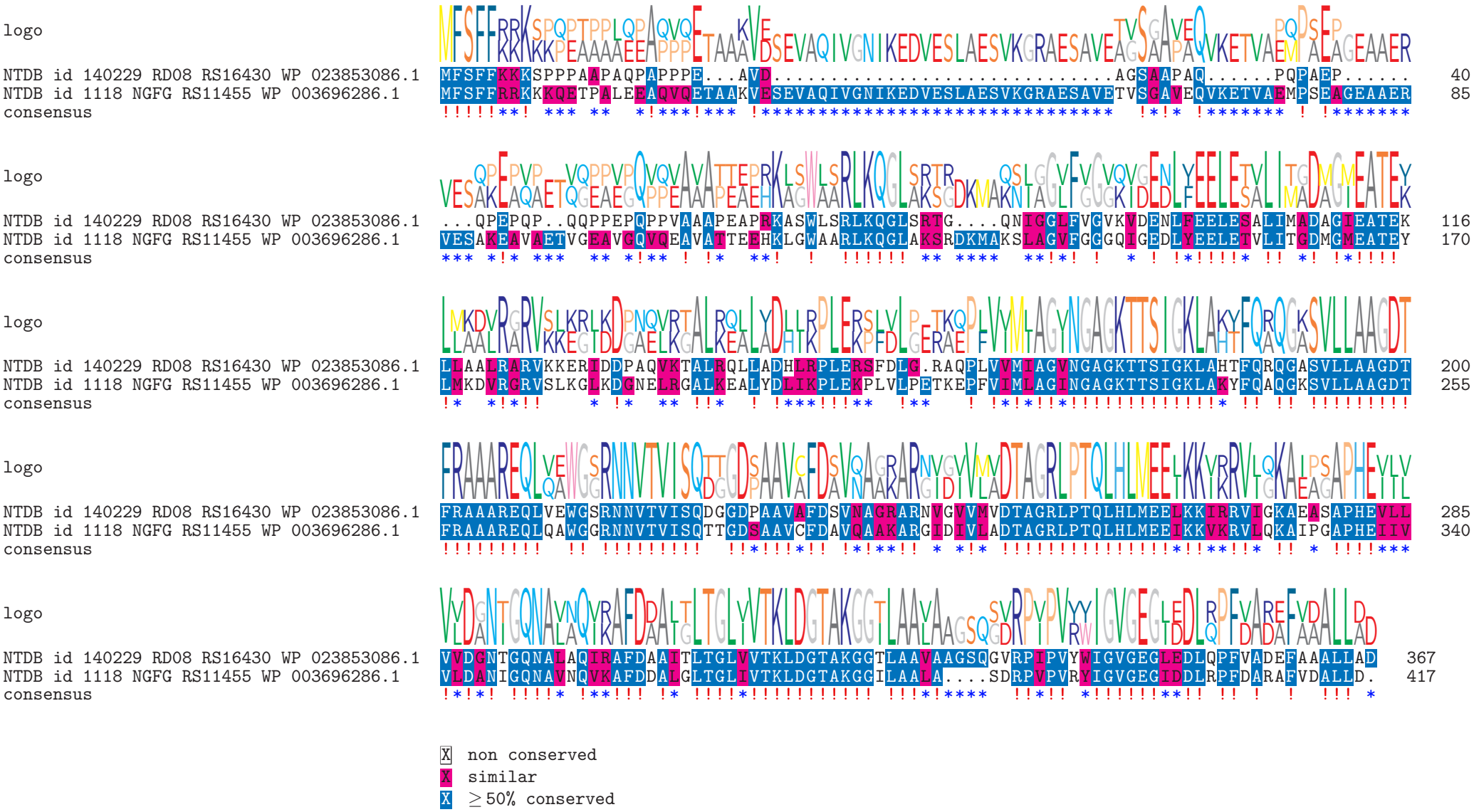

logo
NTDB id 140229 RD08 RS16430 WP 023853086.1 MFSFFKKKSPPPAAPAQPAPPPE...AVD.....AGSAAPAQ......PQPAEP..... 40
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAGEAAER 85
consensus
NTDB id 140229 RD08 RS16430 WP 023853086.1 ...QPEPQP..QQPPEPQPPVAAAPEAPRKASWLSRLKQGLSRTG....QNIGGLFVGVKVDENLFEELESALIMADAGIEATEK 116
NTDB id 1118 NGFG RS11455 WP 003696286.1 VESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLITGDMGMEATEY 170
consensus
NTDB id 140229 RD08 RS16430 WP 023853086.1 LLAALRARVKKERIDDPAQVKTALRQLLADHLRPLERSFDLG.RAQPLVVMIAGVNGAGKTTSIGKLAHTFQRQGASVLLAAGDT 200
NTDB id 1118 NGFG RS11455 WP 003696286.1 LMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKYFQAQGKSVLLAAGDT 255
consensus
NTDB id 140229 RD08 RS16430 WP 023853086.1 FRAAAREQLVEWGSRNNVTVISQDGGDPAAVAFDSVNAGRARNVGVVMVDTAGRLPTQLHLMEELKKIRRVIGKAEASAPHEVLL 285
NTDB id 1118 NGFG RS11455 WP 003696286.1 FRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEIKKVKRVLQKAIPGAPHEIIV 340
consensus
NTDB id 140229 RD08 RS16430 WP 023853086.1 VVDGNTGQNALAQIRAFDAAITLTGLVVTKLDGTAKGGTLAAVAAGSQGVRPIPVYWIGVGEGLEDLQPFVADEFAAALLAD 367
NTDB id 1118 NGFG RS11455 WP 003696286.1 VLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALA....SDRPVPVRYIGVGEGIDDLRPFDARAFVDALLD. 417
consensus
X non conserved
X similar
X ≥ 50% conserved