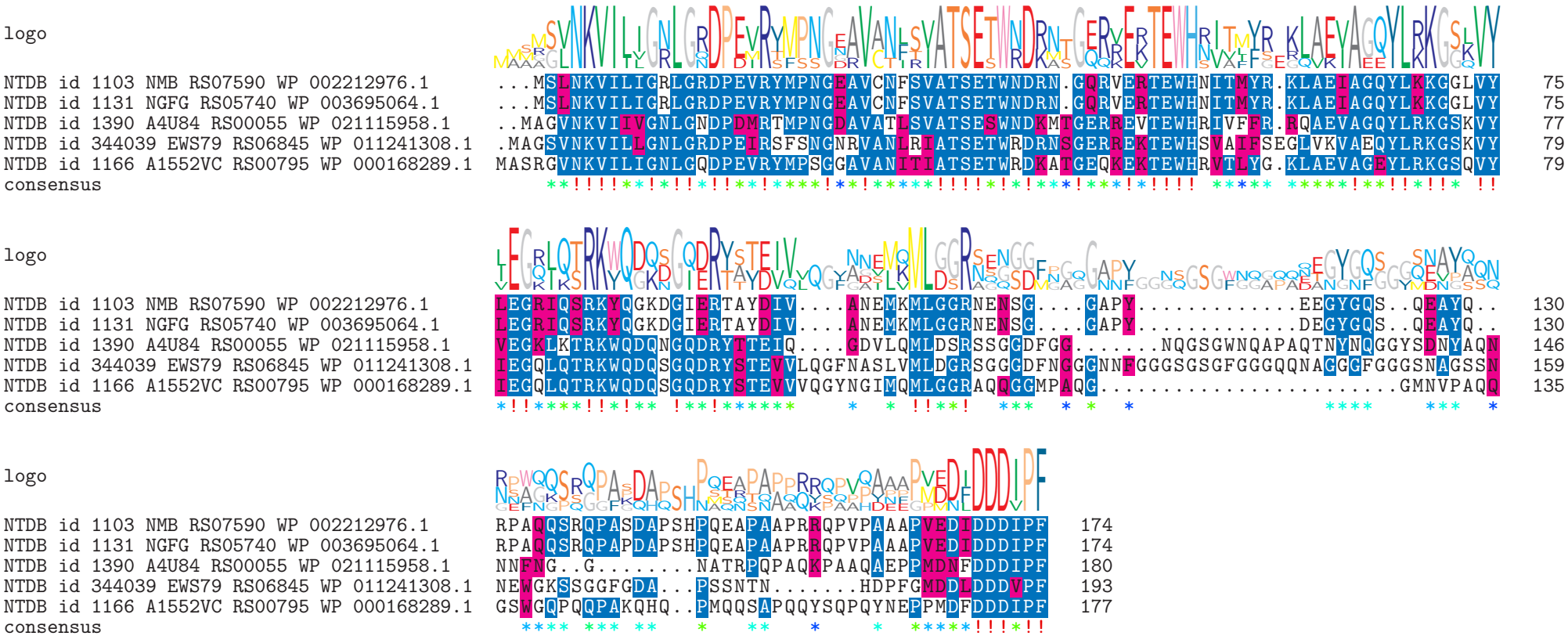

logo
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY 75
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY 75
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY 77
NTDB id 344039 EWS79 RS06845 WP 011241308.1 .MAGSVNKVILLGNLGRDPEIRSFSNGNRVANLRIATSETWRDRNSGERREKTEWHSVAIFSEGLVKVAEQYLRKGSKVY 79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY 79
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 LEGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSG....GAPY.............EEGYGQS..QEAYQ.. 130
NTDB id 1131 NGFG RS05740 WP 003695064.1 LEGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSG....GAPY.............DEGYGQS..QEAYQ.. 130
NTDB id 1390 A4U84 RS00055 WP 021115958.1 VEGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGGDFGG.......NQGSGWNQAPAQTNYNQGGYSDNYAQN 146
NTDB id 344039 EWS79 RS06845 WP 011241308.1 IEGQLQTRKWQDQSGQDRYSTEVVLQGFNASLVMLDGRSGGGDFNGGGNNFGGGSGSGFGGGQQNAGGGFGGGSNAGSSN 159
NTDB id 1166 A1552VC RS00795 WP 000168289.1 IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQG.....................GMNVPAQQ 135
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 RPAQQSRQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1131 NGFG RS05740 WP 003695064.1 RPAQQSRQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF 174
NTDB id 1390 A4U84 RS00055 WP 021115958.1 NNFNG..G.....NATRPQPAQKPAAQAEPPMDNFDDDIPF 180
NTDB id 344039 EWS79 RS06845 WP 011241308.1 NEWGKSSGGFGDA...PSSNTN.....HDPFGMDDLDDDVPF 193
NTDB id 1166 A1552VC RS00795 WP 000168289.1 GSWGQPQQPAKQHQ..PMQQSAPQQYSQPQYNEPPMDFDDDIPF 177
consensus
X non conserved
X similar
X ≥ 50% conserved