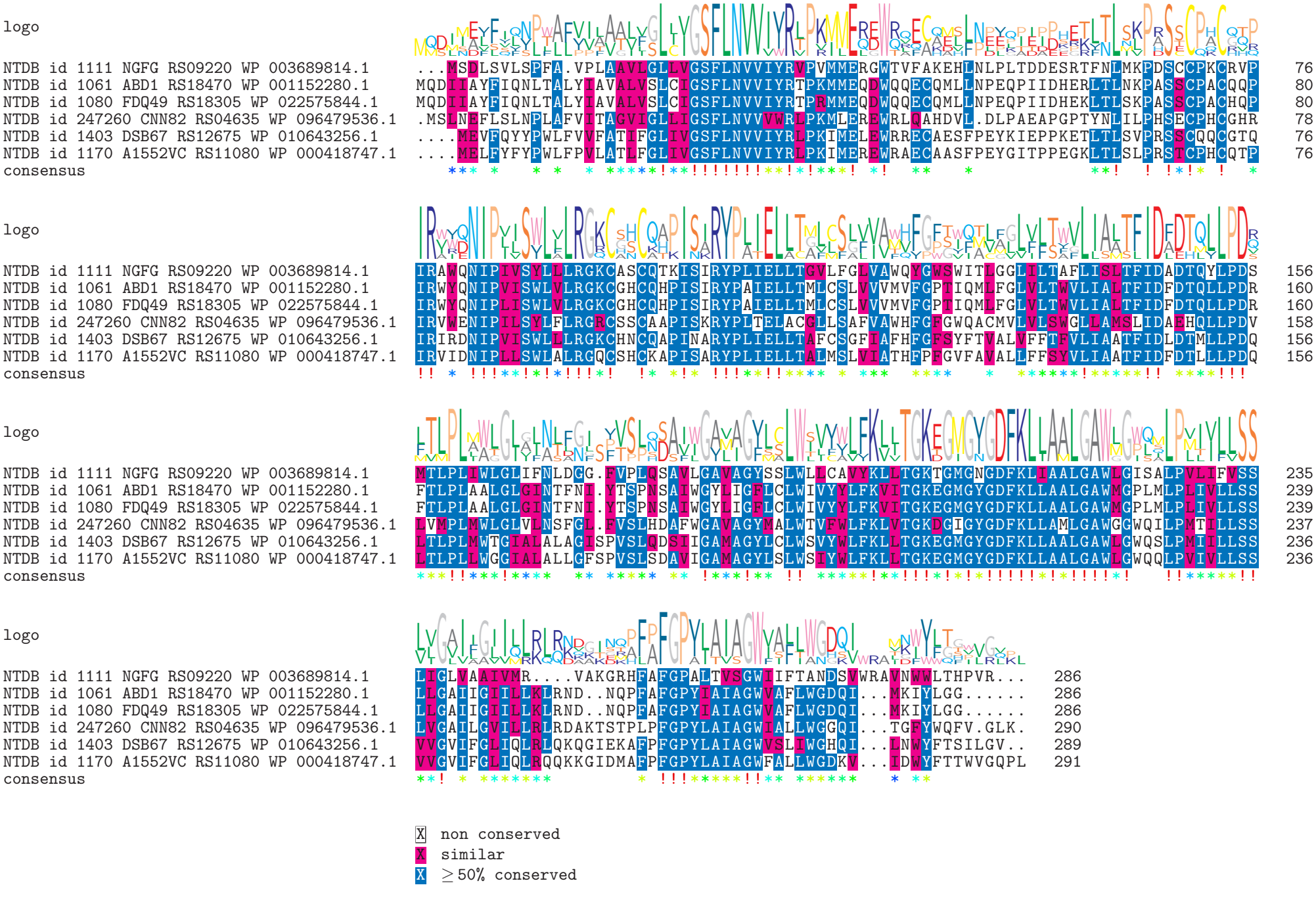
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 ...MSDLSVLSPFA.VPLAAVLGLLVGSFLNVVIYRVPVMMERGWTVFAKEHLNLPLTDDESRTFNLMKPDSCCPKCRVP 76
NTDB id 1061 ABD1 RS18470 WP 001152280.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLNPEQPIIDHERLTLNKPASSCPACQQP 80
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLNPEQPIIDHEKLTLSKPASSCPACHQP 80
NTDB id 247260 CNN82 RS04635 WP 096479536.1 .MSLNEFLSLNPLAFVITAGVIGLLIGSFLNVVVWRLPKMLEREWRLQAHDVL.DLPAEAPGPTYNLILPHSECPHCGHR 78
NTDB id 1403 DSB67 RS12675 WP 010643256.1 ....MEVFQYYPWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFPEYKIEPPKETLTLSVPRSSCQQCGTQ 76
NTDB id 1170 A1552VC RS11080 WP 000418747.1 ....MELFYFYPWLFPVLATLFGLIVGSFLNVVIYRLPKIMEREWRAECAASFPEYGITPPEGKLTLSLPRSTCPHCQTP 76
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 IRAWQNIPIVSYLLLRGKCASCQTKISIRYPLIELLTGVLFGLVAWQYGWSWITLGGLILTAFLISLTFIDADTQYLPDS 156
NTDB id 1061 ABD1 RS18470 WP 001152280.1 IRWYQNIPVISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPDR 160
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 IRWYQNIPLISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPDR 160
NTDB id 247260 CNN82 RS04635 WP 096479536.1 IRVWENIPILSYLFLRGRCSSCAAPISKRYPLTELACGLLSAFVAWHFGFGWQACMVLVLSWGLLAMSLIDAEHQLLPDV 158
NTDB id 1403 DSB67 RS12675 WP 010643256.1 IRIRDNIPVISWLLLRGKCHNCQAPINARYPLIELLTAFCSGFIAFHFGFSYFTVALVFFTFVLIAATFIDLDTMLLPDQ 156
NTDB id 1170 A1552VC RS11080 WP 000418747.1 IRVIDNIPLLSWLALRGQCSHCKAPISARYPLIELLTALMSLVIATHFPFGVFAVALLFFSYVLIAATFIDFDTLLLPDQ 156
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 MTLPLIWLGLIFNLDGG.FVPLQSAVLGAVAGYSSLWLLCAVYKLLTGKTGMGNGDFKLIAALGAWLGISALPVLIFVSS 235
NTDB id 1061 ABD1 RS18470 WP 001152280.1 FTLPLAALGLGINTFNI.YTSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLSS 239
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 FTLPLAALGLGINTFNI.YTSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLSS 239
NTDB id 247260 CNN82 RS04635 WP 096479536.1 LVMPLMWLGLVLNSFGL.FVSLHDAFWGAVAGYMALWTVFWLFKLVTGKDGIGYGDFKLLAMLGAWGGWQILPMTILLSS 237
NTDB id 1403 DSB67 RS12675 WP 010643256.1 LTLPLMWTGIALALAGISPVSLQDSIIGAMAGYLCLWSVYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQSLPMIILLSS 236
NTDB id 1170 A1552VC RS11080 WP 000418747.1 LTLPLLWGGIALALLGFSPVSLSDAVIGAMAGYLSLWSIYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQQLPVIVLLSS 236
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 LIGLVAAIVMR....VAKGRHFAFGPALTVSGWIIFTANDSVWRAVNWWLTHPVR... 286
NTDB id 1061 ABD1 RS18470 WP 001152280.1 LLGAIIGIILLKLRND..NQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 LLGAIIGIILLKLRND..NQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 247260 CNN82 RS04635 WP 096479536.1 LVGAILGVILLRLRDAKTSTPLPFGPYLAIAGWIALLWGGQI...TGFYWQFV.GLK. 290
NTDB id 1403 DSB67 RS12675 WP 010643256.1 VVGVIFGLIQLRLQKQGIEKAFPFGPYLAIAGWVSLIWGHQI...LNWYFTSILGV.. 289
NTDB id 1170 A1552VC RS11080 WP 000418747.1 VVGVIFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKV...IDWYFTTWVGQPL 291
consensus
X non conserved
X similar
X ≥ 50% conserved