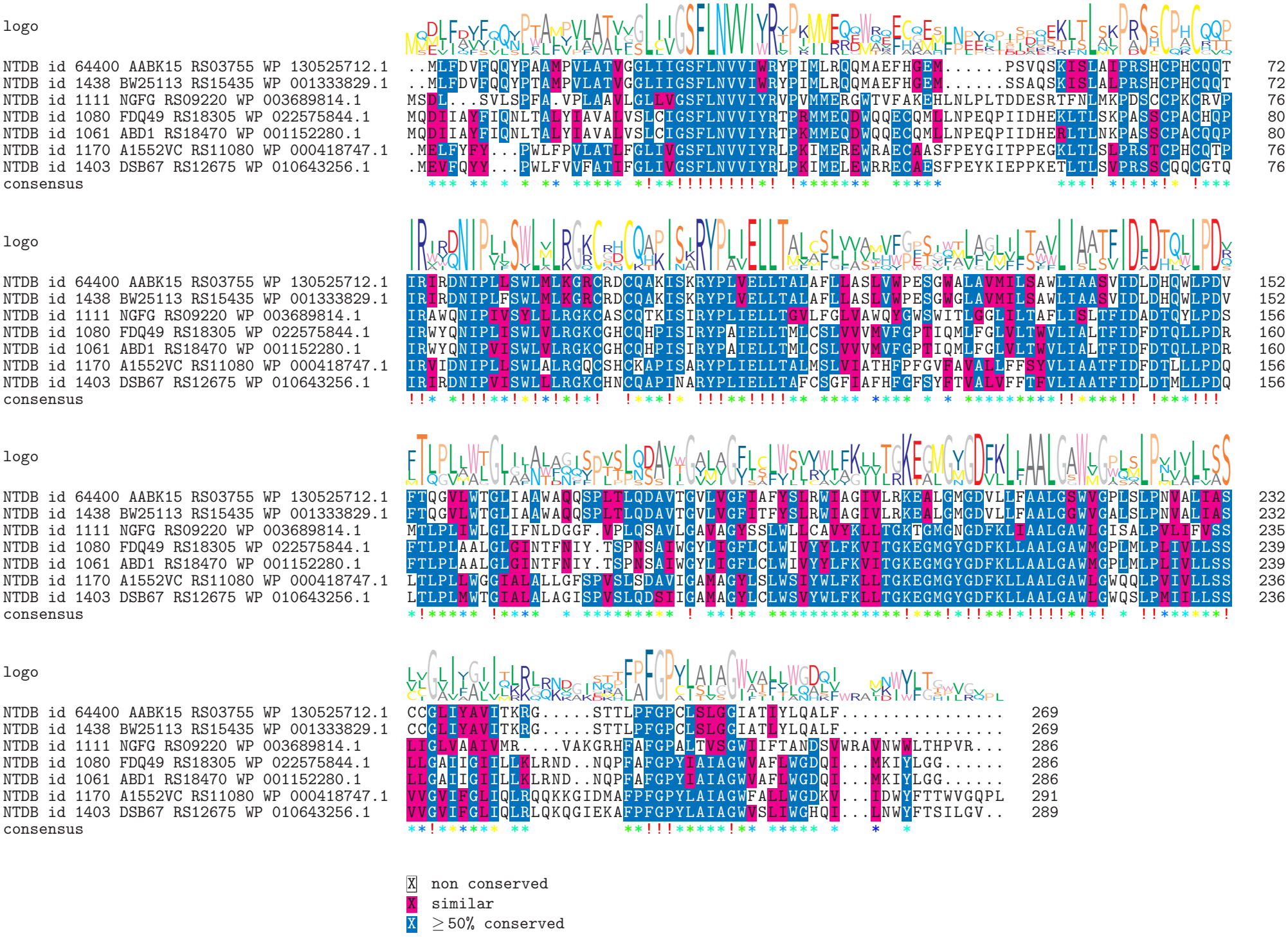
logo
NTDB id 64400 AABK15 RS03755 WP 130525712.1 ..MLFDVFQQYPAAMPVLATVGGLIIGSFLNVVIWRYPIMLRQQMAEFHGEM......PSVQSKISLAIPRSHCPHCQQT 72
NTDB id 1438 BW25113 RS15435 WP 001333829.1 ..MLFDVFQQYPTAMPVLATVGGLIIGSFLNVVIWRYPIMLRQQMAEFHGEM......SSAQSKISLALPRSHCPHCQQT 72
NTDB id 1111 NGFG RS09220 WP 003689814.1 MSDL...SVLSPFA.VPLAAVLGLLVGSFLNVVIYRVPVMMERGWTVFAKEHLNLPLTDDESRTFNLMKPDSCCPKCRVP 76
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLNPEQPIIDHEKLTLSKPASSCPACHQP 80
NTDB id 1061 ABD1 RS18470 WP 001152280.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLNPEQPIIDHERLTLNKPASSCPACQQP 80
NTDB id 1170 A1552VC RS11080 WP 000418747.1 .MELFYFY...PWLFPVLATLFGLIVGSFLNVVIYRLPKIMEREWRAECAASFPEYGITPPEGKLTLSLPRSTCPHCQTP 76
NTDB id 1403 DSB67 RS12675 WP 010643256.1 .MEVFQYY...PWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFPEYKIEPPKETLTLSVPRSSCQQCGTQ 76
consensus
logo
NTDB id 64400 AABK15 RS03755 WP 130525712.1 IRIRDNIPLLSWLMLKGRCRDCQAKISKRYPLVELLTALAFLLASLVWPESGWALAVMILSAWLIAASVIDLDHQWLPDV 152
NTDB id 1438 BW25113 RS15435 WP 001333829.1 IRIRDNIPLFSWLMLKGRCRDCQAKISKRYPLVELLTALAFLLASLVWPESGWGLAVMILSAWLIAASVIDLDHQWLPDV 152
NTDB id 1111 NGFG RS09220 WP 003689814.1 IRAWQNIPIVSYLLLRGKCASCQTKISIRYPLIELLTGVLFGLVAWQYGWSWITLGGLILTAFLISLTFIDADTQYLPDS 156
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 IRWYQNIPLISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPDR 160
NTDB id 1061 ABD1 RS18470 WP 001152280.1 IRWYQNIPVISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPDR 160
NTDB id 1170 A1552VC RS11080 WP 000418747.1 IRVIDNIPLLSWLALRGQCSHCKAPISARYPLIELLTALMSLVIATHFPFGVFAVALLFFSYVLIAATFIDFDTLLLPDQ 156
NTDB id 1403 DSB67 RS12675 WP 010643256.1 IRIRDNIPVISWLLLRGKCHNCQAPINARYPLIELLTAFCSGFIAFHFGFSYFTVALVFFTFVLIAATFIDLDTMLLPDQ 156
consensus
logo
NTDB id 64400 AABK15 RS03755 WP 130525712.1 FTQGVLWTGLIAAWAQQSPLTLQDAVTGVLVGFIAFYSLRWIAGIVLRKEALGMGDVLLFAALGSWVGPLSLPNVALIAS 232
NTDB id 1438 BW25113 RS15435 WP 001333829.1 FTQGVLWTGLIAAWAQQSPLTLQDAVTGVLVGFITFYSLRWIAGIVLRKEALGMGDVLLFAALGGWVGALSLPNVALIAS 232
NTDB id 1111 NGFG RS09220 WP 003689814.1 MTLPLIWLGLIFNLDGGF.VPLQSAVLGAVAGYSSLWLLCAVYKLLTGKTGMGNGDFKLIAALGAWLGISALPVLIFVSS 235
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 FTLPLAALGLGINTFNIY.TSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLSS 239
NTDB id 1061 ABD1 RS18470 WP 001152280.1 FTLPLAALGLGINTFNIY.TSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLSS 239
NTDB id 1170 A1552VC RS11080 WP 000418747.1 LTLPLLWGGIALALLGFSPVSLSDAVIGAMAGYLSLWSIYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQQLPVIVLLSS 236
NTDB id 1403 DSB67 RS12675 WP 010643256.1 LTLPLMWTGIALALAGISPVSLQDSIIGAMAGYLCLWSVYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQSLPMIILLSS 236
consensus
logo
NTDB id 64400 AABK15 RS03755 WP 130525712.1 CCGLIYAVITKRG.....STTLPFGPCLSLGGIATIYLQALF..... 269
NTDB id 1438 BW25113 RS15435 WP 001333829.1 CCGLIYAVITKRG.....STTLPFGPCLSLGGIATLYLQALF..... 269
NTDB id 1111 NGFG RS09220 WP 003689814.1 LIGLVAAIVMR....VAKGRHFAFGPALTVSGWIIFTANDSVWRAVNWWLTHPVR... 286
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 LLGAIIGIILLKLRND..NQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 1061 ABD1 RS18470 WP 001152280.1 LLGAIIGIILLKLRND..NQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 1170 A1552VC RS11080 WP 000418747.1 VVGVIFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKV...IDWYFTTWVGQPL 291
NTDB id 1403 DSB67 RS12675 WP 010643256.1 VVGVIFGLIQLRLQKQGIEKAFPFGPYLAIAGWVSLIWGHQI...LNWYFTSILGV.. 289
consensus
X non conserved
X similar
X ≥ 50% conserved