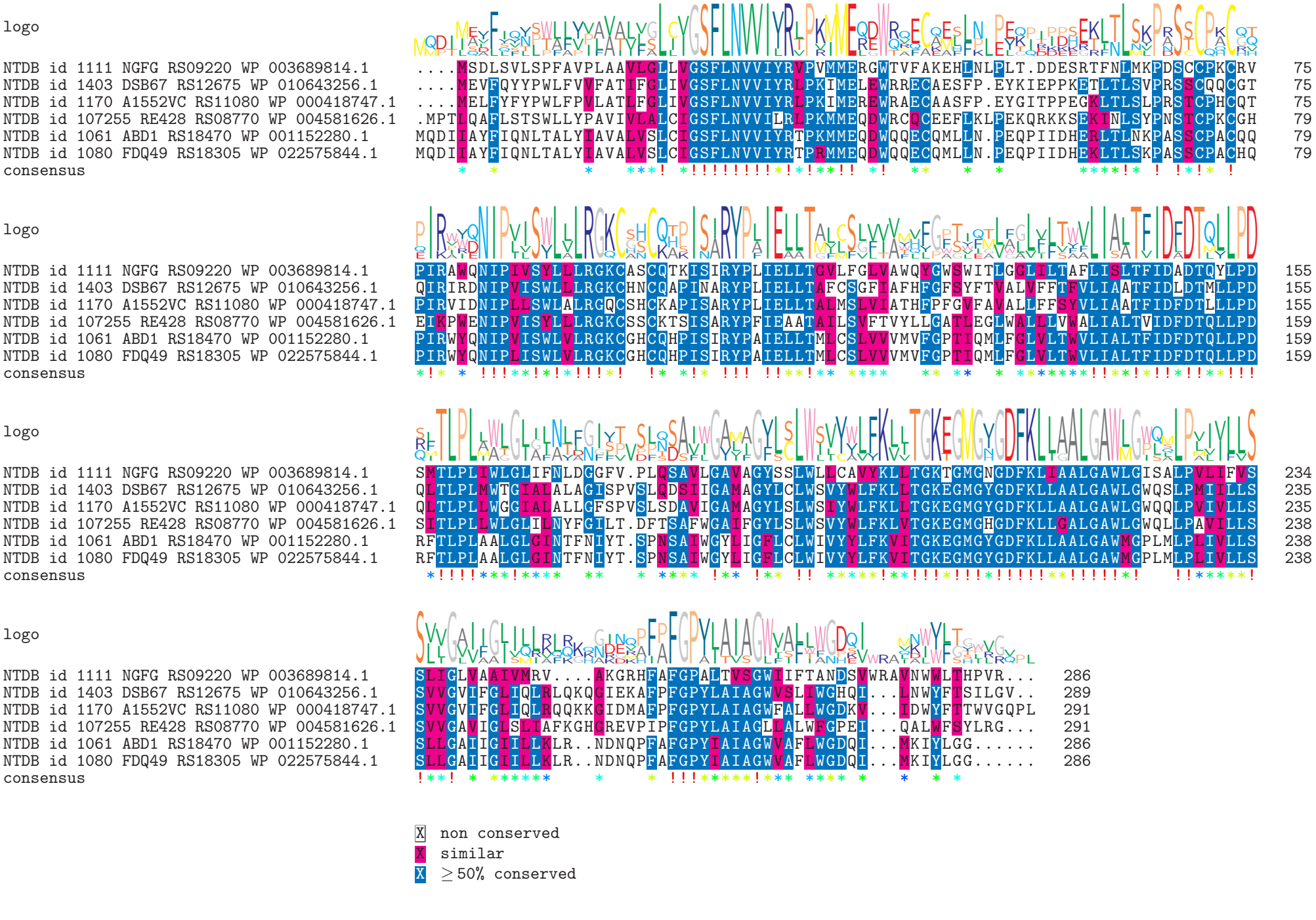
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 ....MSDLSVLSPFAVPLAAVLGLLVGSFLNVVIYRVPVMMERGWTVFAKEHLNLPLT.DDESRTFNLMKPDSCCPKCRV 75
NTDB id 1403 DSB67 RS12675 WP 010643256.1 ....MEVFQYYPWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFP.EYKIEPPKETLTLSVPRSSCQQCGT 75
NTDB id 1170 A1552VC RS11080 WP 000418747.1 ....MELFYFYPWLFPVLATLFGLIVGSFLNVVIYRLPKIMEREWRAECAASFP.EYGITPPEGKLTLSLPRSTCPHCQT 75
NTDB id 107255 RE428 RS08770 WP 004581626.1 .MPTLQAFLSTSWLLYPAVIVLALCIGSFLNVVILRLPKMMEQDWRCQCEEFLKLPEKQRKKSEKINLSYPNSTCPKCGH 79
NTDB id 1061 ABD1 RS18470 WP 001152280.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLN.PEQPIIDHERLTLNKPASSCPACQQ 79
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLN.PEQPIIDHEKLTLSKPASSCPACHQ 79
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 PIRAWQNIPIVSYLLLRGKCASCQTKISIRYPLIELLTGVLFGLVAWQYGWSWITLGGLILTAFLISLTFIDADTQYLPD 155
NTDB id 1403 DSB67 RS12675 WP 010643256.1 QIRIRDNIPVISWLLLRGKCHNCQAPINARYPLIELLTAFCSGFIAFHFGFSYFTVALVFFTFVLIAATFIDLDTMLLPD 155
NTDB id 1170 A1552VC RS11080 WP 000418747.1 PIRVIDNIPLLSWLALRGQCSHCKAPISARYPLIELLTALMSLVIATHFPFGVFAVALLFFSYVLIAATFIDFDTLLLPD 155
NTDB id 107255 RE428 RS08770 WP 004581626.1 EIKPWENIPVISYLLLRGKCSSCKTSISARYPFIEAATAILSVFTVYLLGATLEGLWALLLVWALIALTVIDFDTQLLPD 159
NTDB id 1061 ABD1 RS18470 WP 001152280.1 PIRWYQNIPVISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPD 159
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 PIRWYQNIPLISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPD 159
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 SMTLPLIWLGLIFNLDCGFV.PLQSAVLGAVAGYSSLWLLCAVYKLLTGKTGMGNGDFKLIAALGAWLGISALPVLIFVS 234
NTDB id 1403 DSB67 RS12675 WP 010643256.1 QLTLPLMWTGIALALAGISPVSLQDSIIGAMAGYLCLWSVYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQSLPMIILLS 235
NTDB id 1170 A1552VC RS11080 WP 000418747.1 QLTLPLLWGGIALALLGFSPVSLSDAVIGAMAGYLSLWSIYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQQLPVIVLLS 235
NTDB id 107255 RE428 RS08770 WP 004581626.1 SITLPLLWLGLILNYFGILT.DFTSAFWGAIFGYLSLWSVYWLFKLVTGKEGMGHGDFKLLGALGAWLGWQLLPAVILLS 238
NTDB id 1061 ABD1 RS18470 WP 001152280.1 RFTLPLAALGLGINTFNIYT.SPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLS 238
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 RFTLPLAALGLGINTFNIYT.SPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLS 238
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 SLIGLVAAIVMRV....AKGRHFAFGPALTVSGWIIFTANDSVWRAVNWWLTHPVR... 286
NTDB id 1403 DSB67 RS12675 WP 010643256.1 SVVGVIFGLIQLRLQKQGIEKAFPFGPYLAIAGWVSLIWGHQI...LNWYFTSILGV.. 289
NTDB id 1170 A1552VC RS11080 WP 000418747.1 SVVGVIFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKV...IDWYFTTWVGQPL 291
NTDB id 107255 RE428 RS08770 WP 004581626.1 SVVGAVIGLSLIAFKGHGREVPIPFGPYLAIAGLLALWFGPEI...QALWFSYLRG... 291
NTDB id 1061 ABD1 RS18470 WP 001152280.1 SLLGAIIGIILLKLR..NDNQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 SLLGAIIGIILLKLR..NDNQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
consensus
X non conserved
X similar
X ≥ 50% conserved