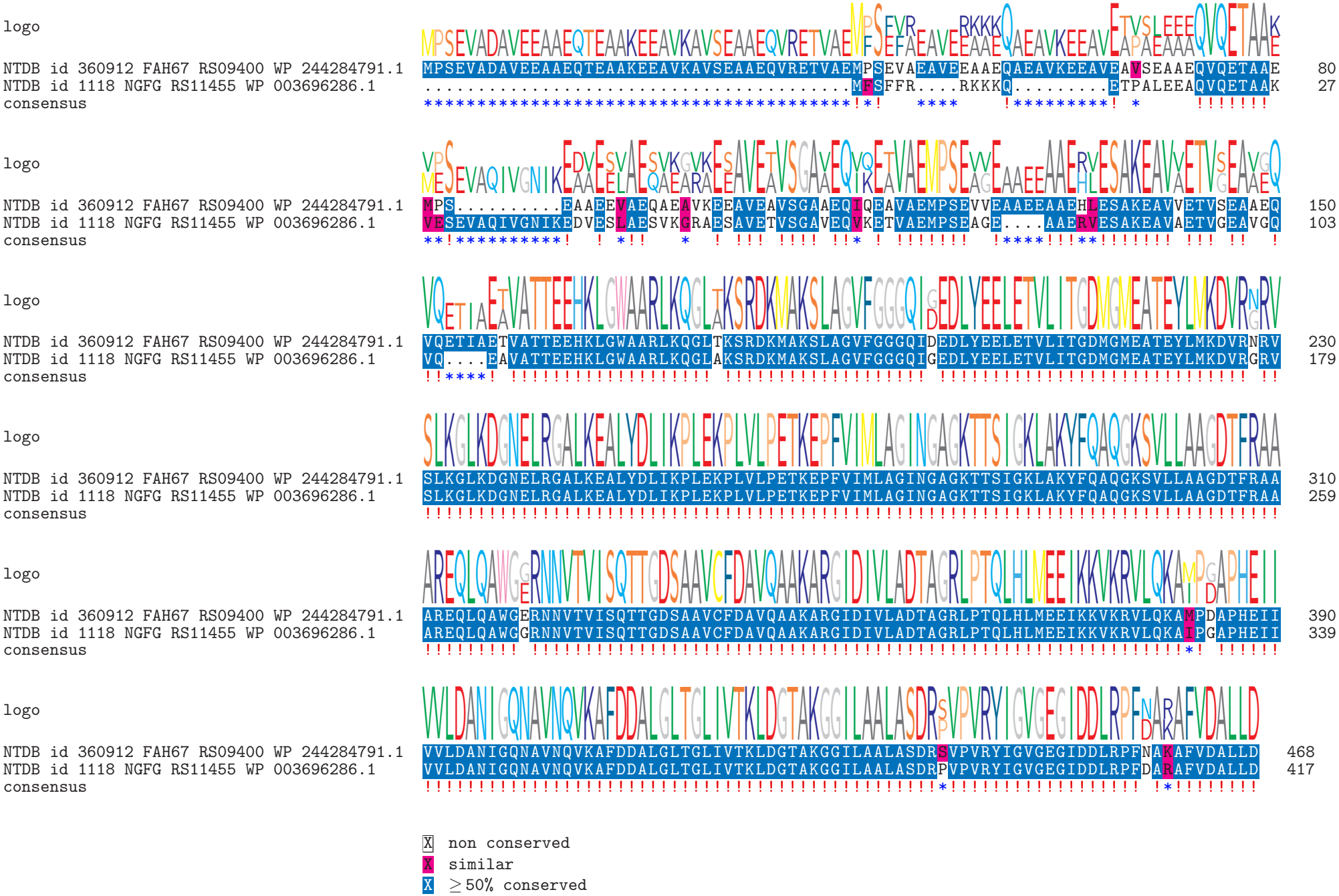
logo
NTDB id 360912 FAH67 RS09400 WP 244284791.1 MPSEVADAVEEAAEQTEAAKEEAVKAVSEAAEQVRETVAEMPSEVAEAVEEAAEQAEAVKEEAVEAVSEAAEQVQETAAE 80
NTDB id 1118 NGFG RS11455 WP 003696286.1 .....MFSFFR....RKKKQ.....ETPALEEAQVQETAAK 27
consensus
NTDB id 360912 FAH67 RS09400 WP 244284791.1 MPS.....EAAEEVAEQAEAVKEEAVEAVSGAAEQIQEAVAEMPSEVVEAAEEAAEHLESAKEAVVETVSEAAEQ 150
NTDB id 1118 NGFG RS11455 WP 003696286.1 VESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAGE....AAERVESAKEAVAETVGEAVGQ 103
consensus
NTDB id 360912 FAH67 RS09400 WP 244284791.1 VQETIAETVATTEEHKLGWAARLKQGLTKSRDKMAKSLAGVFGGGQIGEDLYEELETVLITGDMGMEATEYLMKDVRNRV 230
NTDB id 1118 NGFG RS11455 WP 003696286.1 VQ....EAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLITGDMGMEATEYLMKDVRGRV 179
consensus
NTDB id 360912 FAH67 RS09400 WP 244284791.1 SLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKYFQAQGKSVLLAAGDTFRAA 310
NTDB id 1118 NGFG RS11455 WP 003696286.1 SLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKYFQAQGKSVLLAAGDTFRAA 259
consensus
NTDB id 360912 FAH67 RS09400 WP 244284791.1 AREQLQAWGERNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEIKKVKRVLQKAMPDAPHEII 390
NTDB id 1118 NGFG RS11455 WP 003696286.1 AREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEIKKVKRVLQKAIPGAPHEII 339
consensus
NTDB id 360912 FAH67 RS09400 WP 244284791.1 VVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRSVPVRYIGVGEGIDDLRPFNAKAFVDALLD 468
NTDB id 1118 NGFG RS11455 WP 003696286.1 VVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGIDDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved