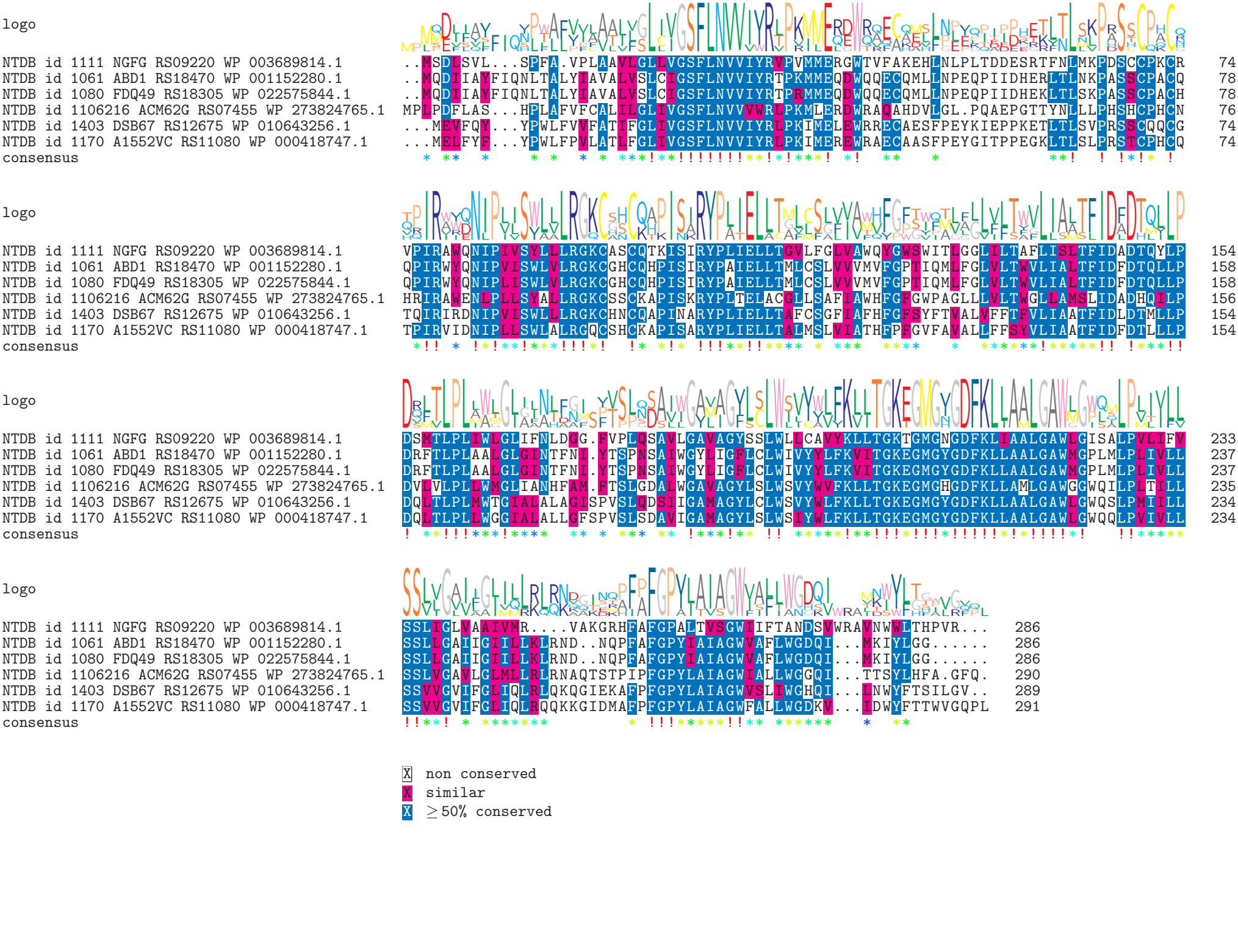
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 ..MSDLSVL...SPFA.VPLAAVLGLLVGSFLNVVIYRVPVMMERGWTVFAKEHLNLPLTDDESRTFNLMKPDSCCPKCR 74
NTDB id 1061 ABD1 RS18470 WP 001152280.1 ..MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLNPEQPIIDHERLTLNKPASSCPACQ 78
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 ..MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLNPEQPIIDHEKLTLSKPASSCPACH 78
NTDB id 1106216 ACM62G RS07455 WP 273824765.1 MPLPDFLAS...HPLAFVFCALILGLIVGSFLNVVVWRLPKMLERDWRAQAHDVLGL.PQAEPGTTYNLLLPHSHCPHCN 76
NTDB id 1403 DSB67 RS12675 WP 010643256.1 ...MEVFQY...YPWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFPEYKIEPPKETLTLSVPRSSCQQCG 74
NTDB id 1170 A1552VC RS11080 WP 000418747.1 ...MELFYF...YPWLFPVLATLFGLIVGSFLNVVIYRLPKIMEREWRAECAASFPEYGITPPEGKLTLSLPRSTCPHCQ 74
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 VPIRAWQNIPIVSYLLLRGKCASCQTKISIRYPLIELLTGVLFGLVAWQYGWSWITLGGLILTAFLISLTFIDADTQYLP 154
NTDB id 1061 ABD1 RS18470 WP 001152280.1 QPIRWYQNIPVISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLP 158
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 QPIRWYQNIPLISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLP 158
NTDB id 1106216 ACM62G RS07455 WP 273824765.1 HRIRAWENLPLLSYALLRGKCSSCKAPISKRYPLTELACGLLSAFIAWHFGFGWPAGLLLVLTWGLLAMSLIDADHQILP 156
NTDB id 1403 DSB67 RS12675 WP 010643256.1 TQIRIRDNIPVISWLLLRGKCHNCQAPINARYPLIELLTAFCSGFIAFHFGFSYFTVALVFFTFVLIAATFIDLDTMLLP 154
NTDB id 1170 A1552VC RS11080 WP 000418747.1 TPIRVIDNIPLLSWLALRGQCSHCKAPISARYPLIELLTALMSLVIATHFPFGVFAVALLFFSYVLIAATFIDFDTLLLP 154
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 DSMTLPLIWLGLIFNLDGG.FVPLQSAVLGAVAGYSSLWLLCAVYKLLTGKTGMGNGDFKLIAALGAWLGISALPVLIFV 233
NTDB id 1061 ABD1 RS18470 WP 001152280.1 DRFTLPLAALGLGINTFNI.YTSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLL 237
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 DRFTLPLAALGLGINTFNI.YTSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLL 237
NTDB id 1106216 ACM62G RS07455 WP 273824765.1 DVLVLPLLWMGLIANHFAM.FTSLGDALWGAVAGYLSLWSVYWVFKLLTGKEGMGHGDFKLLAMLGAWGGWQILPLTILL 235
NTDB id 1403 DSB67 RS12675 WP 010643256.1 DQLTLPLMWTGIALALAGISPVSLQDSIIGAMAGYLCLWSVYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQSLPMIILL 234
NTDB id 1170 A1552VC RS11080 WP 000418747.1 DQLTLPLLWGGIALALLGFSPVSLSDAVIGAMAGYLSLWSIYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQQLPVIVLL 234
consensus
logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 SSLIGLVAAIVMR....VAKGRHFAFGPALTVSGWIIFTANDSVWRAVNWWLTHPVR... 286
NTDB id 1061 ABD1 RS18470 WP 001152280.1 SSLLGAIIGIILLKLRND..NQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 SSLLGAIIGIILLKLRND..NQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 1106216 ACM62G RS07455 WP 273824765.1 SSLVGAVLGLMLLRLRNAQTSTPIPFGPYLAIAGWIALLWGGQI...TTSYLHFA.GFQ. 290
NTDB id 1403 DSB67 RS12675 WP 010643256.1 SSVVGVIFGLIQLRLQKQGIEKAFPFGPYLAIAGWVSLIWGHQI...LNWYFTSILGV.. 289
NTDB id 1170 A1552VC RS11080 WP 000418747.1 SSVVGVIFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKV...IDWYFTTWVGQPL 291
consensus
X non conserved
X similar
X ≥ 50% conserved